

# Create DEMO table as csv file

*mja@statgroup.dk*

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## Create turtle version of example files

This script creates the result and codelist for a simple DEMO table.

```
library(rrdfancillary)
library(rrdfcdisc)
library(rrdfqb)
library(rrdfqbcrnd0)
devtools::load_all(pkg="../..")
```

```
## Loading rrdfqbcrndex
```

All files are stored in the directory

```
targetDir<- system.file("extdata/sample-rdf", package="rrdfqbcrndex")
(targetDir)
```

```
## [1] "/home/ma/projects/rrdfqbcrnd0/rrdfqbcrndex/inst/extdata/sample-rdf"
```

## RDF data cubes specified in workbook

First, get the workbook.

```
RDFCubeWorkbook<- system.file("extdata/sample-cfg", "RDFCubeWorkbook.xlsx", package="rrdfqbcrndex")
(RDFCubeWorkbook)
```

```
## [1] "/home/ma/projects/rrdfqbcrnd0/rrdfqbcrndex/inst/extdata/sample-cfg/RDFCubeWorkbook.xlsx"
```

Create the DM sample cube.

```
dm.cube.fn<- BuildCubeFromWorkbook(RDFCubeWorkbook, "DM" )
```

```

## [1] "prefix code: <http://www.example.org/dc/code/>\nprefix cts: <http://rdf.cdisc.org/ct/schema#>\n"
##      DataStructureDefinition      dimension
## 1      ds:dsd-DM      crnd-dimension:factor
## 2      ds:dsd-DM      crnd-dimension:factor
## 3      ds:dsd-DM      crnd-dimension:factor
## 4      ds:dsd-DM      crnd-dimension:factor
## 5      ds:dsd-DM      crnd-dimension:factor
## 6      ds:dsd-DM      crnd-dimension:factor
## 7      ds:dsd-DM      crnd-dimension:procedure
## 8      ds:dsd-DM      crnd-dimension:procedure
## 9      ds:dsd-DM      crnd-dimension:procedure
## 10     ds:dsd-DM      crnd-dimension:procedure
## 11     ds:dsd-DM      crnd-dimension:procedure
## 12     ds:dsd-DM      crnd-dimension:procedure
## 13     ds:dsd-DM      crnd-dimension:procedure
## 14     ds:dsd-DM      crnd-dimension:race
## 15     ds:dsd-DM      crnd-dimension:race
## 16     ds:dsd-DM      crnd-dimension:race
## 17     ds:dsd-DM      crnd-dimension:race
## 18     ds:dsd-DM      crnd-dimension:race
## 19     ds:dsd-DM      crnd-dimension:race
## 20     ds:dsd-DM      crnd-dimension:race
## 21     ds:dsd-DM      crnd-dimension:saffl
## 22     ds:dsd-DM      crnd-dimension:saffl
## 23     ds:dsd-DM      crnd-dimension:saffl
## 24     ds:dsd-DM      crnd-dimension:sex
## 25     ds:dsd-DM      crnd-dimension:sex
## 26     ds:dsd-DM      crnd-dimension:sex
## 27     ds:dsd-DM      crnd-dimension:sex
## 28     ds:dsd-DM      crnd-dimension:sex
## 29     ds:dsd-DM      crnd-dimension:sex
## 30     ds:dsd-DM      crnd-dimension:trt01a
## 31     ds:dsd-DM      crnd-dimension:trt01a
## 32     ds:dsd-DM      crnd-dimension:trt01a
## 33     ds:dsd-DM      crnd-dimension:trt01a
## 34     ds:dsd-DM      crnd-dimension:trt01a
##      cprefLabel
## 1      Codelist scheme: factor
## 2      Codelist scheme: factor
## 3      Codelist scheme: factor
## 4      Codelist scheme: factor
## 5      Codelist scheme: factor
## 6      Codelist scheme: factor
## 7      Codelist scheme: procedure
## 8      Codelist scheme: procedure
## 9      Codelist scheme: procedure
## 10     Codelist scheme: procedure
## 11     Codelist scheme: procedure
## 12     Codelist scheme: procedure
## 13     Codelist scheme: procedure
## 14     Codelist scheme: race
## 15     Codelist scheme: race
## 16     Codelist scheme: race
## 17     Codelist scheme: race

```

```

## 18      Codelist scheme: race
## 19      Codelist scheme: race
## 20      Codelist scheme: race
## 21      Codelist scheme: saffl
## 22      Codelist scheme: saffl
## 23      Codelist scheme: saffl
## 24      Codelist scheme: sex
## 25      Codelist scheme: sex
## 26      Codelist scheme: sex
## 27      Codelist scheme: sex
## 28      Codelist scheme: sex
## 29      Codelist scheme: sex
## 30      Codelist scheme: trt01a
## 31      Codelist scheme: trt01a
## 32      Codelist scheme: trt01a
## 33      Codelist scheme: trt01a
## 34      Codelist scheme: trt01a
##
##                                     cl
## 1                                     code:factor-AGE
## 2                                     code:factor-WEIGHTBL
## 3                                     code:factor-_ALL_
## 4                                     code:factor-_NONMISS_
## 5                                     code:factor-proportion
## 6                                     code:factor-quantity
## 7                                     code:procedure-count
## 8                                     code:procedure-max
## 9                                     code:procedure-mean
## 10                                    code:procedure-median
## 11                                    code:procedure-min
## 12                                    code:procedure-percent
## 13                                    code:procedure-stdev
## 14      code:race-AMERICAN_INDIAN_OR_ALASKA_NATIVE
## 15                                    code:race-ASIAN
## 16      code:race-BLACK_OR_AFRICAN_AMERICAN
## 17 code:race-NATIVE_HAWAIIAN_OR_OTHER_PACIFIC_ISLANDER
## 18                                    code:race-WHITE
## 19                                    code:race-_ALL_
## 20                                    code:race-_NONMISS_
## 21                                    code:saffl-Y
## 22                                    code:saffl-_ALL_
## 23      code:saffl-_NONMISS_
## 24                                    code:sex-F
## 25                                    code:sex-M
## 26                                    code:sex-U
## 27                                    code:sex-UN
## 28                                    code:sex-_ALL_
## 29                                    code:sex-_NONMISS_
## 30      code:trt01a-Placebo
## 31      code:trt01a-Xanomeline_High_Dose
## 32      code:trt01a-Xanomeline_Low_Dose
## 33      code:trt01a-_ALL_
## 34      code:trt01a-_NONMISS_
##      clprefLabel      vn      vct
## 1      AGE      factor DATA

```

## 2	WEIGHTBL	factor	DATA
## 3	_ALL_	factor	DATA
## 4	_NONMISS_	factor	DATA
## 5	proportion	factor	DATA
## 6	quantity	factor	DATA
## 7	count	procedure	DATA
## 8	max	procedure	DATA
## 9	mean	procedure	DATA
## 10	median	procedure	DATA
## 11	min	procedure	DATA
## 12	percent	procedure	DATA
## 13	stdev	procedure	DATA
## 14	AMERICAN INDIAN OR ALASKA NATIVE	race	SDTM
## 15	ASIAN	race	SDTM
## 16	BLACK OR AFRICAN AMERICAN	race	SDTM
## 17	NATIVE HAWAIIAN OR OTHER PACIFIC ISLANDER	race	SDTM
## 18	WHITE	race	SDTM
## 19	_ALL_	race	SDTM
## 20	_NONMISS_	race	SDTM
## 21	Y	saffl	DATA
## 22	_ALL_	saffl	DATA
## 23	_NONMISS_	saffl	DATA
## 24	F	sex	SDTM
## 25	M	sex	SDTM
## 26	U	sex	SDTM
## 27	UN	sex	SDTM
## 28	_ALL_	sex	SDTM
## 29	_NONMISS_	sex	SDTM
## 30	Placebo	trt01a	DATA
## 31	Xanomeline High Dose	trt01a	DATA
## 32	Xanomeline Low Dose	trt01a	DATA
## 33	_ALL_	trt01a	DATA
## 34	_NONMISS_	trt01a	DATA
##	vnop		vnval
## 1	==		AGE
## 2	==		WEIGHTBL
## 3	<NA>		<NA>
## 4	<NA>		<NA>
## 5	==		proportion
## 6	==		quantity
## 7	==		count
## 8	==		max
## 9	==		mean
## 10	==		median
## 11	==		min
## 12	==		percent
## 13	==		stdev
## 14	rrdfqbcrnd0:ADSL_RACE	AMERICAN INDIAN OR ALASKA NATIVE	
## 15	rrdfqbcrnd0:ADSL_RACE	ASIAN	
## 16	rrdfqbcrnd0:ADSL_RACE	BLACK OR AFRICAN AMERICAN	
## 17	rrdfqbcrnd0:ADSL_RACE	NATIVE HAWAIIAN OR OTHER PACIFIC ISLANDER	
## 18	rrdfqbcrnd0:ADSL_RACE	WHITE	
## 19	rrdfqbcrnd0:ADSL_RACE	<NA>	
## 20	rrdfqbcrnd0:ADSL_RACE	<NA>	

```

## 21 rrdfqbcrnd0:ADSL_SAFFL Y
## 22 rrdfqbcrnd0:ADSL_SAFFL <NA>
## 23 rrdfqbcrnd0:ADSL_SAFFL <NA>
## 24 rrdfqbcrnd0:ADSL_SEX F
## 25 rrdfqbcrnd0:ADSL_SEX M
## 26 rrdfqbcrnd0:ADSL_SEX U
## 27 rrdfqbcrnd0:ADSL_SEX UN
## 28 rrdfqbcrnd0:ADSL_SEX <NA>
## 29 rrdfqbcrnd0:ADSL_SEX <NA>
## 30 rrdfqbcrnd0:ADSL_TRT01A Placebo
## 31 rrdfqbcrnd0:ADSL_TRT01A Xanomeline High Dose
## 32 rrdfqbcrnd0:ADSL_TRT01A Xanomeline Low Dose
## 33 rrdfqbcrnd0:ADSL_TRT01A <NA>
## 34 rrdfqbcrnd0:ADSL_TRT01A <NA>
## crnd-dimension:factor crnd-dimension:factor crnd-dimension:factor crnd-dimension:factor crnd-dimension:factor

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): stdev --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): stdev --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): stdev --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): stdev --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): stdev --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): stdev --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHTBL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

```

```
## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value
```

```
cat("DM cube stored as ", dm.cube.fn, "\n")
```

```
## DM cube stored as /tmp/Rtmpz1JmRR/DC-DM-R-V-0-5-2.ttl
```

```
targetFile<- file.path(targetDir,"DC-DM-sample.ttl")

if (file.copy( dm.cube.fn, targetFile, overwrite=TRUE)) {
  cat("DM cube copied to ", normalizePath(targetFile), "\n")
}
```

```
## DM cube copied to /home/ma/projects/rrdfqbc rnd0/rrdfqbc rndex/inst/extdata/sample-rdf/DC-DM-sample.t
```

Create the AE sample cube.

```
ae.cube.fn<- BuildCubeFromWorkbook(RDFCubeWorkbook, "AE" )
```

```
## [1] "prefix code: <http://www.example.org/dc/code/>\nprefix cts: <http://rdf.cd disc.org/ct/schema#>\n"
##      DataStructureDefinition      dimension
## 1      ds:dsd-AE      crnd-dimension:ae decod
## 2      ds:dsd-AE      crnd-dimension:ae decod
## 3      ds:dsd-AE      crnd-dimension:ae decod
## 4      ds:dsd-AE      crnd-dimension:ae decod
## 5      ds:dsd-AE      crnd-dimension:ae decod
## 6      ds:dsd-AE      crnd-dimension:ae decod
## 7      ds:dsd-AE      crnd-dimension:ae decod
## 8      ds:dsd-AE      crnd-dimension:ae decod
## 9      ds:dsd-AE      crnd-dimension:ae decod
## 10     ds:dsd-AE      crnd-dimension:ae decod
## 11     ds:dsd-AE      crnd-dimension:ae decod
## 12     ds:dsd-AE      crnd-dimension:ae decod
## 13     ds:dsd-AE      crnd-dimension:ae decod
## 14     ds:dsd-AE      crnd-dimension:ae decod
```

[illegible]

