yspec: an R package to create and deploy data set specification objects in a modeling and simulation workflow

R/Pharma 2020

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2020-10-14





Motivation

Data set documentation is ...

- 1. really important
- 2. takes a lot of work and time

Need to create: define.pdf (analysis data sets)

Want to track:

- column name and order
- data source
- data type
- scope
 - range for continuous data
 - levels for discrete data + decode
- other attributes convenience



Motivation (2)

- programmatic handling of data attributes
 - avoid MS Word
- specify data set attributes once (only once)
 - re-use in different contexts
- easy data entry process
 - human readable format



How yspec works

- Use yaml markup language to declare data column names and attributes
- Read yaml code from file into an object (list) in R
- Validate data contents
 - ▶ enforce standards and requirements (column names ≤ 8 characters)
- Extract column info
 - maximize data reuse
- ► Template define.pdf document
 - options for customizing format for different use contexts



Get started

```
spec <- ys_load("nmdata.yml")</pre>
```

Meta data for nmdata data set

```
SETUP__:

description: "Example PopPK analysis data set"

sponsor: "Tru Valu Pharmaceuticals"

projectnumber: "TVP11020F"

use_internal_db: true

character_last: true

comment col: "C"
```



Column data (same file as meta data)

```
C:
  short: "comment character"
  values: {comment: "C", non-comment: "."}
NUM:
USUBJID:
  short: "unique subject identifier"
 type: "character"
TIME:
 unit: "hour"
SEQ:
  short: "data type"
  values: {observation: 0, dose: 1}
WT:
  range: [40, 140]
```

Access

```
spec <- ys_load("nmdata.yml")
spec$WT</pre>
```

- . name value
- . col WT
- . type numeric
- . short subject weight at baseline
- . unit kg
- . range 40 to 140



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Access (2)

name

value

spec\$RF

```
col RF
type character
short renal function stage
value 0 : normal function
1 : mild impairment
2 : moderate impairment
3 : severe impairment
```

Make factor columns in the data set from values + decode (labels)

data <- ys add factors(data, spec)

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Access (3)

spec is a list that we can work on like any other list

```
spec$TIME$unit
. [1] "hour"
labels <- map(spec, ys_get_label)</pre>
names(spec)
summary(spec)
as.list(spec)
spec[c("WT", "ALB", "ACTARM")]
ys_select(spec, WT, ALB, ACTARM)
```

Create define document

- 1. define.pdf for regulatory submission
- 2. mydataset.pdf for working use

```
ys_document(
   spec,
   title = "Data Definitions",
   author = "Kyle Baron",
   sponsor = "Tru Valu Pharmaceuticals",
   type = "regulatory"
)
```



Data Definitions

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2020-10-05

1 Datasets

	Description	Location
ĺ	Example PopPK analysis data set	nmdata.xpt
	Example AE analysis data set	ae.xpt

1.1 Example PopPK analysis data set (nmdata.xpt)

VARIABLE	LABEL	TYPE	CODES
С	comment character	character	C = comment, . = non-comment
NUM	record number	numeric	
ID	subject identifier	numeric	
TIME	time after first dose (unit: hour)	numeric	
SEQ	data type	numeric	0 = observation, 1 = dose
CMT	compartment number	numeric	
EVID	event ID	numeric	values: 0, 1
AMT	dose amount (unit: mg)	numeric	
DV	dependent variable	numeric	
AGE	age (unit: years)	numeric	
WT	subject weight at baseline (unit: kg)	numeric	



Thank you

- R/Pharma Organizers
- MetrumRG
 - Bill Knebel and Marc Gastonguay
 - M&S scientists at MetrumRG

