

# Create DEMO table as csv file

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*2016-04-29*

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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" )
```

```
## Warning in 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/Rtmpw3UFRD/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" )
```

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

```
## Warning in 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

```



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## 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]]):
## 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]]): Placebo --
## CODING ERROR- no decode value

## Warning in 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]]): Placebo --
## CODING ERROR- no decode value

## Warning in 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]]):
## 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]]): Placebo --
## CODING ERROR- no decode value

## Warning in 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

## Warning in 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

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

## Warning in 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

## 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]]):
## 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]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in 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]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in 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]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in 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]]): _ALL_ --
## CODING ERROR- no decode value

## Warning in 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

## 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/Rtmpw3UFRD/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/rrdfqbc rnd0/rrdfqbc rndex/inst/extdata/sample-rdf/DC-AE-sample.t
```

## RDF data cube in csv files



```

demoObsDataCsvFn<- system.file("extdata/sample-cfg", "demo.AR.csv", package="rrdfqbcindex")
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 )

```

```

## Warning in 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]]): 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]]): 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]]): 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/Rtmpw3UFrd/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
```