## 69	ds:dsd-AE	crnd-dimension:aedecod
## 70	ds:dsd-AE	crnd-dimension:aedecod
## 71	ds:dsd-AE	crnd-dimension:aedecod
## 72	ds:dsd-AE	crnd-dimension:aedecod
## 73	ds:dsd-AE	crnd-dimension:aedecod
## 74	ds:dsd-AE	crnd-dimension:aedecod
## 75	ds:dsd-AE	crnd-dimension:aedecod
## 76	ds:dsd-AE	crnd-dimension:aedecod
## 77	ds:dsd-AE	crnd-dimension:aedecod
## 78	ds:dsd-AE	crnd-dimension:aedecod
## 79	ds:dsd-AE	crnd-dimension:aedecod
## 80	ds:dsd-AE	crnd-dimension:aedecod
## 81	ds:dsd-AE	crnd-dimension:aedecod
## 82	ds:dsd-AE	crnd-dimension:aedecod
## 83	ds:dsd-AE	crnd-dimension:aedecod
## 84	ds:dsd-AE	crnd-dimension:aedecod
## 85	ds:dsd-AE	crnd-dimension:aedecod
## 86	ds:dsd-AE	crnd-dimension:aedecod
## 87	ds:dsd-AE	crnd-dimension:aedecod
## 88	ds:dsd-AE	crnd-dimension:aedecod
## 89	ds:dsd-AE	crnd-dimension:aedecod
## 90	ds:dsd-AE	crnd-dimension:aedecod
## 91	ds:dsd-AE	crnd-dimension:aedecod
## 92	ds:dsd-AE	crnd-dimension:aedecod
## 93	ds:dsd-AE	crnd-dimension:aedecod
## 94	ds:dsd-AE	crnd-dimension:aedecod
## 95	ds:dsd-AE	crnd-dimension:aedecod
## 96	ds:dsd-AE	crnd-dimension:aedecod
## 97	ds:dsd-AE	crnd-dimension:aedecod
## 98	ds:dsd-AE	crnd-dimension:aedecod
## 99	ds:dsd-AE	crnd-dimension:aedecod
## 100	ds:dsd-AE	crnd-dimension:aedecod
## 101	ds:dsd-AE	crnd-dimension:aedecod
## 102	ds:dsd-AE	crnd-dimension:aedecod
## 103	ds:dsd-AE	crnd-dimension:aedecod
## 104	ds:dsd-AE	crnd-dimension:aedecod
## 105	ds:dsd-AE	crnd-dimension:aedecod
## 106	ds:dsd-AE	crnd-dimension:aedecod
## 107	ds:dsd-AE	crnd-dimension:aedecod
## 108	ds:dsd-AE	crnd-dimension:aedecod
## 109	ds:dsd-AE	crnd-dimension:aedecod
## 110	ds:dsd-AE	crnd-dimension:aedecod
## 111	ds:dsd-AE	crnd-dimension:aedecod
## 112	ds:dsd-AE	crnd-dimension:aedecod
## 113	ds:dsd-AE	crnd-dimension:aedecod
## 114	ds:dsd-AE	crnd-dimension:aedecod
## 115	ds:dsd-AE	crnd-dimension:aedecod
## 116	ds:dsd-AE	crnd-dimension:aedecod
## 117	ds:dsd-AE	crnd-dimension:aedecod
## 118	ds:dsd-AE	crnd-dimension:aedecod
## 119	ds:dsd-AE	crnd-dimension:aedecod
## 120	ds:dsd-AE	crnd-dimension:aedecod
## 121	ds:dsd-AE	crnd-dimension:aedecod
## 122	ds:dsd-AE	crnd-dimension:aedecod

[illegible]



[illegible]

