# Creating a basic data structure (BDS) Exposure ADaM

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## 2024-12-01

# **Programming workflow**

Get required R packages

Read CDISC pilot SDTM and ADaM datasets	2
EXADJ	2
EXDOSE	2
EXPLDOS	2
Derive numeric datetime, analysis day variables	3
ASTDT	3
AENDT	3
ASTDTM	4
AENDTM	4
ASTDY	5
AENDY	5
EXDURD	5
DOSEO	6
PDOSEO	6
Create 1:1 mapping records	6
PARAMCD	7
AVALC	7
Create Summary Records	8
Create Summary Records	8
Assign PARAMCD, PARAMN, etc. from Reference tables	11
Derive Categorization Variables (AVALCATx)	12
AVALCAT1	12
Assign ASEQ	12
ASEQ	12
Add Labels and Attributes	13
References	12

```
Warning: package 'pharmaversesdtm' was built under R version 4.4.2

Attaching package: 'lubridate'

The following objects are masked from 'package:base':

date, intersect, setdiff, union
```

### Read CDISC pilot SDTM and ADaM datasets

```
adsl_vars <- exprs(TRTSDT, TRTSDTM, TRTEDT, TRTEDTM)

# left join EX and adsl TRTSDT, TRTSDTM, TRTEDT, TRTEDTM on ex.STUDYID=adslSTUDYID and ex.USUB.
adex <- derive_vars_merged(
    dataset=ex
    ,dataset_add = adsl
    ,new_vars = adsl_vars
    ,by_vars = exprs(STUDYID, USUBJID)
    ) # dim(adex) 591 21</pre>
```

The CDISC pilot EX domain data does not contain a dose adjustment flag or the planned dose information. For demonstration purposes, this will be added to the data.

#### **EXADJ**

• Exposure Adjustment?

#### **EXDOSE**

- · exposure dose
- from SDTM.EX.EXDOSE

#### **EXPLDOS**

· Planned Dose

```
adex <- adex %>%
 mutate(
   EXADJ = case_when(
     USUBJID == "01-701-1028" & VISIT %in% c("WEEK 2") ~ "ADVERSE EVENT",
     USUBJID == "01-701-1148" & VISIT %in% c("WEEK 2", "WEEK 24") ~ "MEDICATION ERROR",
     TRUE ~ NA_character_
   ),
   EXDOSE = case when(
     USUBJID == "01-701-1028" & VISIT %in% c("WEEK 2") \sim 0,
     USUBJID == "01-701-1148" & VISIT %in% c("WEEK 2", "WEEK 24") ~ ^{\circ},
     TRUE ~ EXDOSE
   )
 ) %>%
 mutate(EXPLDOS = if_else(EXTRT == "PLACEBO", 0, 54))
adex %>% select(EXTRT, EXPLDOS) %>% distinct()
# A tibble: 2 x 2
 EXTRT EXPLDOS
 <chr>
             <dbl>
1 PLACEBO
                   0
```

## Derive numeric datetime, analysis day variables

54

#### **ASTDT**

2 XANOMELINE

- Analysis Start Date
- Set to a numeric form of EX.EXSTDTC when EX.EXSTDTC consists of a full date.

#### **AENDT**

- · Analysis End Date
- Set to a numeric form of EX.EXENDTC when EX.EXENDTC consists of a full date.

```
# Convert character datetime to numeric datetime
adex <- derive_vars_dt(adex, new_vars_prefix = "AST", dtc = EXSTDTC)
adex <- derive_vars_dt(adex, new_vars_prefix = "AEN", dtc = EXENDTC) # dim(adex) 591 25
adex %>% select(USUBJID, VISIT, EXSTDTC, EXENDTC, ASTDT, AENDT) %>% head()
```

```
# A tibble: 6 x 6
 USUBJID VISIT
                      EXSTDTC
                                 EXENDTC
                                            ASTDT
                                                       AENDT
             <chr>
                      <chr>
  <chr>
                                 <chr>
                                            <date>
                                                       <date>
1 01-701-1015 BASELINE 2014-01-02 2014-01-16 2014-01-02 2014-01-16
2 01-701-1015 WEEK 2 2014-01-17 2014-06-18 2014-01-17 2014-06-18
3 01-701-1015 WEEK 24 2014-06-19 2014-07-02 2014-06-19 2014-07-02
4 01-701-1023 BASELINE 2012-08-05 2012-08-27 2012-08-05 2012-08-27
5 01-701-1023 WEEK 2 2012-08-28 2012-09-01 2012-08-28 2012-09-01
6 01-701-1028 BASELINE 2013-07-19 2013-08-01 2013-07-19 2013-08-01
```

