# Package 'admiralophtha'

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**Type** Package

**Title** ADaM in R Asset Library - Ophthalmology

**Version** 1.1.0

**Description** Aids the programming of Clinical Data Standards Interchange Consortium (CDISC) compliant Ophthalmology Analysis Data Model (ADaM) datasets in R. ADaM datasets are a mandatory part of any New Drug or Biologics License Application submitted to the United States Food and Drug Administration (FDA). Analysis derivations are implemented in accordance with the ``Analysis Data Model Implementation Guide" (CDISC Analysis Data Model Team, 2021, <https://www.cdisc.org/standards/foundational/adam/adamig-v1-3-release-package>).

**License** Apache License (>= 2)

**URL** https://pharmaverse.github.io/admiralophtha/, https://github.com/pharmaverse/admiralophtha/

**BugReports** https://github.com/pharmaverse/admiralophtha/issues/

**Depends** R (>= 4.0)

**Imports** admiral (>= 1.1.1), admiraldev (>= 1.1.0), dplyr (>= 1.0.5), lubridate (>= 1.7.4), magrittr (>= 1.5), purrr (>= 0.3.3), rlang (>= 0.4.4), stringr (>= 1.4.0), hms (>= 0.5.3), tidyr (>= 1.0.2), tidyselect (>= 1.1.0), lifecycle (>= 0.1.0)

**Suggests** pharmaversesdtm (>= 1.0.0), devtools, diffdf, lintr, pkgdown, testthat (>= 3.0.0), knitr, lintr, methods, miniUI, pharmaversesdtm, pkgdown, rmarkdown, roxygen2, spelling, testthat, tibble, usethis

**VignetteBuilder** knitr

**Encoding** UTF-8

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**RoxygenNote** 7.3.1

**NeedsCompilation** no

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`admiralophtha_adbcva` *Best Corrected Visual Acuity Analysis Dataset*

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## Description

An example Best Corrected Visual Acuity (BCVA) analysis dataset

## Usage

```
admiralophtha_adbcva
```

## Format

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 7672 rows and 116 columns.

## Source

Derived from the OE and ADSL datasets using {admiral}, {admiralophtha} and (https://github.com/pharmaverse/admiralophtha/blob/main/inst/templates/ad_adbcva.R)

## See Also

Other datasets: admiralophtha_adoe, admiralophtha_advfq

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# admiralophtha_adoe *Ophthalmology Exam Analysis Dataset*

---

## Description

An example Ophthalmology Exam Analysis dataset

## Usage

```
admiralophtha_adoe
```

## Format

An object of class tbl_df (inherits from tbl, data.frame) with 7672 rows and 98 columns.

## Source

Derived from the OE and ADSL datasets using {admiral}, {admiralophtha} and (https://github.com/pharmaverse/admiralophtha/blob/main/inst/templates/ad_adoe.R)

## See Also

Other datasets: admiralophtha_adbcva, admiralophtha_advfq

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# admiralophtha_advfq *Visual Function Questionnaire Analysis Dataset*

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## Description

An example Visual Function Questionnaire (VFQ) analysis dataset

## Usage

```
admiralophtha_advfq
```

## Format

An object of class tbl_df (inherits from tbl, data.frame) with 28798 rows and 41 columns.

## Source

Derived from the ADSL and QS datasets using {admiral}, {admiralophtha} and (https://github.com/pharmaverse/admiralophtha/blob/main/inst/templates/ad_advfq.R)

## See Also

Other datasets: admiralophtha_adbcva, admiralophtha_adoe

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`convert_etdrs_to_logmar` *ETDRS –> LogMAR conversion*

---

## Description

Convert ETDRS score to LogMAR units

## Usage

```
convert_etdrs_to_logmar(value)
```

## Arguments

| | |
|---|---|
| value | object containing ETDRS score to convert to logMAR |

## Details

ETDRS value converted to logMAR as logMAR = -0.02 * ETDRS + 1.7

Source for conversion formula: Beck, R.W., et al. A computerized method of visual acuity testing. American Journal of Ophthalmology, 135(2), pp.194-205. doi:https://doi.org/10.1016/s0002-9394(02)01825-1.