## 231	ds:dsd-AE	crnd-dimension:aedecod
## 232	ds:dsd-AE	crnd-dimension:aedecod
## 233	ds:dsd-AE	crnd-dimension:aedecod
## 234	ds:dsd-AE	crnd-dimension:aedecod
## 235	ds:dsd-AE	crnd-dimension:aedecod
## 236	ds:dsd-AE	crnd-dimension:aedecod
## 237	ds:dsd-AE	crnd-dimension:aedecod
## 238	ds:dsd-AE	crnd-dimension:aedecod
## 239	ds:dsd-AE	crnd-dimension:aedecod
## 240	ds:dsd-AE	crnd-dimension:aedecod
## 241	ds:dsd-AE	crnd-dimension:aedecod
## 242	ds:dsd-AE	crnd-dimension:aedecod
## 243	ds:dsd-AE	crnd-dimension:aedecod
## 244	ds:dsd-AE	crnd-dimension:aedecod
## 245	ds:dsd-AE	crnd-dimension:aesoc
## 246	ds:dsd-AE	crnd-dimension:aesoc
## 247	ds:dsd-AE	crnd-dimension:aesoc
## 248	ds:dsd-AE	crnd-dimension:aesoc
## 249	ds:dsd-AE	crnd-dimension:aesoc
## 250	ds:dsd-AE	crnd-dimension:aesoc
## 251	ds:dsd-AE	crnd-dimension:aesoc
## 252	ds:dsd-AE	crnd-dimension:aesoc
## 253	ds:dsd-AE	crnd-dimension:aesoc
## 254	ds:dsd-AE	crnd-dimension:aesoc
## 255	ds:dsd-AE	crnd-dimension:aesoc
## 256	ds:dsd-AE	crnd-dimension:aesoc
## 257	ds:dsd-AE	crnd-dimension:aesoc
## 258	ds:dsd-AE	crnd-dimension:aesoc
## 259	ds:dsd-AE	crnd-dimension:aesoc
## 260	ds:dsd-AE	crnd-dimension:aesoc
## 261	ds:dsd-AE	crnd-dimension:aesoc
## 262	ds:dsd-AE	crnd-dimension:aesoc
## 263	ds:dsd-AE	crnd-dimension:aesoc
## 264	ds:dsd-AE	crnd-dimension:aesoc
## 265	ds:dsd-AE	crnd-dimension:aesoc
## 266	ds:dsd-AE	crnd-dimension:aesoc
## 267	ds:dsd-AE	crnd-dimension:aesoc
## 268	ds:dsd-AE	crnd-dimension:aesoc
## 269	ds:dsd-AE	crnd-dimension:aesoc
## 270	ds:dsd-AE	crnd-dimension:factor
## 271	ds:dsd-AE	crnd-dimension:factor
## 272	ds:dsd-AE	crnd-dimension:factor
## 273	ds:dsd-AE	crnd-dimension:factor
## 274	ds:dsd-AE	crnd-dimension:procedure
## 275	ds:dsd-AE	crnd-dimension:procedure
## 276	ds:dsd-AE	crnd-dimension:saffl
## 277	ds:dsd-AE	crnd-dimension:saffl
## 278	ds:dsd-AE	crnd-dimension:saffl
## 279	ds:dsd-AE	crnd-dimension:trta
## 280	ds:dsd-AE	crnd-dimension:trta
## 281	ds:dsd-AE	crnd-dimension:trta
## 282	ds:dsd-AE	crnd-dimension:trta
## 283	ds:dsd-AE	crnd-dimension:trta
##	cprefLabel	

```
## 1      Codelist scheme: aeecod
## 2      Codelist scheme: aeecod
## 3      Codelist scheme: aeecod
## 4      Codelist scheme: aeecod
## 5      Codelist scheme: aeecod
## 6      Codelist scheme: aeecod
## 7      Codelist scheme: aeecod
## 8      Codelist scheme: aeecod
## 9      Codelist scheme: aeecod
## 10     Codelist scheme: aeecod
## 11     Codelist scheme: aeecod
## 12     Codelist scheme: aeecod
## 13     Codelist scheme: aeecod
## 14     Codelist scheme: aeecod
## 15     Codelist scheme: aeecod
## 16     Codelist scheme: aeecod
## 17     Codelist scheme: aeecod
## 18     Codelist scheme: aeecod
## 19     Codelist scheme: aeecod
## 20     Codelist scheme: aeecod
## 21     Codelist scheme: aeecod
## 22     Codelist scheme: aeecod
## 23     Codelist scheme: aeecod
## 24     Codelist scheme: aeecod
## 25     Codelist scheme: aeecod
## 26     Codelist scheme: aeecod
## 27     Codelist scheme: aeecod
## 28     Codelist scheme: aeecod
## 29     Codelist scheme: aeecod
## 30     Codelist scheme: aeecod
## 31     Codelist scheme: aeecod
## 32     Codelist scheme: aeecod
## 33     Codelist scheme: aeecod
## 34     Codelist scheme: aeecod
## 35     Codelist scheme: aeecod
## 36     Codelist scheme: aeecod
## 37     Codelist scheme: aeecod
## 38     Codelist scheme: aeecod
## 39     Codelist scheme: aeecod
## 40     Codelist scheme: aeecod
## 41     Codelist scheme: aeecod
## 42     Codelist scheme: aeecod
## 43     Codelist scheme: aeecod
## 44     Codelist scheme: aeecod
## 45     Codelist scheme: aeecod
## 46     Codelist scheme: aeecod
## 47     Codelist scheme: aeecod
## 48     Codelist scheme: aeecod
## 49     Codelist scheme: aeecod
## 50     Codelist scheme: aeecod
## 51     Codelist scheme: aeecod
## 52     Codelist scheme: aeecod
## 53     Codelist scheme: aeecod
## 54     Codelist scheme: aeecod
```

```
## 55    Codelist scheme: aeecod
## 56    Codelist scheme: aeecod
## 57    Codelist scheme: aeecod
## 58    Codelist scheme: aeecod
## 59    Codelist scheme: aeecod
## 60    Codelist scheme: aeecod
## 61    Codelist scheme: aeecod
## 62    Codelist scheme: aeecod
## 63    Codelist scheme: aeecod
## 64    Codelist scheme: aeecod
## 65    Codelist scheme: aeecod
## 66    Codelist scheme: aeecod
## 67    Codelist scheme: aeecod
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## 163 Codelist scheme: aedecod
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## 270 Codelist scheme: factor
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## 271      Codelist scheme: factor
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## 274 Codelist scheme: procedure
## 275 Codelist scheme: procedure
## 276      Codelist scheme: saffl
## 277      Codelist scheme: saffl
## 278      Codelist scheme: saffl
## 279      Codelist scheme: trta
## 280      Codelist scheme: trta
## 281      Codelist scheme: trta
## 282      Codelist scheme: trta
## 283      Codelist scheme: trta
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##
## 1          code:aedecod-ABDOMINAL_DISCOMFORT
## 2          code:aedecod-ABDOMINAL_PAIN
## 3          code:aedecod-ACROCHORDON_EXCISION
## 4          code:aedecod-ACTINIC_KERATOSIS
## 5          code:aedecod-AGITATION
## 6          code:aedecod-ALCOHOL_USE
## 7          code:aedecod-ALLERGIC_GRANULOMATOUS_ANGIITIS
## 8          code:aedecod-ALOPECIA
## 9          code:aedecod-AMNESIA
## 10         code:aedecod-ANXIETY
## 11         code:aedecod-APPLICATION_SITE_BLEEDING
## 12         code:aedecod-APPLICATION_SITE_DERMATITIS
## 13         code:aedecod-APPLICATION_SITE_DESQUMATION
## 14         code:aedecod-APPLICATION_SITE_DISCHARGE
## 15         code:aedecod-APPLICATION_SITE_DISCOLOURATION
## 16         code:aedecod-APPLICATION_SITE_ERYTHEMA
## 17         code:aedecod-APPLICATION_SITE_INDURATION
## 18         code:aedecod-APPLICATION_SITE_IRRITATION
## 19         code:aedecod-APPLICATION_SITE_PAIN
## 20         code:aedecod-APPLICATION_SITE_PERSPIRATION
## 21         code:aedecod-APPLICATION_SITE_PRURITUS
## 22         code:aedecod-APPLICATION_SITE_REACTION
## 23         code:aedecod-APPLICATION_SITE_SWELLING
## 24         code:aedecod-APPLICATION_SITE_URTICARIA
## 25         code:aedecod-APPLICATION_SITE_VESICLES
## 26         code:aedecod-APPLICATION_SITE_WARMTH
## 27         code:aedecod-ARTHRALGIA
## 28         code:aedecod-ARTHRITIS
## 29         code:aedecod-ASTHENIA
## 30         code:aedecod-ATRIAL_FIBRILLATION
## 31         code:aedecod-ATRIAL_FLUTTER
## 32         code:aedecod-ATRIAL_HYPERTROPHY
## 33         code:aedecod-ATRIOVENTRICULAR_BLOCK_FIRST_DEGREE
## 34         code:aedecod-ATRIOVENTRICULAR_BLOCK_SECOND_DEGREE
## 35         code:aedecod-BACK_PAIN
## 36         code:aedecod-BALANCE_DISORDER
## 37         code:aedecod-BENIGN_PROSTATIC_HYPERPLASIA
## 38         code:aedecod-BIOPSY
## 39         code:aedecod-BIOPSY_PROSTATE
## 40         code:aedecod-BLISTER

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## 41         code:aedecod-BLOOD_ALKALINE_PHOSPHATASE_INCREASED
## 42         code:aedecod-BLOOD_CHOLESTEROL_INCREASED
## 43 code:aedecod-BLOOD_CREATINE_PHOSPHOKINASE_INCREASED
## 44         code:aedecod-BLOOD_GLUCOSE_INCREASED
## 45         code:aedecod-BLOOD_URINE_PRESENT
## 46         code:aedecod-BODY_TEMPERATURE_INCREASED
## 47         code:aedecod-BRADYCARDIA
## 48         code:aedecod-BRONCHITIS
## 49         code:aedecod-BUNDLE_BRANCH_BLOCK_LEFT
## 50         code:aedecod-BUNDLE_BRANCH_BLOCK_RIGHT
## 51         code:aedecod-BURNING_SENSATION
## 52         code:aedecod-CALCULUS_URETHRAL
## 53         code:aedecod-CARDIAC_DISORDER
## 54         code:aedecod-CARDIAC_FAILURE_CONGESTIVE
## 55         code:aedecod-CATARACT_OPERATION
## 56         code:aedecod-CELLULITIS
## 57         code:aedecod-CERUMEN_IMPACTION
## 58         code:aedecod-CERVICITIS
## 59         code:aedecod-CHEST_DISCOMFORT
## 60         code:aedecod-CHEST_PAIN
## 61         code:aedecod-CHILLS
## 62         code:aedecod-COGNITIVE_DISORDER
## 63         code:aedecod-COLD_SWEAT
## 64         code:aedecod-COLON_CANCER
## 65         code:aedecod-COMPLETED_SUICIDE
## 66         code:aedecod-COMPLEX_PARTIAL_SEIZURES
## 67         code:aedecod-CONFUSIONAL_STATE
## 68         code:aedecod-CONJUNCTIVAL_HAEMORRHAGE
## 69         code:aedecod-CONJUNCTIVITIS
## 70         code:aedecod-CONSTIPATION
## 71         code:aedecod-CONTUSION
## 72         code:aedecod-COORDINATION_ABNORMAL
## 73         code:aedecod-COUGH
## 74         code:aedecod-CYST
## 75         code:aedecod-CYSTITIS
## 76         code:aedecod-CYSTOSCOPY
## 77         code:aedecod-DECREASED_APPETITE
## 78         code:aedecod-DEHYDRATION
## 79         code:aedecod-DELIRIUM
## 80         code:aedecod-DELUSION
## 81         code:aedecod-DEPRESSED_MOOD
## 82         code:aedecod-DERMATITIS_ATOPIC
## 83         code:aedecod-DERMATITIS_CONTACT
## 84         code:aedecod-DIABETES_MELLITUS
## 85         code:aedecod-DIARRHOEA
## 86         code:aedecod-DISORIENTATION
## 87         code:aedecod-DIZZINESS
## 88         code:aedecod-DRUG_ERUPTION
## 89         code:aedecod-DYSPEPSIA
## 90         code:aedecod-DYSPHAGIA
## 91         code:aedecod-DYSPHONIA
## 92         code:aedecod-DYSPNOEA
## 93         code:aedecod-DYSURIA
## 94         code:aedecod-EAR_INFECTION

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## 95	code: aedecod-EAR_PAIN
## 96	code: aedecod-ELECTROCARDIOGRAM_ST_SEGMENT_DEPRESSION
## 97	code: aedecod-ELECTROCARDIOGRAM_T_WAVE_AMPLITUDE_DECREASED
## 98	code: aedecod-ELECTROCARDIOGRAM_T_WAVE_INVERSION
## 99	code: aedecod-EMPHYSEMA
## 100	code: aedecod-ENURESIS
## 101	code: aedecod-EPISTAXIS
## 102	code: aedecod-ERYTHEMA
## 103	code: aedecod-EXCORIATION
## 104	code: aedecod-EYE_ALLERGY
## 105	code: aedecod-EYE_LASER_SURGERY
## 106	code: aedecod-EYE_PRURITUS
## 107	code: aedecod-EYE_SWELLING
## 108	code: aedecod-FACIAL_BONES_FRACTURE
## 109	code: aedecod-FALL
## 110	code: aedecod-FATIGUE
## 111	code: aedecod-FEELING_ABNORMAL
## 112	code: aedecod-FEELING_COLD
## 113	code: aedecod-FLANK_PAIN
## 114	code: aedecod-FLATULENCE
## 115	code: aedecod-FOOD_CRAVING
## 116	code: aedecod-GASTROENTERITIS_VIRAL
## 117	code: aedecod-GASTROINTESTINAL_HAEMORRHAGE
## 118	code: aedecod-GASTROOESOPHAGEAL_REFLUX_DISEASE
## 119	code: aedecod-GLAUCOMA
## 120	code: aedecod-GLOSSITIS
## 121	code: aedecod-HAEMOPTYSIS
## 122	code: aedecod-HALLUCINATION
## 123	code: aedecod-HALLUCINATION_VISUAL
## 124	code: aedecod-HEADACHE
## 125	code: aedecod-HEART_RATE_INCREASED
## 126	code: aedecod-HEART_RATE_IRREGULAR
## 127	code: aedecod-HEMIANOPIA_HOMONYMOUS
## 128	code: aedecod-HIATUS_HERNIA
## 129	code: aedecod-HIP_FRACTURE
## 130	code: aedecod-HORDEOLUM
## 131	code: aedecod-HOT_FLUSH
## 132	code: aedecod-HYPERBILIRUBINAEMIA
## 133	code: aedecod-HYPERCHOLESTEROLAEMIA
## 134	code: aedecod-HYPERHIDROSIS
## 135	code: aedecod-HYPERSENSITIVITY
## 136	code: aedecod-HYPERSOMNIA
## 137	code: aedecod-HYPERTENSION
## 138	code: aedecod-HYPONATRAEMIA
## 139	code: aedecod-HYPOTENSION
## 140	code: aedecod-INCONTINENCE
## 141	code: aedecod-INCREASED_APPETITE
## 142	code: aedecod-INFLAMMATION
## 143	code: aedecod-INFLUENZA
## 144	code: aedecod-INSOMNIA
## 145	code: aedecod-IRRITABILITY
## 146	code: aedecod-JOINT_DISLOCATION
## 147	code: aedecod-LETHARGY
## 148	code: aedecod-LIBIDO_DECREASED

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## 149                                code:aedecod-LISTLESS
## 150                                code:aedecod-LOCALISED_INFECTION
## 151                                code:aedecod-LOWER_RESPIRATORY_TRACT_INFECTION
## 152                                code:aedecod-MALaise
## 153                                code:aedecod-MALIGNANT_FIBROUS_HISTIOCYTOMA
## 154                                code:aedecod-MICTURITION_URGENCY
## 155                                code:aedecod-MUSCLE_SPASMS
## 156                                code:aedecod-MUSCULAR_WEAKNESS
## 157                                code:aedecod-MYALGIA
## 158                                code:aedecod-MYOCARDIAL_INFARCTION
## 159                                code:aedecod-NASAL_CONGESTION
## 160                                code:aedecod-NASAL_MUCOSA_BIOPSY
## 161                                code:aedecod-NASOPHARYNGITIS
## 162                                code:aedecod-NAUSEA
## 163                                code:aedecod-NEPHROLITHIASIS
## 164                                code:aedecod-NEUTROPHIL_COUNT_INCREASED
## 165                                code:aedecod-NIGHTMARE
## 166                                code:aedecod-OEDEMA
## 167                                code:aedecod-OEDEMA_PERIPHERAL
## 168                                code:aedecod-ONYCHOMYCOSIS
## 169                                code:aedecod-ORTHOSTATIC_HYPOTENSION
## 170                                code:aedecod-PAIN
## 171                                code:aedecod-PAIN_IN_EXTREMITY
## 172                                code:aedecod-PALPITATIONS
## 173                                code:aedecod-PARAESTHESIA
## 174                                code:aedecod-PARAESTHESIA_ORAL
## 175                                code:aedecod-PARKINSON_S_DISEASE
## 176                                code:aedecod-PAROSMIA
## 177                                code:aedecod-PARTIAL_SEIZURES_WITH_SECONDARY_GENERALISATION
## 178                                code:aedecod-PELVIC_PAIN
## 179                                code:aedecod-PHARYNGEAL_ERYTHEMA
## 180                                code:aedecod-PHARYNGOLARYNGEAL_PAIN
## 181                                code:aedecod-PNEUMONIA
## 182                                code:aedecod-POLLAKIURIA
## 183                                code:aedecod-POSTNASAL_DRIP
## 184                                code:aedecod-PRODUCTIVE_COUGH
## 185                                code:aedecod-PROSTATE_CANCER
## 186                                code:aedecod-PRURITUS
## 187                                code:aedecod-PRURITUS_GENERALISED
## 188                                code:aedecod-PSYCHOMOTOR_HYPERACTIVITY
## 189                                code:aedecod-PYREXIA
## 190                                code:aedecod-RALES
## 191                                code:aedecod-RASH
## 192                                code:aedecod-RASH_ERYTHEMATOUS
## 193                                code:aedecod-RASH_MACULO-PAPULAR
## 194                                code:aedecod-RASH_PAPULAR
## 195                                code:aedecod-RASH_PRURITIC
## 196                                code:aedecod-RECTAL_HAEMORRHAGE
## 197                                code:aedecod-RESPIRATORY_TRACT_CONGESTION
## 198                                code:aedecod-RESTLESSNESS
## 199                                code:aedecod-RHINITIS
## 200                                code:aedecod-RHINORRHOEA
## 201                                code:aedecod-SALIVARY_HYPERSECRETION
## 202                                code:aedecod-SEASONAL_ALLERGY

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## 203 code:aedecod-SECRETION_DISCHARGE
## 204 code:aedecod-SHOULDER_PAIN
## 205 code:aedecod-SINUS_ARRHYTHMIA
## 206 code:aedecod-SINUS_BRADYCARDIA
## 207 code:aedecod-SKIN_EXFOLIATION
## 208 code:aedecod-SKIN_IRRITATION
## 209 code:aedecod-SKIN_LACERATION
## 210 code:aedecod-SKIN_LESION_EXCISION
## 211 code:aedecod-SKIN_ODOUR_ABNORMAL
## 212 code:aedecod-SKIN_ULCER
## 213 code:aedecod-SOMNOLENCE
## 214 code:aedecod-STOMACH_DISCOMFORT
## 215 code:aedecod-STUPOR
## 216 code:aedecod-SUDDEN_DEATH
## 217 code:aedecod-SUPRAVENTRICULAR_EXTRASYSTOLES
## 218 code:aedecod-SUPRAVENTRICULAR_TACHYCARDIA
## 219 code:aedecod-SWELLING
## 220 code:aedecod-SYNCOPE
## 221 code:aedecod-SYNCOPE_VASOVAGAL
## 222 code:aedecod-TACHYCARDIA
## 223 code:aedecod-TINNITUS
## 224 code:aedecod-TRANSIENT_ISCHAEMIC_ATTACK
## 225 code:aedecod-ULCER
## 226 code:aedecod-UPPER_RESPIRATORY_TRACT_INFECTION
## 227 code:aedecod-URINARY_TRACT_INFECTION
## 228 code:aedecod-URINE_ANALYSIS_ABNORMAL
## 229 code:aedecod-URTICARIA
## 230 code:aedecod-VAGINAL_MYCOSIS
## 231 code:aedecod-VENTRICULAR_EXTRASYSTOLES
## 232 code:aedecod-VENTRICULAR_HYPERTROPHY
## 233 code:aedecod-VENTRICULAR_SEPTAL_DEFECT
## 234 code:aedecod-VERTIGO
## 235 code:aedecod-VIRAL_INFECTION
## 236 code:aedecod-VISION_BLURRED
## 237 code:aedecod-VOMITING
## 238 code:aedecod-WEIGHT_DECREASED
## 239 code:aedecod-WHITE_BLOOD_CELL_COUNT_INCREASED
## 240 code:aedecod-WOLFF-PARKINSON-WHITE_SYNDROME
## 241 code:aedecod-WOUND
## 242 code:aedecod-WOUND_HAEMORRHAGE
## 243 code:aedecod-_ALL_
## 244 code:aedecod-_NONMISS_
## 245 code:aesoc-CARDIAC_DISORDERS
## 246 code:aesoc-CONGENITAL__FAMILIAL_AND_GENETIC_DISORDERS
## 247 code:aesoc-EAR_AND_LABYRINTH_DISORDERS
## 248 code:aesoc-EYE_DISORDERS
## 249 code:aesoc-GASTROINTESTINAL_DISORDERS
## 250 code:aesoc-GENERAL_DISORDERS_AND_ADMINISTRATION_SITE_CONDITIONS
## 251 code:aesoc-HEPATOBIILIARY_DISORDERS
## 252 code:aesoc-IMMUNE_SYSTEM_DISORDERS
## 253 code:aesoc-INFECTIONS_AND_INFESTATIONS
## 254 code:aesoc-INJURY__POISONING_AND_PROCEDURAL_COMPLICATIONS
## 255 code:aesoc-INVESTIGATIONS
## 256 code:aesoc-METABOLISM_AND_NUTRITION_DISORDERS

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## 257             code:aesoc-MUSCULOSKELETAL_AND_CONNECTIVE_TISSUE_DISORDERS
## 258 code:aesoc-NEOPLASMS_BENIGN__MALIGNANT_AND_UNSPECIFIED__INCL_CYSTS_AND_POLYPS_
## 259             code:aesoc-NERVOUS_SYSTEM_DISORDERS
## 260             code:aesoc-PSYCHIATRIC_DISORDERS
## 261             code:aesoc-RENAL_AND_URINARY_DISORDERS
## 262             code:aesoc-REPRODUCTIVE_SYSTEM_AND_BREAST_DISORDERS
## 263             code:aesoc-RESPIRATORY__THORACIC_AND_MEDIASTINAL_DISORDERS
## 264             code:aesoc-SKIN_AND_SUBCUTANEOUS_TISSUE_DISORDERS
## 265             code:aesoc-SOCIAL_CIRCUMSTANCES
## 266             code:aesoc-SURGICAL_AND_MEDICAL_PROCEDURES
## 267             code:aesoc-VASCULAR_DISORDERS
## 268             code:aesoc-_ALL_
## 269             code:aesoc-_NONMISS_
## 270             code:factor-USUBJID
## 271             code:factor-_ALL_
## 272             code:factor-_NONMISS_
## 273             code:factor-quantity
## 274             code:procedure-count
## 275             code:procedure-countdistinct
## 276             code:saffl-Y
## 277             code:saffl-_ALL_
## 278             code:saffl-_NONMISS_
## 279             code:trta-Placebo
## 280             code:trta-Xanomeline_High_Dose
## 281             code:trta-Xanomeline_Low_Dose
## 282             code:trta-_ALL_
## 283             code:trta-_NONMISS_
##             clprefLabel
## 1             ABDOMINAL DISCOMFORT
## 2             ABDOMINAL PAIN
## 3             ACROCHORDON EXCISION
## 4             ACTINIC KERATOSIS
## 5             AGITATION
## 6             ALCOHOL USE
## 7             ALLERGIC GRANULOMATOUS ANGIITIS
## 8             ALOPECIA
## 9             AMNESIA
## 10            ANXIETY
## 11            APPLICATION SITE BLEEDING
## 12            APPLICATION SITE DERMATITIS
## 13            APPLICATION SITE DESQUAMATION
## 14            APPLICATION SITE DISCHARGE
## 15            APPLICATION SITE DISCOLOURATION
## 16            APPLICATION SITE ERYTHEMA
## 17            APPLICATION SITE INDURATION
## 18            APPLICATION SITE IRRITATION
## 19            APPLICATION SITE PAIN
## 20            APPLICATION SITE PERSPIRATION
## 21            APPLICATION SITE PRURITUS
## 22            APPLICATION SITE REACTION
## 23            APPLICATION SITE SWELLING
## 24            APPLICATION SITE URTICARIA
## 25            APPLICATION SITE VESICLES
## 26            APPLICATION SITE WARMTH

