

Join SDTM with its SUPP dataset (function)

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```
library(safetyData) # a package with sample CDISC data sets
library(tidyverse)
```

NOTES:

- with tidyverse, column names are case sensitive
- tidyr
 - pivot_wider() : transpose from long to wide
 - mutate() : change column from character to numeric
- dplyr
 - select()
 - * keep/drop columns
 - select(-any_of(vars)) : drop columns
 - * reorder columns
 - everything() : columns not explicitly named
 - slice_head(n=) : keep the first n rows
- References:
 - select(-any_of(vars)): https://tidyselect.r-lib.org/reference/all_of.html

UPDATES

- 2025-05-22: Added parameters ds_name and supp_name to the function

```
## Get list of datasets
dataset_list <- data(package = "safetyData")$results[ , "Item"]
dataset_list
```

##	[1]	"adam_adae"	"adam_adlbc"	"adam_adlbh"	"adam_adlbhy"
##	[5]	"adam_adqsadas"	"adam_adqscibc"	"adam_adqsnpix"	"adam_ads1"
##	[9]	"adam_adtte"	"adam_advsv"	"sdtm_ae"	"sdtm_cm"
##	[13]	"sdtm_dm"	"sdtm_ds"	"sdtm_ex"	"sdtm_lb"
##	[17]	"sdtm_mh"	"sdtm_qs"	"sdtm_relrec"	"sdtm_sc"
##	[21]	"sdtm_se"	"sdtm_suppae"	"sdtm_suppdm"	"sdtm_suppds"
##	[25]	"sdtm_supplb"	"sdtm_sv"	"sdtm_ta"	"sdtm_te"
##	[29]	"sdtm_ti"	"sdtm_ts"	"sdtm_tv"	"sdtm_vs"

The function

Parameters

- domain: SDTM domain abbreviation

- ds_name: name of main dataset
- supp_name (default: NULL): name of existing SUPP- dataset

```
join_domain_with_supp_by_seq <- function(domain, ds_name, supp_name=NULL){

## concatenate string to get name of datasets to join and of the SEQ variable
name_main <- paste0(ds_name)

if(!is.null(supp_name)){
  name_supp <- paste0(supp_name)
  name_seq <- paste0(toupper(domain), "SEQ")
}

## get main dataset
if(is.null(supp_name)){
  output <- get(name_main) %>%
    as_tibble()
}
else if(!is.null(supp_name)){
  main <- get(name_main) %>%
    as_tibble() %>%
    rename("SEQ"=name_seq) ## rename --SEQ as SEQ, to simplify join statement below

## Columns in SUPP-- not needed for ADaMs or TLFs
cols_to_drop <- c("STUDYID", "RDOMAIN", "IDVAR", "QLABEL", "QORIG", "QEVAL")

## get SUPP-- dataset and transpose
supp_t <- get(name_supp) %>%
  as_tibble() %>%
  pivot_wider(names_from=QNAM, values_from=QVAL) %>%
  mutate(SEQ = as.numeric(IDVARVAL)) %>%
  select(-any_of(cols_to_drop)) %>%
  select(SEQ, everything())

## join datasets
output <- left_join(main, supp_t, by=c("USUBJID"="USUBJID", "SEQ"="SEQ")) %>%
  rename(!name_seq := "SEQ") ## rename SEQ back to --SEQ, now that the join is done
}

## output result
return(output)
}
```

Function calls

DS domain

```
ds <- join_domain_with_supp_by_seq(domain = "ds", ds_name = "sdtm_ds", supp_name = "sdtm_suplds")

## Warning: Using an external vector in selections was deprecated in tidyselect 1.1.0.
## i Please use `all_of()` or `any_of()` instead.
## # Was:
## data %>% select(name_seq)
```

```
##
## # Now:
## data %>% select(all_of(name_seq))
##
## See <https://tidyselect.r-lib.org/reference/faq-external-vector.html>.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

Warning appears even though any_of() is being used.

Documented workaround shows an example for all_of(): <https://tidyselect.r-lib.org/reference/faq-external-vector.html>

```
# show the first few rows of a tibble
ds %>%
  slice_head(n=3)
```

```
## # A tibble: 3 x 15
##   STUDYID DOMAIN USUBJID DSSEQ DSSPID DSTERM DSDECOD DSCAT VISITNUM VISIT DSDTC
##   <chr>    <chr>  <chr>  <dbl>  <int> <chr>  <chr>  <chr>    <dbl> <chr> <chr>
## 1 CDISCPIL~ DS      01-701~     1    NA PROTO~ COMPLE~ DISP~     13 WEEK~ 2014~
## 2 CDISCPIL~ DS      01-701~     2    NA FINAL~ FINAL ~ OTHE~     13 WEEK~ 2014~
## 3 CDISCPIL~ DS      01-701~     1    24 ADVER~ ADVERS~ DISP~      5 WEEK~ 2012~
## # i 4 more variables: DSSTDTC <chr>, DSSTDY <int>, IDVARVAL <int>,
## #   ENTCRIT <int>
```

```
# show as a dataframe rather than as a tibble (which gets truncated)
ds %>%
  as.data.frame() %>%
  head(3)
```

```
##           STUDYID DOMAIN      USUBJID DSSEQ DSSPID           DSTERM
## 1 CDISCPILOT01      DS 01-701-1015      1      NA PROTOCOL COMPLETED
## 2 CDISCPILOT01      DS 01-701-1015      2      NA      FINAL LAB VISIT
## 3 CDISCPILOT01      DS 01-701-1023      1      24      ADVERSE EVENT
##           DSDECOD           DSCAT VISITNUM VISIT           DSDTC
## 1      COMPLETED DISPOSITION EVENT      13 WEEK 26      2014-07-02
## 2      FINAL LAB VISIT      OTHER EVENT      13 WEEK 26 2014-07-02T11:45
## 3      ADVERSE EVENT DISPOSITION EVENT      5 WEEK 4      2012-09-02
##           DSSTDTC DSSTDY IDVARVAL ENTCRIT
## 1 2014-07-02      182      NA      NA
## 2 2014-07-02      182      NA      NA
## 3 2012-09-02       29      NA      NA
```