## Value

The input value converted converted to logMAR units

## Author(s)

Rachel Linacre

## Examples

```
library(tibble)
library(dplyr)
library(admiral)
library(admiraldev)

adbcva <- tribble(
  ~STUDYID, ~USUBJID, ~AVAL,
```

```
  "XXX001", "P01", 5,
  "XXX001", "P02", 10,
  "XXX001", "P03", 15,
  "XXX001", "P04", 20,
  "XXX001", "P05", 25
)

adbcva <- adbcva %>% mutate(AVAL = convert_etdrs_to_logmar(AVAL))
```

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`convert_logmar_to_etdrs` *LogMAR –> ETDRS conversion*

---

## Description

Convert LogMAR score to ETDRS units

## Usage

```
convert_logmar_to_etdrs(value)
```

## Arguments

| | |
|---|---|
| `value` | object containing logMAR score to convert to ETDRS |

## Details

logMAR value converted to ETDRS as ETDRS = -(logMAR - 1.7) / 0.02

Source for conversion formula: Beck, R.W., et al. A computerized method of visual acuity testing. American Journal of Ophthalmology, 135(2), pp.194-205. doi:https://doi.org/10.1016/s0002-9394(02)01825-1.

## Value

The input value converted to ETDRS units

## Author(s)

Nandini R Thampi

## Examples

```
library(tibble)
library(dplyr)
library(admiral)

oe <- tribble(
  ~STUDYID, ~USUBJID, ~OETESTCD, ~OEMETHOD, ~OESTRESN,
  "XXX001", "P01", "VACSCORE", "logMAR EYE CHART", 1.08,
```

```
  "XXX001", "P02", "VACSCORE", "logMAR EYE CHART", 1.66,
  "XXX001", "P03", "VACSCORE", "logMAR EYE CHART", 1.60,
  "XXX001", "P04", "VACSCORE", "ETDRS EYE CHART", 57,
  "XXX001", "P05", "VACSCORE", "ETDRS EYE CHART", 1
)

adbcva <- oe %>%
  filter(OETESTCD == "VACSCORE" & toupper(OEMETHOD) == "LOGMAR EYE CHART") %>%
  mutate(OESTRESN = convert_logmar_to_etdrs(OESTRESN))
```

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`derive_var_afeye` *Derive Affected Eye*

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## Description

Derive Affected Eye (`AFEYE`) in occurrence datasets

## Usage

```
derive_var_afeye(
  dataset,
  dataset_occ,
  loc_var,
  lat_var,
  lat_vals,
  loc_vals = "EYE"
)
```

## Arguments

| | |
|---|---|
| `dataset` | Input dataset |
| `dataset_occ` | Input dataset<br>**[Deprecated]** Please use `dataset` instead. |
| `loc_var` | Location variable |
| `lat_var` | Laterality variable |
| `lat_vals` | `xxLAT`values for which `AFEYE` is derived<br>**[Deprecated]** Please simply ensure `xxLAT` values are contained in `c("LEFT", "RIGHT", "BILATERAL")`. |
| `loc_vals` | `xxLOC` values for which `AFEYE` is derived |

## Details

Affected Eye is derived in the occurrence dataset using laterality and Study Eye. This assumes Study Eye has already been added from ADSL.

## Value

The input occurrence dataset with Affected Eye (AFEYE) added.