#### **ASTDTM**

#### **AENDTM**

The next examples demonstrates the datetime imputation features available in the derive\_vars\_dtm() function, where the time is imputed as "00:00:00":

```
adex <- derive_vars_dtm(</pre>
  adex
  ,dtc = EXSTDTC
  # Impute dtc date to the first day of the month
  ,highest_imputation = "M"
  ,date_imputation = "first"
  ,new_vars_prefix = "AST"
)
adex <- derive_vars_dtm(</pre>
  adex,
 dtc = EXENDTC,
  # Impute dtc date to the last day of the month
 highest_imputation = "M",
 date_imputation = "last",
 new_vars_prefix = "AEN"
adex %>% select(EXSTDTC,EXENDTC,ASTDTM,AENDTM) %>% head()
```

#### **ASTDY**

- Analysis Start Day
- 'ASTDT-TRTSDT+1'

#### **AENDY**

- Analysis End Day
- 'AENDT-TRTSDT+1'

```
adex <- derive_vars_dy(
  dataset=adex
  ,reference_date = TRTSDT
  ,source_vars = exprs(ASTDT, AENDT)
  ) # dim(adex) 591 33
adex %>% select(TRTSDT, ASTDT, ASTDY, AENDT, AENDY) %>% head()
```

```
# A tibble: 6 x 5
 TRTSDT ASTDT
                  ASTDY AENDT
                               AENDY
         <date>
 <date>
                 <dbl> <date>
                               <dbl>
1 2014-01-02 2014-01-02 1 2014-01-16
                                15
3 2014-01-02 2014-06-19 169 2014-07-02 182
4 2012-08-05 2012-08-05 1 2012-08-27 23
5 2012-08-05 2012-08-28 24 2012-09-01
                                  28
6 2013-07-19 2013-07-19 1 2013-08-01
                                14
```

#### **EXDURD**

- Duration of treatment or exposure
- 'EXDURD=AENDT ASTDT +1'

```
adex <- adex %>%
  derive_vars_duration(
    new_var = EXDURD
    ,start_date = ASTDT
    ,end_date = AENDT
    # duration unit can be "years", "months", "weeks", "days", "hours", "minutes", "seconds"
    ,out_unit = "DAYS")

adex %>% select(ASTDT, AENDT, EXDURD) %>% head()
```

```
# A tibble: 6 x 3
 ASTDT AENDT
                        EXDURD
  <date>
             <date>
                         <dbl>
1 2014-01-02 2014-01-16
                            15
2 2014-01-17 2014-06-18
                           153
3 2014-06-19 2014-07-02
                            14
4 2012-08-05 2012-08-27
                            23
5 2012-08-28 2012-09-01
                             5
6 2013-07-19 2013-08-01
                            14
```

#### **DOSEO**

- Dose Overall (?). Refers to the actual dose of the study drug administered, standardized (e.g., per unit of body weight, such as mg/kg).
- 'EXDOSE \* EXDURD'

#### **PDOSEO**

- Planned Dose Overall (?) Represents the planned dose per protocol, similarly normalized
- 'EXPLDOS \* EXDURD'

```
adex <- adex %>%
  mutate(
    DOSEO = EXDOSE * EXDURD
    ,PDOSEO = EXPLDOS * EXDURD)

adex %>% select(USUBJID, EXDOSE, EXPLDOS, EXDURD, DOSEO, PDOSEO) %>% head()
```

```
# A tibble: 6 x 6
              EXDOSE EXPLDOS EXDURD DOSEO PDOSEO
 USUBJID
  <chr>
               <dbl>
                        <dbl>
                               <dbl> <dbl>
                                             <dbl>
1 01-701-1015
                    0
                            0
                                  15
                                          0
                                                 0
2 01-701-1015
                            0
                                  153
                    0
                                          0
                                                 0
3 01-701-1015
                    0
                            0
                                  14
                                          0
                                                 0
4 01-701-1023
                    0
                            0
                                   23
                                          0
                                                 0
5 01-701-1023
                            0
                                          0
                                                 0
                    0
                                    5
6 01-701-1028
                   54
                           54
                                   14
                                        756
                                               756
```

#### Create 1:1 mapping records

The first set of exposure records to create will be records mapped 1:1 to an existing collected exposure record in SDTM. For these records, the AVAL or AVALC would be calculated using columns that exist on the data and no summarizing of records would be necessary.