AE domain, with SUPPAE

```
ae <- join_domain_with_supp_by_seq(domain = "ae", ds_name = "sdm_ae", supp_name = "sdm_suppa")

# show as a dataframe rather than as a tibble (which gets truncated)
ae %>%
  as.data.frame() %>%
  head(3)
```

```
##           STUDYID DOMAIN      USUBJID AESEQ AESPID           AETERM
## 1 CDISCPILOT01      AE 01-701-1015      1      E07 APPLICATION SITE ERYTHEMA
## 2 CDISCPILOT01      AE 01-701-1015      2      E08 APPLICATION SITE PRURITUS
```

```

## 3 CDISCPILOT01      AE 01-701-1015      3      E06      DIARRHOEA
##      AELLT AELLTCD      AEDECOD AEPTCD      AEHLT
## 1 APPLICATION SITE REDNESS      NA APPLICATION SITE ERYTHEMA      NA HLT_0617
## 2 APPLICATION SITE ITCHING      NA APPLICATION SITE PRURITUS      NA HLT_0317
## 3      DIARRHEA      NA      DIARRHOEA      NA HLT_0148
##      AEHLTCD      AEHLGT AEHLGTCD
## 1      NA HLGT_0152      NA
## 2      NA HLGT_0338      NA
## 3      NA HLGT_0588      NA
##
##      AEBODSYS AEBDSYCD
## 1 GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS      NA
## 2 GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS      NA
## 3      GASTROINTESTINAL DISORDERS      NA
##
##      AESOC AESOCCD AESEV AESER
## 1 GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS      NA MILD      N
## 2 GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS      NA MILD      N
## 3      GASTROINTESTINAL DISORDERS      NA MILD      N
##      AEACN      AEREL      AEOUT AESCAN AESCONG AESDISAB AESDTH
## 1      NA PROBABLE NOT RECOVERED/NOT RESOLVED      N      N      N      N
## 2      NA PROBABLE NOT RECOVERED/NOT RESOLVED      N      N      N      N
## 3      NA      REMOTE      RECOVERED/RESOLVED      N      N      N      N
##      AESHOSP AESLIFE AESOD      AEDTC      AESTDTC      AEENDTC AESTDY AEENDY IDVARVAL
## 1      N      N      N 2014-01-16 2014-01-03      <NA>      2      NA      1
## 2      N      N      N 2014-01-16 2014-01-03      <NA>      2      NA      2
## 3      N      N      N 2014-01-16 2014-01-09 2014-01-11      8      10      3
##      AETRTEM
## 1      Y
## 2      Y
## 3      Y

```

QS domain, without SUPP-

```

qs <- join_domain_with_supp_by_seq(domain = "qs", ds_name = "sdm_qs")

# show as a dataframe rather than as a tibble (which gets truncated)
qs %>%
  as.data.frame() %>%
  head(3)

```

```

##      STUDYID DOMAIN      USUBJID QSSEQ QSTESTCD      QSTEST
## 1 CDISCPILOT01      QS 01-701-1015  5001 ACITM01 WORD RECALL TASK
## 2 CDISCPILOT01      QS 01-701-1015  5016 ACITM01 WORD RECALL TASK
## 3 CDISCPILOT01      QS 01-701-1015  5031 ACITM01 WORD RECALL TASK
##
##      QSCAT QSSCAT QSORRES QSORRESU QSSTRESC
## 1 ALZHEIMER'S DISEASE ASSESSMENT SCALE      <NA>      3      <NA>      3
## 2 ALZHEIMER'S DISEASE ASSESSMENT SCALE      <NA>      2      <NA>      2
## 3 ALZHEIMER'S DISEASE ASSESSMENT SCALE      <NA>      4      <NA>      4
##      QSSTRESN QSSTRESU QSBFL QSDRVFL VISITNUM      VISIT VISITDY      QSDTC QSDY
## 1      3      <NA>      Y      <NA>      3 BASELINE      1 2014-01-02      1
## 2      2      <NA>      <NA>      <NA>      8 WEEK 8      56 2014-03-05      63
## 3      4      <NA>      <NA>      <NA>      10 WEEK 16      112 2014-05-07      126

```