```

## 27	ARTHRALGIA
## 28	ARTHRITIS
## 29	ASTHENIA
## 30	ATRIAL FIBRILLATION
## 31	ATRIAL FLUTTER
## 32	ATRIAL HYPERTROPHY
## 33	ATRIOVENTRICULAR BLOCK FIRST DEGREE
## 34	ATRIOVENTRICULAR BLOCK SECOND DEGREE
## 35	BACK PAIN
## 36	BALANCE DISORDER
## 37	BENIGN PROSTATIC HYPERPLASIA
## 38	BIOPSY
## 39	BIOPSY PROSTATE
## 40	BLISTER
## 41	BLOOD ALKALINE PHOSPHATASE INCREASED
## 42	BLOOD CHOLESTEROL INCREASED
## 43	BLOOD CREATINE PHOSPHOKINASE INCREASED
## 44	BLOOD GLUCOSE INCREASED
## 45	BLOOD URINE PRESENT
## 46	BODY TEMPERATURE INCREASED
## 47	BRADYCARDIA
## 48	BRONCHITIS
## 49	BUNDLE BRANCH BLOCK LEFT
## 50	BUNDLE BRANCH BLOCK RIGHT
## 51	BURNING SENSATION
## 52	CALCULUS URETHRAL
## 53	CARDIAC DISORDER
## 54	CARDIAC FAILURE CONGESTIVE
## 55	CATARACT OPERATION
## 56	CELLULITIS
## 57	CERUMEN IMPACTION
## 58	CERVICITIS
## 59	CHEST DISCOMFORT
## 60	CHEST PAIN
## 61	CHILLS
## 62	COGNITIVE DISORDER
## 63	COLD SWEAT
## 64	COLON CANCER
## 65	COMPLETED SUICIDE
## 66	COMPLEX PARTIAL SEIZURES
## 67	CONFUSIONAL STATE
## 68	CONJUNCTIVAL HAEMORRHAGE
## 69	CONJUNCTIVITIS
## 70	CONSTIPATION
## 71	CONTUSION
## 72	COORDINATION ABNORMAL
## 73	COUGH
## 74	CYST
## 75	CYSTITIS
## 76	CYSTOSCOPY
## 77	DECREASED APPETITE
## 78	DEHYDRATION
## 79	DELIRIUM
## 80	DELUSION

## 81	DEPRESSED MOOD
## 82	DERMATITIS ATOPIC
## 83	DERMATITIS CONTACT
## 84	DIABETES MELLITUS
## 85	DIARRHOEA
## 86	DISORIENTATION
## 87	DIZZINESS
## 88	DRUG ERUPTION
## 89	DYSPEPSIA
## 90	DYSPHAGIA
## 91	DYSPHONIA
## 92	DYSPNOEA
## 93	DYSURIA
## 94	EAR INFECTION
## 95	EAR PAIN
## 96	ELECTROCARDIOGRAM ST SEGMENT DEPRESSION
## 97	ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED
## 98	ELECTROCARDIOGRAM T WAVE INVERSION
## 99	EMPHYSEMA
## 100	ENURESIS
## 101	EPISTAXIS
## 102	ERYTHEMA
## 103	EXCORIATION
## 104	EYE ALLERGY
## 105	EYE LASER SURGERY
## 106	EYE PRURITUS
## 107	EYE SWELLING
## 108	FACIAL BONES FRACTURE
## 109	FALL
## 110	FATIGUE
## 111	FEELING ABNORMAL
## 112	FEELING COLD
## 113	FLANK PAIN
## 114	FLATULENCE
## 115	FOOD CRAVING
## 116	GASTROENTERITIS VIRAL
## 117	GASTROINTESTINAL HAEMORRHAGE
## 118	GASTROOESOPHAGEAL REFLUX DISEASE
## 119	GLAUCOMA
## 120	GLOSSITIS
## 121	HAEMOPTYSIS
## 122	HALLUCINATION
## 123	HALLUCINATION, VISUAL
## 124	HEADACHE
## 125	HEART RATE INCREASED
## 126	HEART RATE IRREGULAR
## 127	HEMIANOPIA HOMONYMOUS
## 128	HIATUS HERNIA
## 129	HIP FRACTURE
## 130	HORDEOLUM
## 131	HOT FLUSH
## 132	HYPERBILIRUBINAEMIA
## 133	HYPERCHOLESTEROLAEMIA
## 134	HYPERHIDROSIS

## 135	HYPERSENSITIVITY
## 136	HYPERSOMNIA
## 137	HYPERTENSION
## 138	HYPONATRAEMIA
## 139	HYPOTENSION
## 140	INCONTINENCE
## 141	INCREASED APPETITE
## 142	INFLAMMATION
## 143	INFLUENZA
## 144	INSOMNIA
## 145	IRRITABILITY
## 146	JOINT DISLOCATION
## 147	LETHARGY
## 148	LIBIDO DECREASED
## 149	LISTLESS
## 150	LOCALISED INFECTION
## 151	LOWER RESPIRATORY TRACT INFECTION
## 152	MALaise
## 153	MALIGNANT FIBROUS HISTIOCYTOMA
## 154	MICTURITION URGENCY
## 155	MUSCLE SPASMS
## 156	MUSCULAR WEAKNESS
## 157	MYALGIA
## 158	MYOCARDIAL INFARCTION
## 159	NASAL CONGESTION
## 160	NASAL MUCOSA BIOPSY
## 161	NASOPHARYNGITIS
## 162	NAUSEA
## 163	NEPHROLITHIASIS
## 164	NEUTROPHIL COUNT INCREASED
## 165	NIGHTMARE
## 166	OEDEMA
## 167	OEDEMA PERIPHERAL
## 168	ONYCHOMYCOSIS
## 169	ORTHOSTATIC HYPOTENSION
## 170	PAIN
## 171	PAIN IN EXTREMITY
## 172	PALPITATIONS
## 173	PARAESTHESIA
## 174	PARAESTHESIA ORAL
## 175	PARKINSON'S DISEASE
## 176	PAROSMIA
## 177	PARTIAL SEIZURES WITH SECONDARY GENERALISATION
## 178	PELVIC PAIN
## 179	PHARYNGEAL ERYTHEMA
## 180	PHARYNGOLARYNGEAL PAIN
## 181	PNEUMONIA
## 182	POLLAKIURIA
## 183	POSTNASAL DRIP
## 184	PRODUCTIVE COUGH
## 185	PROSTATE CANCER
## 186	PRURITUS
## 187	PRURITUS GENERALISED
## 188	PSYCHOMOTOR HYPERACTIVITY



## 189	PYREXIA
## 190	RALES
## 191	RASH
## 192	RASH ERYTHEMATOUS
## 193	RASH MACULO-PAPULAR
## 194	RASH PAPULAR
## 195	RASH PRURITIC
## 196	RECTAL HAEMORRHAGE
## 197	RESPIRATORY TRACT CONGESTION
## 198	RESTLESSNESS
## 199	RHINITIS
## 200	RHINORRHOEA
## 201	SALIVARY HYPERSECRETION
## 202	SEASONAL ALLERGY
## 203	SECRETION DISCHARGE
## 204	SHOULDER PAIN
## 205	SINUS ARRHYTHMIA
## 206	SINUS BRADYCARDIA
## 207	SKIN EXFOLIATION
## 208	SKIN IRRITATION
## 209	SKIN LACERATION
## 210	SKIN LESION EXCISION
## 211	SKIN ODOUR ABNORMAL
## 212	SKIN ULCER
## 213	SOMNOLENCE
## 214	STOMACH DISCOMFORT
## 215	STUPOR
## 216	SUDDEN DEATH
## 217	SUPRAVENTRICULAR EXTRASYSTOLES
## 218	SUPRAVENTRICULAR TACHYCARDIA
## 219	SWELLING
## 220	SYNCOPE
## 221	SYNCOPE VASOVAGAL
## 222	TACHYCARDIA
## 223	TINNITUS
## 224	TRANSIENT ISCHAEMIC ATTACK
## 225	ULCER
## 226	UPPER RESPIRATORY TRACT INFECTION
## 227	URINARY TRACT INFECTION
## 228	URINE ANALYSIS ABNORMAL
## 229	URTICARIA
## 230	VAGINAL MYCOSIS
## 231	VENTRICULAR EXTRASYSTOLES
## 232	VENTRICULAR HYPERTROPHY
## 233	VENTRICULAR SEPTAL DEFECT
## 234	VERTIGO
## 235	VIRAL INFECTION
## 236	VISION BLURRED
## 237	VOMITING
## 238	WEIGHT DECREASED
## 239	WHITE BLOOD CELL COUNT INCREASED
## 240	WOLFF-PARKINSON-WHITE SYNDROME
## 241	WOUND
## 242	WOUND HAEMORRHAGE

```

## 243                                     _ALL_
## 244                                     _NONMISS_
## 245                                CARDIAC DISORDERS
## 246                        CONGENITAL, FAMILIAL AND GENETIC DISORDERS
## 247                                EAR AND LABYRINTH DISORDERS
## 248                                EYE DISORDERS
## 249                                GASTROINTESTINAL DISORDERS
## 250                GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS
## 251                                HEPATOBILIARY DISORDERS
## 252                                IMMUNE SYSTEM DISORDERS
## 253                                INFECTIONS AND INFESTATIONS
## 254                        INJURY, POISONING AND PROCEDURAL COMPLICATIONS
## 255                                INVESTIGATIONS
## 256                                METABOLISM AND NUTRITION DISORDERS
## 257                MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS
## 258    NEOPLASMS BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS)
## 259                                NERVOUS SYSTEM DISORDERS
## 260                                PSYCHIATRIC DISORDERS
## 261                                RENAL AND URINARY DISORDERS
## 262                REPRODUCTIVE SYSTEM AND BREAST DISORDERS
## 263                RESPIRATORY, THORACIC AND MEDIASTINAL DISORDERS
## 264                SKIN AND SUBCUTANEOUS TISSUE DISORDERS
## 265                                SOCIAL CIRCUMSTANCES
## 266                                SURGICAL AND MEDICAL PROCEDURES
## 267                                VASCULAR DISORDERS
## 268                                     _ALL_
## 269                                     _NONMISS_
## 270                                     USUBJID
## 271                                     _ALL_
## 272                                     _NONMISS_
## 273                                     quantity
## 274                                     count
## 275                                countdistinct
## 276                                     Y
## 277                                     _ALL_
## 278                                     _NONMISS_
## 279                                     Placebo
## 280                                Xanomeline High Dose
## 281                                Xanomeline Low Dose
## 282                                     _ALL_
## 283                                     _NONMISS_
##          vn  vct                                vnop
## 1    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 2    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 3    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 4    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 5    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 6    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 7    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 8    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 9    aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 10   aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 11   aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 12   aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD

```

[illegible]

[illegible]

[illegible]

[illegible]

```

## 229  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 230  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 231  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 232  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 233  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 234  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 235  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 236  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 237  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 238  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 239  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 240  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 241  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 242  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 243  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 244  aedecod DATA rrdfqbcrnd0:ADAE_AEDECOD
## 245    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 246    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 247    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 248    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 249    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 250    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 251    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 252    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 253    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 254    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 255    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 256    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 257    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 258    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 259    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 260    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 261    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 262    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 263    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 264    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 265    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 266    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 267    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 268    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 269    aesoc DATA rrdfqbcrnd0:ADAE_AESOC
## 270    factor DATA ==
## 271    factor DATA <NA>
## 272    factor DATA <NA>
## 273    factor DATA ==
## 274 procedure DATA ==
## 275 procedure DATA ==
## 276    saffl DATA rrdfqbcrnd0:ADAE_SAFFL
## 277    saffl DATA rrdfqbcrnd0:ADAE_SAFFL
## 278    saffl DATA rrdfqbcrnd0:ADAE_SAFFL
## 279    trta DATA rrdfqbcrnd0:ADAE_TRTA
## 280    trta DATA rrdfqbcrnd0:ADAE_TRTA
## 281    trta DATA rrdfqbcrnd0:ADAE_TRTA
## 282    trta DATA rrdfqbcrnd0:ADAE_TRTA

```

```

## 283      trta DATA      rrdfqbcrnd0:ADAE_TRTA
##
## 1          ABDOMINAL DISCOMFORT
## 2          ABDOMINAL PAIN
## 3          ACROCHORDON EXCISION
## 4          ACTINIC KERATOSIS
## 5          AGITATION
## 6          ALCOHOL USE
## 7          ALLERGIC GRANULOMATOUS ANGIITIS
## 8          ALOPECIA
## 9          AMNESIA
## 10         ANXIETY
## 11         APPLICATION SITE BLEEDING
## 12         APPLICATION SITE DERMATITIS
## 13         APPLICATION SITE DESQUAMATION
## 14         APPLICATION SITE DISCHARGE
## 15         APPLICATION SITE DISCOLOURATION
## 16         APPLICATION SITE ERYTHEMA
## 17         APPLICATION SITE INDURATION
## 18         APPLICATION SITE IRRITATION
## 19         APPLICATION SITE PAIN
## 20         APPLICATION SITE PERSPIRATION
## 21         APPLICATION SITE PRURITUS
## 22         APPLICATION SITE REACTION
## 23         APPLICATION SITE SWELLING
## 24         APPLICATION SITE URTICARIA
## 25         APPLICATION SITE VESICLES
## 26         APPLICATION SITE WARMTH
## 27         ARTHRALGIA
## 28         ARTHRITIS
## 29         ASTHENIA
## 30         ATRIAL FIBRILLATION
## 31         ATRIAL FLUTTER
## 32         ATRIAL HYPERTROPHY
## 33         ATRIOVENTRICULAR BLOCK FIRST DEGREE
## 34         ATRIOVENTRICULAR BLOCK SECOND DEGREE
## 35         BACK PAIN
## 36         BALANCE DISORDER
## 37         BENIGN PROSTATIC HYPERPLASIA
## 38         BIOPSY
## 39         BIOPSY PROSTATE
## 40         BLISTER
## 41         BLOOD ALKALINE PHOSPHATASE INCREASED
## 42         BLOOD CHOLESTEROL INCREASED
## 43         BLOOD CREATINE PHOSPHOKINASE INCREASED
## 44         BLOOD GLUCOSE INCREASED
## 45         BLOOD URINE PRESENT
## 46         BODY TEMPERATURE INCREASED
## 47         BRADYCARDIA
## 48         BRONCHITIS
## 49         BUNDLE BRANCH BLOCK LEFT
## 50         BUNDLE BRANCH BLOCK RIGHT
## 51         BURNING SENSATION
## 52         CALCULUS URETHRAL

```



## 53	CARDIAC DISORDER
## 54	CARDIAC FAILURE CONGESTIVE
## 55	CATARACT OPERATION
## 56	CELLULITIS
## 57	CERUMEN IMPACTION
## 58	CERVICITIS
## 59	CHEST DISCOMFORT
## 60	CHEST PAIN
## 61	CHILLS
## 62	COGNITIVE DISORDER
## 63	COLD SWEAT
## 64	COLON CANCER
## 65	COMPLETED SUICIDE
## 66	COMPLEX PARTIAL SEIZURES
## 67	CONFUSIONAL STATE
## 68	CONJUNCTIVAL HAEMORRHAGE
## 69	CONJUNCTIVITIS
## 70	CONSTIPATION
## 71	CONTUSION
## 72	COORDINATION ABNORMAL
## 73	COUGH
## 74	CYST
## 75	CYSTITIS
## 76	CYSTOSCOPY
## 77	DECREASED APPETITE
## 78	DEHYDRATION
## 79	DELIRIUM
## 80	DELUSION
## 81	DEPRESSED MOOD
## 82	DERMATITIS ATOPIC
## 83	DERMATITIS CONTACT
## 84	DIABETES MELLITUS
## 85	DIARRHOEA
## 86	DISORIENTATION
## 87	DIZZINESS
## 88	DRUG ERUPTION
## 89	DYSPEPSIA
## 90	DYSPHAGIA
## 91	DYSPHONIA
## 92	DYSPNOEA
## 93	DYSURIA
## 94	EAR INFECTION
## 95	EAR PAIN
## 96	ELECTROCARDIOGRAM ST SEGMENT DEPRESSION
## 97	ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED
## 98	ELECTROCARDIOGRAM T WAVE INVERSION
## 99	EMPHYSEMA
## 100	ENURESIS
## 101	EPISTAXIS
## 102	ERYTHEMA
## 103	EXCORIATION
## 104	EYE ALLERGY
## 105	EYE LASER SURGERY
## 106	EYE PRURITUS

## 107	EYE SWELLING
## 108	FACIAL BONES FRACTURE
## 109	FALL
## 110	FATIGUE
## 111	FEELING ABNORMAL
## 112	FEELING COLD
## 113	FLANK PAIN
## 114	FLATULENCE
## 115	FOOD CRAVING
## 116	GASTROENTERITIS VIRAL
## 117	GASTROINTESTINAL HAEMORRHAGE
## 118	GASTROOESOPHAGEAL REFLUX DISEASE
## 119	GLAUCOMA
## 120	GLOSSITIS
## 121	HAEMOPTYSIS
## 122	HALLUCINATION
## 123	HALLUCINATION, VISUAL
## 124	HEADACHE
## 125	HEART RATE INCREASED
## 126	HEART RATE IRREGULAR
## 127	HEMIANOPIA HOMONYMOUS
## 128	HIATUS HERNIA
## 129	HIP FRACTURE
## 130	HORDEOLUM
## 131	HOT FLUSH
## 132	HYPERBILIRUBINAEMIA
## 133	HYPERCHOLESTEROLAEMIA
## 134	HYPERHIDROSIS
## 135	HYPERSENSITIVITY
## 136	HYPERSONNIA
## 137	HYPERTENSION
## 138	HYPONATRAEMIA
## 139	HYPOTENSION
## 140	INCONTINENCE
## 141	INCREASED APPETITE
## 142	INFLAMMATION
## 143	INFLUENZA
## 144	INSOMNIA
## 145	IRRITABILITY
## 146	JOINT DISLOCATION
## 147	LETHARGY
## 148	LIBIDO DECREASED
## 149	LISTLESS
## 150	LOCALISED INFECTION
## 151	LOWER RESPIRATORY TRACT INFECTION
## 152	MALADISE
## 153	MALIGNANT FIBROUS HISTIOCYTOMA
## 154	MICTURITION URGENCY
## 155	MUSCLE SPASMS
## 156	MUSCULAR WEAKNESS
## 157	MYALGIA
## 158	MYOCARDIAL INFARCTION
## 159	NASAL CONGESTION
## 160	NASAL MUCOSA BIOPSY

## 161	NASOPHARYNGITIS
## 162	NAUSEA
## 163	NEPHROLITHIASIS
## 164	NEUTROPHIL COUNT INCREASED
## 165	NIGHTMARE
## 166	OEDEMA
## 167	OEDEMA PERIPHERAL
## 168	ONYCHOMYCOSIS
## 169	ORTHOSTATIC HYPOTENSION
## 170	PAIN
## 171	PAIN IN EXTREMITY
## 172	PALPITATIONS
## 173	PARAESTHESIA
## 174	PARAESTHESIA ORAL
## 175	PARKINSON'S DISEASE
## 176	PAROSMIA
## 177	PARTIAL SEIZURES WITH SECONDARY GENERALISATION
## 178	PELVIC PAIN
## 179	PHARYNGEAL ERYTHEMA
## 180	PHARYNGOLARYNGEAL PAIN
## 181	PNEUMONIA
## 182	POLLAKIURIA
## 183	POSTNASAL DRIP
## 184	PRODUCTIVE COUGH
## 185	PROSTATE CANCER
## 186	PRURITUS
## 187	PRURITUS GENERALISED
## 188	PSYCHOMOTOR HYPERACTIVITY
## 189	PYREXIA
## 190	RALES
## 191	RASH
## 192	RASH ERYTHEMATOUS
## 193	RASH MACULO-PAPULAR
## 194	RASH PAPULAR
## 195	RASH PRURITIC
## 196	RECTAL HAEMORRHAGE
## 197	RESPIRATORY TRACT CONGESTION
## 198	RESTLESSNESS
## 199	RHINITIS
## 200	RHINORRHOEA
## 201	SALIVARY HYPERSECRETION
## 202	SEASONAL ALLERGY
## 203	SECRETION DISCHARGE
## 204	SHOULDER PAIN
## 205	SINUS ARRHYTHMIA
## 206	SINUS BRADYCARDIA
## 207	SKIN EXFOLIATION
## 208	SKIN IRRITATION
## 209	SKIN LACERATION
## 210	SKIN LESION EXCISION
## 211	SKIN ODOUR ABNORMAL
## 212	SKIN ULCER
## 213	SOMNOLENCE
## 214	STOMACH DISCOMFORT