These records may be used for input into summary records or be used individually for summarization in outputs. Some examples may be exposure duration, dose administered, dose adjusted, etc. based on one exposure record in SDTM.

These records can be derived using simple dplyr::mutate assignments and then combined

#### **PARAMCD**

Parameter Code

#### **AVALC**

• Analysis Value Character?

```
adex_durd <- adex %>%
 mutate(
   PARAMCD = "DURD",
   AVAL = EXDURD)
adex_dose <- adex %>%
 mutate(
   PARAMCD = "DOSE",
   AVAL = DOSEO)
adex_pldos <- adex %>%
 mutate(
   PARAMCD = "PLDOSE",
   AVAL = PDOSEO)
adex_adj <- adex %>%
 mutate(
   PARAMCD = "ADJ",
   AVALC = if_else(!is.na(EXADJ), "Y", NA_character_)
  )
adex_adjae <- adex %>%
 mutate(
   PARAMCD = "ADJAE",
   AVALC = if_else(EXADJ == "ADVERSE EVENT", "Y", NA_character_)
  )
adex <- bind_rows(</pre>
 adex_durd,
 adex_dose,
 adex_pldos,
  adex_adj,
```

```
adex_adjae) %>%
mutate(PARCAT1 = "INDIVIDUAL") # dim(adex) 2955 40
adex %>% select(USUBJID, VISIT, ASTDT, AENDT, PARAMCD, AVAL, AVALC) %>% head()
```

```
# A tibble: 6 x 7
 USUBJID
              VISIT
                       ASTDT
                                  AENDT
                                              PARAMCD
                                                       AVAL AVALC
  <chr>
              <chr>
                       <date>
                                  <date>
                                              <chr>
                                                      <dbl> <chr>
1 01-701-1015 BASELINE 2014-01-02 2014-01-16 DURD
                                                         15 <NA>
2 01-701-1015 WEEK 2
                       2014-01-17 2014-06-18 DURD
                                                        153 <NA>
3 01-701-1015 WEEK 24 2014-06-19 2014-07-02 DURD
                                                         14 <NA>
4 01-701-1023 BASELINE 2012-08-05 2012-08-27 DURD
                                                         23 <NA>
5 01-701-1023 WEEK 2
                       2012-08-28 2012-09-01 DURD
                                                          5 <NA>
6 01-701-1028 BASELINE 2013-07-19 2013-08-01 DURD
                                                         14 <NA>
```

#### **Create Summary Records**

Exposure is commonly analyzed by a timing interval (e.g. APHASE, APERIOD, AVISIT, etc.). For these types of calculations, the derive\_param\_exposure() function may be used. In addition to creating a summarized AVAL, the function will also compute minimum and maximum dates for the record. For example, to calculate the total dose by subject and treatment

```
adex <- derive_param_exposure(
  dataset = adex
  ,dataset_add = adex
  ,by_vars = exprs(STUDYID, USUBJID, !!!adsl_vars)
  ,input_code = "DOSE"
  ,set_values_to = exprs(
    PARAMCD = "TDOSE",
    PARCAT1 = "OVERALL",
    AVAL = sum(AVAL, na.rm = TRUE)
  )
) # dim(adex) 3209 40</pre>
```

#### **Create Summary Records**

Exposure is commonly analyzed by a timing interval (e.g. APHASE, APERIOD, AVISIT, etc.). For these types of calculations, the derive\_param\_exposure() function may be used. In addition to creating a summarized AVAL, the function will also compute minimum and maximum dates for the record. For example, to calculate the total dose by subject and treatment

A record with PARAMCD == "TDOSE" is created with PARCAT1 set to "OVERALL" using the records in ADEX where PARAMCD == "DOSE" by summing AVAL. In addition, the ASTDT, and AENDT are created as the minimum and maximum date/times associated with each by\_vars grouping. Note that, in addition to PARAMCD, PARCAT1, AVAL, ASTDT and AENDT, only those variables specified in the by\_vars argument will be populated in the new records.

```
adsl_vars <- exprs(TRTSDT, TRTSDTM, TRTEDT, TRTEDTM)

# Remove existing TDOSE records
adex <- adex %>% filter(PARAMCD != "TDOSE")

# Derive new records for the parameter TDOSE (Total Dose) by summarizing exposure data
adex <- derive_param_exposure(
   dataset = adex
   ,dataset_add = adex
   ,by_vars = exprs(STUDYID, USUBJID, !!!adsl_vars)

# Specifies which parameter (DOSE) to summarize. This tells the function to use rows with PAI
   ,input_code = "DOSE"
   ,set_values_to = exprs(
    PARAMCD = "TDOSE"
    ,PARCAT1 = "OVERALL"
   ,AVAL = sum(AVAL, na.rm = TRUE)
   )
   ) # dim(adex) 3209 40</pre>
```

Multiple parameters (records) may be created at one time using the call\_derivation() function:

```
"DOSE" "PLDOSE" "ADJ"
unique(adex$PARAMCD) # [1] "DURD"
                                                             "ADJAE" "TDOSE"
[1] "DURD"
                      "PLDOSE" "ADJ"
             "DOSE"
                                        "ADJAE"
                                                 "TDOSE"
adex <- adex %>%
 filter(!PARAMCD %in% c("TDOSE", "TPDOSE", "TDURD", "TADJ", "TADJAE")) # dim(adex) 2955 40
# Derive new records for multiple parameters by summarizing exposure data
adex <- adex %>%
 call_derivation(
   derivation = derive_param_exposure,
   variable_params = list(
     params(
        set_values_to = exprs(
         PARAMCD = "TDOSE",
         PARCAT1 = "OVERALL",
         AVAL = sum(AVAL, na.rm = TRUE)
       ),
       input_code = "DOSE"
      ),
      params(
       set_values_to = exprs(
         PARAMCD = "TPDOSE",
         PARCAT1 = "OVERALL",
          AVAL = sum(AVAL, na.rm = TRUE)
```

```
input_code = "PLDOSE"
      ),
     params(
       set_values_to = exprs(
        PARAMCD = "TDURD",
         PARCAT1 = "OVERALL",
         AVAL = sum(AVAL, na.rm = TRUE)
       ),
       input_code = "DURD"
      ),
     params(
        set_values_to = exprs(
        PARAMCD = "TADJ",
         PARCAT1 = "OVERALL",
         AVALC = if_else(sum(!is.na(AVALC)) > 0, "Y", NA_character_)
       ),
       input_code = "ADJ"
      ),
     params(
       set_values_to = exprs(
        PARAMCD = "TADJAE",
         PARCAT1 = "OVERALL",
         AVALC = if_else(sum(!is.na(AVALC)) > 0, "Y", NA_character_)
       ),
       input_code = "ADJAE"
     )
   ),
   dataset_add = adex,
   by_vars = exprs(STUDYID, USUBJID, !!!adsl_vars)
 )
count(adex, PARAMCD, PARCAT1)
# A tibble: 10 x 3
  PARAMCD PARCAT1
                        n
  <chr>
         <chr>
                     <int>
 1 ADJ
          INDIVIDUAL
                       591
 2 ADJAE
          INDIVIDUAL
                       591
```

```
3 DOSE
          INDIVIDUAL
                      591
4 DURD
          INDIVIDUAL
                      591
5 PLDOSE INDIVIDUAL
                      591
6 TADJ
          OVERALL
                      254
7 TADJAE OVERALL
                      254
8 TDOSE OVERALL
                      254
9 TDURD OVERALL
                      254
10 TPDOSE OVERALL
                      254
```

```
# Count rows per by group using dplyr, producing the same result as count(adex, PARAMCD, PARCA'
adex %>% group_by(PARAMCD, PARCAT1) %>% summarize(n = n(), .groups = "drop")
```

```
# A tibble: 10 x 3
  PARAMCD PARCAT1
                          n
   <chr>
           <chr>
                      <int>
 1 ADJ
           INDIVIDUAL
                        591
 2 ADJAE
           INDIVIDUAL
                        591
 3 DOSE
           INDIVIDUAL
                        591
4 DURD
           INDIVIDUAL
                        591
5 PLDOSE INDIVIDUAL
                        591
 6 TADJ
           OVERALL
                        254
7 TADJAE OVERALL
                        254
 8 TDOSE
           OVERALL
                        254
 9 TDURD
           OVERALL
                        254
10 TPDOSE OVERALL
                        254
```

Dose intensity can be calculated using the function derive\_param\_doseint(). The planned dose and administered dose are passed into the function and a new record is created with the dose intensity calculation. Again, only those variables specified in the by\_vars argument will be populated in this new record.

```
dim(adex) # 4225 40
```

[1] 4225 40

```
adex <- adex %>%
  derive_param_doseint(
    by_vars = exprs(STUDYID, USUBJID, !!!adsl_vars),
    set_values_to = exprs(PARAMCD = "TNDOSINT"),
    tadm_code = "TDOSE",
    tpadm_code = "TPDOSE") # dim(adex) 4479 40
```

The default calculation for dose intensity is: Administered Doses / Planned Doses \* 100. Please see the derive\_param\_doseint() documentation to see how planned doses of 0 or NA are handled.