##	215	STUPOR
##	216	SUDDEN DEATH
##	217	SUPRAVENTRICULAR EXTRASYSTOLES
##	218	SUPRAVENTRICULAR TACHYCARDIA
##	219	SWELLING
##	220	SYNCOPE
##	221	SYNCOPE VASOVAGAL
##	222	TACHYCARDIA
##	223	TINNITUS
##	224	TRANSIENT ISCHAEMIC ATTACK
##	225	ULCER
##	226	UPPER RESPIRATORY TRACT INFECTION
##	227	URINARY TRACT INFECTION
##	228	URINE ANALYSIS ABNORMAL
##	229	URTICARIA
##	230	VAGINAL MYCOSIS
##	231	VENTRICULAR EXTRASYSTOLES
##	232	VENTRICULAR HYPERTROPHY
##	233	VENTRICULAR SEPTAL DEFECT
##	234	VERTIGO
##	235	VIRAL INFECTION
##	236	VISION BLURRED
##	237	VOMITING
##	238	WEIGHT DECREASED
##	239	WHITE BLOOD CELL COUNT INCREASED
##	240	WOLFF-PARKINSON-WHITE SYNDROME
##	241	WOUND
##	242	WOUND HAEMORRHAGE
##	243	<NA>
##	244	<NA>
##	245	CARDIAC DISORDERS
##	246	CONGENITAL, FAMILIAL AND GENETIC DISORDERS
##	247	EAR AND LABYRINTH DISORDERS
##	248	EYE DISORDERS
##	249	GASTROINTESTINAL DISORDERS
##	250	GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS
##	251	HEPATOBIILIARY DISORDERS
##	252	IMMUNE SYSTEM DISORDERS
##	253	INFECTIONS AND INFESTATIONS
##	254	INJURY, POISONING AND PROCEDURAL COMPLICATIONS
##	255	INVESTIGATIONS
##	256	METABOLISM AND NUTRITION DISORDERS
##	257	MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS
##	258	NEOPLASMS BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS)
##	259	NERVOUS SYSTEM DISORDERS
##	260	PSYCHIATRIC DISORDERS
##	261	RENAL AND URINARY DISORDERS
##	262	REPRODUCTIVE SYSTEM AND BREAST DISORDERS
##	263	RESPIRATORY, THORACIC AND MEDIASTINAL DISORDERS
##	264	SKIN AND SUBCUTANEOUS TISSUE DISORDERS
##	265	SOCIAL CIRCUMSTANCES
##	266	SURGICAL AND MEDICAL PROCEDURES
##	267	VASCULAR DISORDERS
##	268	<NA>

```

## 269                                     <NA>
## 270                                     USUBJID
## 271                                     <NA>
## 272                                     <NA>
## 273                                     quantity
## 274                                     count
## 275                                     countdistinct
## 276                                     Y
## 277                                     <NA>
## 278                                     <NA>
## 279                                     Placebo
## 280                                     Xanomeline High Dose
## 281                                     Xanomeline Low Dose
## 282                                     <NA>
## 283                                     <NA>
## crnd-dimension:aedecod crnd-dimension:aedecod crnd-dimension:aedecod crnd-dimension:aedecod crnd-dim

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## HYPERTROPHY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## HYPERTROPHY -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK FIRST DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK FIRST DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK SECOND DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK SECOND DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRADYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRADYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK LEFT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK LEFT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK RIGHT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK RIGHT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## FAILURE CONGESTIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## FAILURE CONGESTIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## ARRHYTHMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## ARRHYTHMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TACHYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TACHYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## HYPERTROPHY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## HYPERTROPHY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## CONJUNCTIVITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## CONJUNCTIVITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE ALLERGY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE ALLERGY
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE PRURITUS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE PRURITUS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE SWELLING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE SWELLING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLAUCOMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLAUCOMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONSTIPATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONSTIPATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLATULENCE
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLATULENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROESOPHAGEAL REFLUX DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROESOPHAGEAL REFLUX DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLOSSITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLOSSITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIATUS
## HERNIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIATUS
## HERNIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE INDURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE INDURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE REACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE REACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ASTHENIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ASTHENIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PYREXIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PYREXIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBIILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBIILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBIILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERBILIRUBINAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBIILIARY DISORDERS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERBILIRUBINAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRONCHITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRONCHITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERVICITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERVICITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROENTERITIS VIRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROENTERITIS VIRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOCALISED
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOCALISED
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINARY
## TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINARY
## TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VAGINAL
## MYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VAGINAL
## MYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIP FRACTURE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIP FRACTURE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## LACERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## LACERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## ALKALINE PHOSPHATASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## ALKALINE PHOSPHATASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CREATINE PHOSPHOKINASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CREATINE PHOSPHOKINASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD URINE
## PRESENT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD URINE
## PRESENT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTOSCOPY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTOSCOPY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM ST SEGMENT DEPRESSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM ST SEGMENT DEPRESSION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

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## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## IRREGULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## IRREGULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

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## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DECREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DECREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEHYDRATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEHYDRATION
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIABETES
## MELLITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIABETES
## MELLITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FOOD CRAVING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FOOD CRAVING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPONATRAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



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## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPONATRAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN IN
## EXTREMITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN IN
## EXTREMITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SHOULDER
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SHOULDER
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARKINSON'S
## DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARKINSON'S
## DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHOMOTOR
## HYPERACTIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHOMOTOR
## HYPERACTIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ANXIETY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ANXIETY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLETED
## SUICIDE -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLETED
## SUICIDE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## DISORIENTATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## DISORIENTATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INSOMNIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INSOMNIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IRRITABILITY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IRRITABILITY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSURIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSURIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NEPHROLITHIASIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NEPHROLITHIASIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POLLAKIURIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POLLAKIURIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BENIGN
## PROSTATIC HYPERPLASIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BENIGN
## PROSTATIC HYPERPLASIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PELVIC PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PELVIC PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EMPHYSEMA --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EMPHYSEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HAEMOPTYSIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HAEMOPTYSIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POSTNASAL
## DRIP -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POSTNASAL
## DRIP -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RALES --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RALES --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALOPECIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALOPECIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLD SWEAT
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLD SWEAT
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## ATOPIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## ATOPIC -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DRUG
## ERUPTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DRUG
## ERUPTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ULCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ULCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CATARACT
## OPERATION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CATARACT
## OPERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE LASER
## SURGERY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE LASER
## SURGERY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPOTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPOTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ORTHOSTATIC
## HYPOTENSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ORTHOSTATIC
## HYPOTENSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FLUTTER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FLUTTER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK SECOND DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK SECOND DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## SEPTAL DEFECT -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## SEPTAL DEFECT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VERTIGO --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VERTIGO --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VISION
## BLURRED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VISION
## BLURRED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SALIVARY
## HYPERSECRETION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SALIVARY
## HYPERSECRETION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STOMACH
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STOMACH
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PERSPIRATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PERSPIRATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE REACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE REACTION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE SWELLING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE SWELLING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE URTICARIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE URTICARIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ASTHENIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ASTHENIA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING COLD
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING COLD
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALAISE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALAISE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PYREXIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PYREXIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SEASONAL
## ALLERGY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SEASONAL
## ALLERGY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTITIS --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HORDEOLUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HORDEOLUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOWER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOWER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINARY
## TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINARY
## TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FACIAL BONES
## FRACTURE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FACIAL BONES
## FRACTURE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIP FRACTURE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIP FRACTURE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY
## PROSTATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY
## PROSTATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CHOLESTEROL INCREASED -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CHOLESTEROL INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## GLUCOSE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## GLUCOSE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHT
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHT
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DECREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DECREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERCHOLESTEROLAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERCHOLESTEROLAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLANK PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLANK PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCLE
## SPASMS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCLE
## SPASMS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYALGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYALGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PROSTATE
## CANCER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PROSTATE
## CANCER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMNESIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMNESIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BURNING
## SENSATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BURNING
## SENSATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COGNITIVE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COGNITIVE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERSOMNIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERSOMNIA
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LETHARGY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LETHARGY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAROSMIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAROSMIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARTIAL
## SEIZURES WITH SECONDARY GENERALISATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARTIAL
## SEIZURES WITH SECONDARY GENERALISATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE
## VASOVAGAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE
## VASOVAGAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TRANSIENT
## ISCHAEMIC ATTACK -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TRANSIENT
## ISCHAEMIC ATTACK -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELIRIUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELIRIUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEPRESSED
## MOOD -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEPRESSED
## MOOD -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION, VISUAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION, VISUAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INSOMNIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INSOMNIA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LIBIDO
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LIBIDO
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LISTLESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LISTLESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NIGHTMARE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NIGHTMARE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CALCULUS
## URETHRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CALCULUS
## URETHRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NEPHROLITHIASIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NEPHROLITHIASIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BENIGN
## PROSTATIC HYPERPLASIA -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BENIGN
## PROSTATIC HYPERPLASIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALLERGIC
## GRANULOMATOUS ANGIITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALLERGIC
## GRANULOMATOUS ANGIITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EPISTAXIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EPISTAXIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PHARYNGEAL
## ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PHARYNGEAL
## ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## PHARYNGOLARYNGEAL PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## PHARYNGOLARYNGEAL PAIN -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRODUCTIVE
## COUGH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRODUCTIVE
## COUGH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY
## TRACT CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY
## TRACT CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINORRHOEA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINORRHOEA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACTINIC
## KERATOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACTINIC
## KERATOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLISTER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLISTER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS
## GENERALISED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS
## GENERALISED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH MACULO-
## PAPULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH MACULO-
## PAPULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH PAPULAR
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH PAPULAR
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## PRURITIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## PRURITIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ODOUR
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ODOUR
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URTICARIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URTICARIA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALCOHOL USE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALCOHOL USE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACROCHORDON
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACROCHORDON
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN LESION
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN LESION
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FLUTTER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FLUTTER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK FIRST DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK FIRST DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK RIGHT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK RIGHT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PALPITATIONS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PALPITATIONS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR TACHYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR TACHYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOLFF-
## PARKINSON-WHITE SYNDROME -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOLFF-
## PARKINSON-WHITE SYNDROME -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## SEPTAL DEFECT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## SEPTAL DEFECT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERUMEN
## IMPACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERUMEN
## IMPACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TINNITUS --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TINNITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VERTIGO --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VERTIGO --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONJUNCTIVAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONJUNCTIVAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VISION
## BLURRED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VISION
## BLURRED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHAGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHAGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RECTAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RECTAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE BLEEDING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE BLEEDING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DESQUAMATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DESQUAMATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCOLOURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCOLOURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE SWELLING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE SWELLING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE URTICARIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE URTICARIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE WARMTH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE WARMTH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYST --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYST --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLAMMATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLAMMATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALAISE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALAISE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SECRETION
## DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SECRETION
## DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SUDDEN DEATH
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SUDDEN DEATH
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SWELLING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SWELLING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ULCER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ULCER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERSENSITIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERSENSITIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CELLULITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CELLULITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOCALISED
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOCALISED
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ONYCHOMYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ONYCHOMYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PNEUMONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PNEUMONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VIRAL
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VIRAL
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): JOINT
## DISLOCATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): JOINT
## DISLOCATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## LACERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## LACERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## GLUCOSE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## GLUCOSE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BODY
## TEMPERATURE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BODY
## TEMPERATURE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM ST SEGMENT DEPRESSION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM ST SEGMENT DEPRESSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL MUCOSA
## BIOPSY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL MUCOSA
## BIOPSY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEUTROPHIL
## COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEUTROPHIL
## COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINE
## ANALYSIS ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINE
## ANALYSIS ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE BLOOD
## CELL COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE BLOOD
## CELL COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FOOD CRAVING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FOOD CRAVING
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCLE
## SPASMS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCLE
## SPASMS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCULAR
## WEAKNESS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCULAR
## WEAKNESS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SHOULDER
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SHOULDER
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLON CANCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLON CANCER
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALIGNANT
## FIBROUS HISTIOCYTOMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALIGNANT
## FIBROUS HISTIOCYTOMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BALANCE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BALANCE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLEX
## PARTIAL SEIZURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLEX
## PARTIAL SEIZURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COORDINATION
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COORDINATION
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEMIANOPIA
## HOMONYMOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEMIANOPIA
## HOMONYMOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LETHARGY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LETHARGY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## ORAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## ORAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STUPOR --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STUPOR --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TRANSIENT
## ISCHAEMIC ATTACK -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TRANSIENT
## ISCHAEMIC ATTACK -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ANXIETY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ANXIETY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEPRESSED
## MOOD -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEPRESSED
## MOOD -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IRRITABILITY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IRRITABILITY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESTLESSNESS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESTLESSNESS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSURIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSURIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ENURESIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ENURESIS --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCONTINENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCONTINENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EPISTAXIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EPISTAXIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## PHARYNGOLARYNGEAL PAIN -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## PHARYNGOLARYNGEAL PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINORRHOEA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINORRHOEA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLISTER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLISTER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## CONTACT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## CONTACT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS
## GENERALISED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS
## GENERALISED -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## ERYTHEMATOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## ERYTHEMATOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## PRURITIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## PRURITIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## EXFOLIATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## EXFOLIATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URTICARIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URTICARIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CATARACT
## OPERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CATARACT
## OPERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HOT FLUSH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HOT FLUSH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPOTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPOTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FIBRILLATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FLUTTER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## FLUTTER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## HYPERTROPHY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ATRIAL
## HYPERTROPHY -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK FIRST DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK FIRST DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK SECOND DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ATRIOVENTRICULAR BLOCK SECOND DEGREE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRADYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRADYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK LEFT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK LEFT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK RIGHT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BUNDLE
## BRANCH BLOCK RIGHT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## FAILURE CONGESTIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## FAILURE CONGESTIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYOCARDIAL
## INFARCTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PALPITATIONS
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PALPITATIONS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## ARRHYTHMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## ARRHYTHMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SINUS
## BRADYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR TACHYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## SUPRAVENTRICULAR TACHYCARDIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TACHYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TACHYCARDIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## EXTRASYSTOLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## HYPERTROPHY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## HYPERTROPHY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOLFF-
## PARKINSON-WHITE SYNDROME -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CARDIAC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOLFF-
## PARKINSON-WHITE SYNDROME -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## SEPTAL DEFECT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONGENITAL,
## FAMILIAL AND GENETIC DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VENTRICULAR
## SEPTAL DEFECT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERUMEN
## IMPACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERUMEN
## IMPACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TINNITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TINNITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VERTIGO --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR AND
## LABYRINTH DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VERTIGO --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONJUNCTIVAL
## HAEMORRHAGE -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONJUNCTIVAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## CONJUNCTIVITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## CONJUNCTIVITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE ALLERGY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE ALLERGY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE PRURITUS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE PRURITUS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE SWELLING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE SWELLING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLAUCOMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLAUCOMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VISION
## BLURRED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VISION
## BLURRED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## DISCOMFORT -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ABDOMINAL
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONSTIPATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONSTIPATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIARRHOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPEPSIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHAGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHAGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLATULENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLATULENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROOESOPHAGEAL REFLUX DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROOESOPHAGEAL REFLUX DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLOSSITIS --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GLOSSITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIATUS
## HERNIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIATUS
## HERNIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NAUSEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RECTAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RECTAL
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SALIVARY
## HYPERSECRETION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SALIVARY
## HYPERSECRETION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STOMACH
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STOMACH
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROINTESTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VOMITING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE BLEEDING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE BLEEDING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DERMATITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DESQUAMATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DESQUAMATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCOLOURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE DISCOLOURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE INDURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE INDURATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PERSPIRATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PERSPIRATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE PRURITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE REACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE REACTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE SWELLING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE SWELLING -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE URTICARIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE URTICARIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE VESICLES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE WARMTH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): APPLICATION
## SITE WARMTH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ASTHENIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ASTHENIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST
## DISCOMFORT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHEST PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CHILLS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYST --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYST --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FATIGUE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING COLD
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FEELING COLD
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLAMMATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLAMMATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALAISE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALAISE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): OEDEMA
## PERIPHERAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PYREXIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PYREXIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SECRETION
## DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SECRETION
## DISCHARGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SUDDEN DEATH
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SUDDEN DEATH
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SWELLING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SWELLING --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ULCER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): GENERAL
## DISORDERS AND ADMINISTRATION SITE CONDITIONS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ULCER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERBILIRUBINAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HEPATOBILIARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERBILIRUBINAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERSENSITIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERSENSITIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SEASONAL
## ALLERGY -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IMMUNE
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SEASONAL
## ALLERGY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRONCHITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BRONCHITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CELLULITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CELLULITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERVICITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CERVICITIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EAR
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROENTERITIS VIRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## GASTROENTERITIS VIRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HORDEOLUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HORDEOLUM --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFLUENZA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOCALISED
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOCALISED
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOWER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LOWER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NASOPHARYNGITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ONYCHOMYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ONYCHOMYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PNEUMONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PNEUMONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): UPPER
## RESPIRATORY TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINARY
## TRACT INFECTION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINARY
## TRACT INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VAGINAL
## MYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VAGINAL
## MYCOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VIRAL
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INFECTIONS
## AND INFESTATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VIRAL
## INFECTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONTUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EXCORIATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FACIAL BONES
## FRACTURE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FACIAL BONES
## FRACTURE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FALL --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIP FRACTURE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HIP FRACTURE
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): JOINT
## DISLOCATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): JOINT
## DISLOCATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## LACERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## LACERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INJURY,
## POISONING AND PROCEDURAL COMPLICATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY
## PROSTATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BIOPSY
## PROSTATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## ALKALINE PHOSPHATASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## ALKALINE PHOSPHATASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CHOLESTEROL INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CHOLESTEROL INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CREATINE PHOSPHOKINASE INCREASED -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## CREATINE PHOSPHOKINASE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## GLUCOSE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD
## GLUCOSE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD URINE
## PRESENT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLOOD URINE
## PRESENT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BODY
## TEMPERATURE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BODY
## TEMPERATURE INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTOSCOPY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CYSTOSCOPY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM ST SEGMENT DEPRESSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM ST SEGMENT DEPRESSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE AMPLITUDE DECREASED -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## ELECTROCARDIOGRAM T WAVE INVERSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## IRREGULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEART RATE
## IRREGULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL MUCOSA
## BIOPSY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL MUCOSA
## BIOPSY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEUTROPHIL
## COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEUTROPHIL
## COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINE
## ANALYSIS ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URINE
## ANALYSIS ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHT
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WEIGHT
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE BLOOD
## CELL COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## INVESTIGATIONS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE BLOOD
## CELL COUNT INCREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DECREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DECREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEHYDRATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEHYDRATION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIABETES
## MELLITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIABETES
## MELLITUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FOOD CRAVING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FOOD CRAVING
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERCHOLESTEROLAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERCHOLESTEROLAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPONATRAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPONATRAEMIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): METABOLISM
## AND NUTRITION DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCREASED
## APPETITE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRALGIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ARTHRITIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BACK PAIN --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLANK PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): FLANK PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCLE
## SPASMS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCLE
## SPASMS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCULAR
## WEAKNESS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MUSCULAR
## WEAKNESS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYALGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MYALGIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN IN
## EXTREMITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAIN IN
## EXTREMITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SHOULDER
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS -- CODING ERROR- no decode
## value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SHOULDER
## PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLON CANCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLON CANCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALIGNANT
## FIBROUS HISTIOCYTOMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MALIGNANT
## FIBROUS HISTIOCYTOMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PROSTATE
## CANCER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NEOPLASMS
## BENIGN, MALIGNANT AND UNSPECIFIED (INCL CYSTS AND POLYPS) -- CODING ERROR-
## no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PROSTATE
## CANCER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMNESIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMNESIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BALANCE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BALANCE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BURNING
## SENSATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BURNING
## SENSATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COGNITIVE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COGNITIVE
## DISORDER -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLEX
## PARTIAL SEIZURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLEX
## PARTIAL SEIZURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COORDINATION
## ABNORMAL -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COORDINATION
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DIZZINESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEADACHE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEMIANOPIA
## HOMONYMOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HEMIANOPIA
## HOMONYMOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERSOMNIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERSOMNIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LETHARGY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LETHARGY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## ORAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARAESTHESIA
## ORAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARKINSON'S
## DISEASE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARKINSON'S
## DISEASE -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAROSMIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PAROSMIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARTIAL
## SEIZURES WITH SECONDARY GENERALISATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PARTIAL
## SEIZURES WITH SECONDARY GENERALISATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHOMOTOR
## HYPERACTIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHOMOTOR
## HYPERACTIVITY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOMNOLENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STUPOR --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): STUPOR --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE
## VASOVAGAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SYNCOPE
## VASOVAGAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TRANSIENT
## ISCHAEMIC ATTACK -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NERVOUS
## SYSTEM DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): TRANSIENT
## ISCHAEMIC ATTACK -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AGITATION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ANXIETY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ANXIETY --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLETED
## SUICIDE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COMPLETED
## SUICIDE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CONFUSIONAL
## STATE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELIRIUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELIRIUM --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DELUSION --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEPRESSED
## MOOD -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DEPRESSED
## MOOD -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## DISORIENTATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## DISORIENTATION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION, VISUAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HALLUCINATION, VISUAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INSOMNIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INSOMNIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IRRITABILITY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): IRRITABILITY
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LIBIDO
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LIBIDO
## DECREASED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LISTLESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): LISTLESS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NIGHTMARE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NIGHTMARE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESTLESSNESS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PSYCHIATRIC
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESTLESSNESS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CALCULUS
## URETHRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CALCULUS
## URETHRAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSURIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSURIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ENURESIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ENURESIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCONTINENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): INCONTINENCE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): MICTURITION
## URGENCY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NEPHROLITHIASIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## NEPHROLITHIASIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POLLAKIURIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RENAL AND
## URINARY DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POLLAKIURIA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BENIGN
## PROSTATIC HYPERPLASIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BENIGN
## PROSTATIC HYPERPLASIA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PELVIC PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): REPRODUCTIVE
## SYSTEM AND BREAST DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PELVIC PAIN
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALLERGIC
## GRANULOMATOUS ANGIITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALLERGIC
## GRANULOMATOUS ANGIITIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COUGH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPHONIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DYSPNOEA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EMPHYSEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EMPHYSEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EPISTAXIS --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EPISTAXIS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HAEMOPTYSIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HAEMOPTYSIS
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NASAL
## CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PHARYNGEAL
## ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PHARYNGEAL
## ERYTHEMA -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## PHARYNGOLARYNGEAL PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## PHARYNGOLARYNGEAL PAIN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POSTNASAL
## DRIP -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): POSTNASAL
## DRIP -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRODUCTIVE
## COUGH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRODUCTIVE
## COUGH -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RALES --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RALES --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY
## TRACT CONGESTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY
## TRACT CONGESTION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINORRHOEA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RESPIRATORY,
## THORACIC AND MEDIASTINAL DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RHINORRHOEA
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACTINIC
## KERATOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACTINIC
## KERATOSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALOPECIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALOPECIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLISTER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLISTER --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLD SWEAT
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): COLD SWEAT
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## ATOPIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## ATOPIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## CONTACT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DERMATITIS
## CONTACT -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DRUG
## ERUPTION -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): DRUG
## ERUPTION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