#### Assign PARAMCD, PARAMN, etc. from Reference tables

To assign parameter level values such as PARAM, PARAMN, PARCAT1, etc., a lookup can be created to join to the source data.

```
# param_lookup not found

# adex <- derive_vars_merged(
# adex,</pre>
```

```
# dataset_add = param_lookup,
# by_vars = exprs(PARAMCD)
# )
#count(adex, PARAMCD, PARAM, PARAMN)
```

## Derive Categorization Variables (AVALCATx)

#### **AVALCAT1**

{admiral} does not currently have a generic function to aid in assigning AVALCATX/ AVALCAXN values. Below is a simple example of how these values may be assigned using the dplyr::mutate function:

```
adex <- adex %>%
mutate(
    AVALCAT1 = case_when(
        PARAMCD %in% c("TDURD") & AVAL < 30 ~ "< 30 days",
        PARAMCD %in% c("TDURD") & AVAL >= 30 & AVAL < 90 ~ ">= 30 and < 90 days",
        PARAMCD %in% c("TDURD") & AVAL >= 90 ~ ">=90 days",
        PARAMCD %in% c("TDURD") & AVAL >= 90 ~ ">=90 days",
        PARAMCD %in% c("TDOSE", "TPDOSE") & AVAL < 1000 ~ "< 1000 mg",
        PARAMCD %in% c("TDOSE", "TPDOSE") & AVAL >= 1000 ~ ">= 1000 mg",
        TRUE ~ NA_character_
    )
    )
    adex %>% select(USUBJID, VISIT, PARCAT1, PARAMCD, AVAL, AVALCAT1) %>% filter(!is.na(AVALCAT1)) %>%;
```

```
# A tibble: 6 x 6
 USUBJID VISIT PARCAT1 PARAMCD AVAL AVALCAT1
  <chr>
              <chr> <chr> <chr> <chr> <dbl> <chr>
1 01-701-1015 <NA> OVERALL TDOSE
                                         0 < 1000 \text{ mg}
2 01-701-1023 <NA> OVERALL TDOSE
                                         0 < 1000 \text{ mg}
3 01-701-1028 <NA> OVERALL TDOSE
                                     1188 >= 1000 mg
4 01-701-1033 <NA> OVERALL TDOSE
                                      756 < 1000 mg
5 01-701-1034 <NA> OVERALL TDOSE
                                     14067 >= 1000 \text{ mg}
6 01-701-1047 <NA> OVERALL TDOSE
                                          0 < 1000 \text{ mg}
```

#### Assign ASEQ

#### **ASEQ**

Analysis Sequence

The {admiral} function derive var obs number() can be used to derive ASEQ. An example call is:

```
# Not working as PARAMN is not found
#adex <- derive_var_obs_number(
# adex,
# new_var = ASEQ,
# by_vars = exprs(STUDYID, USUBJID),
# order = exprs(PARCAT1, ASTDT, VISIT, VISITNUM, EXSEQ, PARAMN),
# check_type = "error"
#)
#adex %>% select(USUBJID,,VISIT,PARCAT1,PARAMCD,AVAL,ASTDT,ASEQ) %>% head()
```

If needed, the other ADSL variables can now be added:

```
# Deselect variables using negate_vars()
#adex <- adex %>%
# derive_vars_merged(
# dataset_add = select(adsl, !!!negate_vars(adsl_vars)),
# by_vars = exprs(STUDYID, USUBJID)
# )
```

#### **Add Labels and Attributes**

Adding labels and attributes for SAS transport files is supported by the following packages:

- metacore: establish a common foundation for the use of metadata within an R session.
- metatools: enable the use of metacore objects. Metatools can be used to build datasets or enhance columns in existing datasets as well as checking datasets against the metadata.
- xportr: functionality to associate all metadata information to a local R data frame, perform data set level validation checks and convert into a transport v5 file(xpt).

#### References

Creating a BDS Exposure ADaM

ADaM Subject-level Analysis - ADSL Dataset