```



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## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ERYTHEMA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## HYPERHIDROSIS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS
## GENERALISED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): PRURITUS
## GENERALISED -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## ERYTHEMATOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## ERYTHEMATOUS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH MACULO-
## PAPULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH MACULO-
## PAPULAR -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH PAPULAR
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH PAPULAR
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## PRURITIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): RASH
## PRURITIC -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## EXFOLIATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## EXFOLIATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN
## IRRITATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ODOUR
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ODOUR
## ABNORMAL -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ULCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN ULCER
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URTICARIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN AND
## SUBCUTANEOUS TISSUE DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): URTICARIA --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALCOHOL USE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SOCIAL
## CIRCUMSTANCES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ALCOHOL USE
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACROCHORDON
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ACROCHORDON
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CATARACT
## OPERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): CATARACT
## OPERATION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE LASER
## SURGERY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): EYE LASER
## SURGERY -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN LESION
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SURGICAL AND
## MEDICAL PROCEDURES -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): SKIN LESION
## EXCISION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HOT FLUSH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HOT FLUSH --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPERTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPOTENSION
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HYPOTENSION
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ORTHOSTATIC
## HYPOTENSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): ORTHOSTATIC
## HYPOTENSION -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

```

```
## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Y -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): VASCULAR
## DISORDERS -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WOUND
## HAEMORRHAGE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]):
## countdistinct -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): USUBJID --
## CODING ERROR- no decode value
```

```
cat("AE cube stored as ", ae.cube.fn, "\n")
```

```
## AE cube stored as /tmp/Rtmpz1JmRR/DC-AE-R-V-0-5-2.ttl
```

```
targetFile<- file.path(targetDir,"DC-AE-sample.ttl")
if (file.copy( ae.cube.fn, targetFile, overwrite=TRUE)) {
  cat("AE cube copied to ", normalizePath(targetFile), "\n")
}
```

```
## AE cube copied to /home/ma/projects/rrdfqbcrrnd0/rrdfqbcrrndex/inst/extdata/sample-rdf/DC-AE-sample.t
```

## RDF data cube in csv files

```
demoObsDataCsvFn<- system.file("extdata/sample-cfg", "demo.AR.csv", package="rrdfqbcrrndex")
demoObsData <- read.csv(demoObsDataCsvFn,stringsAsFactors=FALSE)

##TODO add measurefmt; quick hack - affects vignettes/cube-from-workbook.Rmd and
##TODO inst/data-raw/create-qb-examples-as-ttl.Rmd
if (!( "measurefmt" %in% names(demoObsData))) {
  demoObsData$measurefmt<- "%6.1f"
  demoObsData$measurefmt[ demoObsData$procedure %in% c("n", "nmiss", "count") ]<- "%6.0f"
```

```
## sprintf( demoObsData$measurefmt, demoObsData$measure)
}

demoMetaDataCsvFn<- system.file("extdata/sample-cfg", "DEMO-Components.csv", package="rrdfqbcindex")
demoMetaData <- read.csv(demoMetaDataCsvFn,stringsAsFactors=FALSE)

demo.cube.fn<- BuildCubeFromDataFrames(demoMetaData, demoObsData )
```

```
## [1] "prefix rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>\nprefix skos: <http://www.w3.org/2004/02/22-testsuite-rdf-syntax-ns#>"
##      DataStructureDefinition      dimension
## 1      ds:dsd-DEMO      crnd-dimension:agegr1
## 2      ds:dsd-DEMO      crnd-dimension:agegr1
## 3      ds:dsd-DEMO      crnd-dimension:agegr1
## 4      ds:dsd-DEMO      crnd-dimension:agegr1
## 5      ds:dsd-DEMO      crnd-dimension:agegr1
## 6      ds:dsd-DEMO      crnd-dimension:ethnic
## 7      ds:dsd-DEMO      crnd-dimension:ethnic
## 8      ds:dsd-DEMO      crnd-dimension:ethnic
## 9      ds:dsd-DEMO      crnd-dimension:ethnic
## 10     ds:dsd-DEMO      crnd-dimension:factor
## 11     ds:dsd-DEMO      crnd-dimension:factor
## 12     ds:dsd-DEMO      crnd-dimension:factor
## 13     ds:dsd-DEMO      crnd-dimension:factor
## 14     ds:dsd-DEMO      crnd-dimension:factor
## 15     ds:dsd-DEMO      crnd-dimension:factor
## 16     ds:dsd-DEMO      crnd-dimension:procedure
## 17     ds:dsd-DEMO      crnd-dimension:procedure
## 18     ds:dsd-DEMO      crnd-dimension:procedure
## 19     ds:dsd-DEMO      crnd-dimension:procedure
## 20     ds:dsd-DEMO      crnd-dimension:procedure
## 21     ds:dsd-DEMO      crnd-dimension:procedure
## 22     ds:dsd-DEMO      crnd-dimension:procedure
## 23     ds:dsd-DEMO      crnd-dimension:procedure
## 24     ds:dsd-DEMO      crnd-dimension:procedure
## 25     ds:dsd-DEMO      crnd-dimension:procedure
## 26     ds:dsd-DEMO      crnd-dimension:race
## 27     ds:dsd-DEMO      crnd-dimension:race
## 28     ds:dsd-DEMO      crnd-dimension:race
## 29     ds:dsd-DEMO      crnd-dimension:race
## 30     ds:dsd-DEMO      crnd-dimension:race
## 31     ds:dsd-DEMO      crnd-dimension:race
## 32     ds:dsd-DEMO      crnd-dimension:race
## 33     ds:dsd-DEMO      crnd-dimension:sex
## 34     ds:dsd-DEMO      crnd-dimension:sex
## 35     ds:dsd-DEMO      crnd-dimension:sex
## 36     ds:dsd-DEMO      crnd-dimension:sex
## 37     ds:dsd-DEMO      crnd-dimension:sex
## 38     ds:dsd-DEMO      crnd-dimension:sex
## 39     ds:dsd-DEMO      crnd-dimension:trt01a
## 40     ds:dsd-DEMO      crnd-dimension:trt01a
## 41     ds:dsd-DEMO      crnd-dimension:trt01a
## 42     ds:dsd-DEMO      crnd-dimension:trt01a
## 43     ds:dsd-DEMO      crnd-dimension:trt01a
```

```

##          cprefLabel
## 1   Codelist scheme: agegr1
## 2   Codelist scheme: agegr1
## 3   Codelist scheme: agegr1
## 4   Codelist scheme: agegr1
## 5   Codelist scheme: agegr1
## 6   Codelist scheme: ethnic
## 7   Codelist scheme: ethnic
## 8   Codelist scheme: ethnic
## 9   Codelist scheme: ethnic
## 10  Codelist scheme: factor
## 11  Codelist scheme: factor
## 12  Codelist scheme: factor
## 13  Codelist scheme: factor
## 14  Codelist scheme: factor
## 15  Codelist scheme: factor
## 16  Codelist scheme: procedure
## 17  Codelist scheme: procedure
## 18  Codelist scheme: procedure
## 19  Codelist scheme: procedure
## 20  Codelist scheme: procedure
## 21  Codelist scheme: procedure
## 22  Codelist scheme: procedure
## 23  Codelist scheme: procedure
## 24  Codelist scheme: procedure
## 25  Codelist scheme: procedure
## 26   Codelist scheme: race
## 27   Codelist scheme: race
## 28   Codelist scheme: race
## 29   Codelist scheme: race
## 30   Codelist scheme: race
## 31   Codelist scheme: race
## 32   Codelist scheme: race
## 33   Codelist scheme: sex
## 34   Codelist scheme: sex
## 35   Codelist scheme: sex
## 36   Codelist scheme: sex
## 37   Codelist scheme: sex
## 38   Codelist scheme: sex
## 39  Codelist scheme: trt01a
## 40  Codelist scheme: trt01a
## 41  Codelist scheme: trt01a
## 42  Codelist scheme: trt01a
## 43  Codelist scheme: trt01a
##
##                                     cl
## 1   code:agegr1-65-80
## 2   code:agegr1-_65
## 3   code:agegr1-_80
## 4   code:agegr1-_ALL_
## 5   code:agegr1-_NONMISS_
## 6   code:ethnic-HISPANIC_OR_LATINO
## 7   code:ethnic-NOT_HISPANIC_OR_LATINO
## 8   code:ethnic-_ALL_
## 9   code:ethnic-_NONMISS_

```

```

## 10             code:factor-_ALL_
## 11             code:factor-_NONMISS_
## 12             code:factor-age
## 13             code:factor-proportion
## 14             code:factor-quantity
## 15             code:factor-weightbl
## 16             code:procedure-count
## 17             code:procedure-max
## 18             code:procedure-mean
## 19             code:procedure-median
## 20             code:procedure-min
## 21             code:procedure-n
## 22             code:procedure-percent
## 23             code:procedure-q1
## 24             code:procedure-q3
## 25             code:procedure-std
## 26             code:race-AMERICAN_INDIAN_OR_ALASKA_NATIVE
## 27             code:race-ASIAN
## 28             code:race-BLACK_OR_AFRICAN_AMERICAN
## 29             code:race-NATIVE_HAWAIIAN_OR_OTHER_PACIFIC_ISLANDER
## 30             code:race-WHITE
## 31             code:race-_ALL_
## 32             code:race-_NONMISS_
## 33             code:sex-F
## 34             code:sex-M
## 35             code:sex-U
## 36             code:sex-UN
## 37             code:sex-_ALL_
## 38             code:sex-_NONMISS_
## 39             code:trt01a-Placebo
## 40             code:trt01a-Xanomeline_High_Dose
## 41             code:trt01a-Xanomeline_Low_Dose
## 42             code:trt01a-_ALL_
## 43             code:trt01a-_NONMISS_
##             clprefLabel      vn  vct
## 1             65-80      agegr1 DATA
## 2             <65      agegr1 DATA
## 3             >80      agegr1 DATA
## 4             _ALL_      agegr1 DATA
## 5             _NONMISS_      agegr1 DATA
## 6             HISPANIC OR LATINO      ethnic DATA
## 7             NOT HISPANIC OR LATINO      ethnic DATA
## 8             _ALL_      ethnic DATA
## 9             _NONMISS_      ethnic DATA
## 10            _ALL_      factor DATA
## 11            _NONMISS_      factor DATA
## 12            age      factor DATA
## 13            proportion      factor DATA
## 14            quantity      factor DATA
## 15            weightbl      factor DATA
## 16            count procedure DATA
## 17            max procedure DATA
## 18            mean procedure DATA
## 19            median procedure DATA

```

```

## 20                                min procedure DATA
## 21                                n procedure DATA
## 22                                percent procedure DATA
## 23                                q1 procedure DATA
## 24                                q3 procedure DATA
## 25                                std procedure DATA
## 26                AMERICAN INDIAN OR ALASKA NATIVE        race SDTM
## 27                                ASIAN                    race SDTM
## 28                                BLACK OR AFRICAN AMERICAN    race SDTM
## 29    NATIVE HAWAIIAN OR OTHER PACIFIC ISLANDER        race SDTM
## 30                                WHITE                    race SDTM
## 31                                _ALL_                    race SDTM
## 32                                _NONMISS_                race SDTM
## 33                                F                        sex SDTM
## 34                                M                        sex SDTM
## 35                                U                        sex SDTM
## 36                                UN                      sex SDTM
## 37                                _ALL_                    sex SDTM
## 38                                _NONMISS_                sex SDTM
## 39                                Placebo                  trt01a DATA
## 40                                Xanomeline High Dose      trt01a DATA
## 41                                Xanomeline Low Dose       trt01a DATA
## 42                                _ALL_                    trt01a DATA
## 43                                _NONMISS_                trt01a DATA
##                                vnop                                vnval
## 1  rrdfqbcrnd0:ADSL_AGEGR1                                65-80
## 2  rrdfqbcrnd0:ADSL_AGEGR1                                <65
## 3  rrdfqbcrnd0:ADSL_AGEGR1                                >80
## 4  rrdfqbcrnd0:ADSL_AGEGR1                                <NA>
## 5  rrdfqbcrnd0:ADSL_AGEGR1                                <NA>
## 6  rrdfqbcrnd0:ADSL_ETHNIC                                HISPANIC OR LATINO
## 7  rrdfqbcrnd0:ADSL_ETHNIC                                NOT HISPANIC OR LATINO
## 8  rrdfqbcrnd0:ADSL_ETHNIC                                <NA>
## 9  rrdfqbcrnd0:ADSL_ETHNIC                                <NA>
## 10                                <NA>                                <NA>
## 11                                <NA>                                <NA>
## 12                                ==                                age
## 13                                ==                                proportion
## 14                                ==                                quantity
## 15                                ==                                weightbl
## 16                                ==                                count
## 17                                ==                                max
## 18                                ==                                mean
## 19                                ==                                median
## 20                                ==                                min
## 21                                ==                                n
## 22                                ==                                percent
## 23                                ==                                q1
## 24                                ==                                q3
## 25                                ==                                std
## 26  rrdfqbcrnd0:ADSL_RACE                AMERICAN INDIAN OR ALASKA NATIVE
## 27  rrdfqbcrnd0:ADSL_RACE                ASIAN
## 28  rrdfqbcrnd0:ADSL_RACE                BLACK OR AFRICAN AMERICAN
## 29  rrdfqbcrnd0:ADSL_RACE    NATIVE HAWAIIAN OR OTHER PACIFIC ISLANDER

```

```

## 30   rrdfqbcrnd0:ADSL_RACE                WHITE
## 31   rrdfqbcrnd0:ADSL_RACE                <NA>
## 32   rrdfqbcrnd0:ADSL_RACE                <NA>
## 33   rrdfqbcrnd0:ADSL_SEX                 F
## 34   rrdfqbcrnd0:ADSL_SEX                 M
## 35   rrdfqbcrnd0:ADSL_SEX                 U
## 36   rrdfqbcrnd0:ADSL_SEX                 UN
## 37   rrdfqbcrnd0:ADSL_SEX                 <NA>
## 38   rrdfqbcrnd0:ADSL_SEX                 <NA>
## 39   rrdfqbcrnd0:ADSL_TRT01A              Placebo
## 40   rrdfqbcrnd0:ADSL_TRT01A              Xanomeline High Dose
## 41   rrdfqbcrnd0:ADSL_TRT01A              Xanomeline Low Dose
## 42   rrdfqbcrnd0:ADSL_TRT01A              <NA>
## 43   rrdfqbcrnd0:ADSL_TRT01A              <NA>
## crnd-dimension:agegr1 crnd-dimension:agegr1 crnd-dimension:agegr1 crnd-dimension:agegr1 crnd-dimension:agegr1

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): F -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): M -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): std --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): std --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): std --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): n -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): n -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): n -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q3 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q3 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q3 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q1 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q1 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q1 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): age --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): <65 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): <65 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): <65 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): <65 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): <65 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): <65 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): 65-80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): 65-80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): 65-80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): 65-80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): 65-80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): 65-80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): >80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): >80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): >80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): >80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): >80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): >80 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): WHITE --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): BLACK OR
## AFRICAN AMERICAN -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): AMERICAN
## INDIAN OR ALASKA NATIVE -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): std --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): std --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): std --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): n -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): n -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): n -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): median --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): mean --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q3 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q3 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q3 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q1 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q1 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightb1 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): q1 -- CODING
## ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightb1 --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): max --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): min --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): weightbl --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _NONMISS_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NOT HISPANIC
## OR LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

```



```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NOT HISPANIC
## OR LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NOT HISPANIC
## OR LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NOT HISPANIC
## OR LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NOT HISPANIC
## OR LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): NOT HISPANIC
## OR LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HISPANIC OR
## LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HISPANIC OR
## LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

```

```

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HISPANIC OR
## LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): count --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): quantity --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Placebo --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HISPANIC OR
## LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## Low Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HISPANIC OR
## LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

```

```
## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): Xanomeline
## High Dose -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): HISPANIC OR
## LATINO -- CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): percent --
## CODING ERROR- no decode value

## Warning in ph.recode(obsData[i, qbdim], recode.list[[qbdim]]): proportion
## -- CODING ERROR- no decode value
```

```
cat("DEMO cube stored as ", normalizePath(demo.cube.fn), "\n")
```

```
## DEMO cube stored as /tmp/Rtmpz1JmRR/DC-DEMO-R-V-0-5-2.ttl
```

```
targetFile<- file.path(targetDir,"DC-DEMO-sample.ttl")

if (file.copy( demo.cube.fn, targetFile, overwrite=TRUE)) {
  cat("DEMO cube copied to ", normalizePath(targetFile), "\n")
}
```

```
## DEMO cube copied to /home/ma/projects/rrdfqbc rnd0/rrdfqbc rndex/inst/extdata/sample-rdf/DC-DEMO-samp
```