## Author(s)

Lucy Palmen

## Examples

```
library(tibble)
library(admiral)

adae1 <- tribble(
  ~STUDYID, ~USUBJID, ~STUDYEYE, ~AELOC, ~AELAT,
  "XXX001", "P01", "RIGHT", "EYE", "RIGHT",
  "XXX001", "P01", "RIGHT", "EYE", "LEFT",
  "XXX001", "P01", "RIGHT", "EYE", "",
  "XXX001", "P01", "RIGHT", "", "RIGHT",
  "XXX001", "P02", "LEFT", "", "",
  "XXX001", "P02", "LEFT", "EYE", "LEFT",
  "XXX001", "P04", "BILATERAL", "EYE", "RIGHT",
  "XXX001", "P05", "RIGHT", "EYE", "RIGHT",
  "XXX001", "P05", "RIGHT", "EYE", "BILATERAL",
  "XXX001", "P06", "BILATERAL", "", "",
  "XXX001", "P06", "BILATERAL", "", "RIGHT",
  "XXX001", "P07", "BILATERAL", "EYE", "BILATERAL",
  "XXX001", "P08", "", "EYE", "BILATERAL",
  "XXX001", "P09", "NONSENSE", "EYE", "BILATERAL",
  "XXX001", "P09", "BILATERAL", "EYE", "NONSENSE",
  "XXX001", "P09", "BILATERAL", "NONSENSE", "BILATERAL",
  "XXX001", "P10", "RIGHT", "EYE", "BOTH"
)

derive_var_afeye(adae1, loc_var = AELOC, lat_var = AELAT)

adae2 <- tribble(
  ~STUDYID, ~USUBJID, ~STUDYEYE, ~AELOC, ~AELAT,
  "XXX001", "P01", "RIGHT", "EYES", "RIGHT",
  "XXX001", "P02", "RIGHT", "RETINA", "LEFT",
  "XXX001", "P03", "LEFT", "", ""
)

derive_var_afeye(adae2, loc_var = AELOC, lat_var = AELAT, loc_vals = c("EYES", "RETINA"))
```

---

# derive_var_bcvacritxfl

*Adds* CRITx/CRITxFL *pairs to BCVA dataset*

---

## Description

Adds a criterion variables CRITx and their corresponding flags CRITxFL to a dataset containing BCVA records

## Usage

```
derive_var_bcvacritxfl(
  dataset,
  crit_var,
  bcva_ranges = NULL,
  bcva_uplims = NULL,
  bcva_lowlims = NULL,
  additional_text = "",
  critxfl_index = NULL
)
```

## Arguments

| | |
|---|---|
| dataset | Input dataset containing BCVA data (usually ADBCVA). |
| crit_var | Variable with respect to which CRITx/CRITxFL are derived (usually CHG or AVAL). |
| bcva_ranges | List containing one or more numeric vectors of length 2. For each vector c(a,b) in bcva_ranges, a pair of variables CRITx, CRITxFL is created with the condition: a <= crit_var <= b. If criterion flags of that type are not required, then leave as NULL. |
| bcva_uplims | List containing one or more numeric elements. For each element a in bcva_uplims, a pair of variables CRITx, CRITxFL is created with the condition: crit_var <= a. If criterion flags of that type are not required, then leave as NULL. |
| bcva_lowlims | List containing one or more numeric elements. For each element b in bcva_lowlims, a pair of variables CRITx, CRITxFL is created with the condition: crit_var >= b. If criterion flags of that type are not required, then leave as NULL. |
| additional_text | string containing additional text to append to CRITx |
| critxfl_index | positive integer detailing the first value of x to use in CRITxFL. If not supplied, the function takes the first available value of x, counting up from x = 1. |

## Details

This function works by calling derive_var_bcvacritxfl() once for each of the elements in bcva_ranges, bcva_uplims and bcva_lowlims. NOTE: if crit_var is equal to NA, then the resulting criterion flag is also marked as NA.

## Value

The input BCVA dataset with additional column pairsCRITx, CRITxFL.

## Author(s)

Edoardo Mancini

## Examples

```
library(tibble)
library(admiral)
library(admiraldev)

adbcva1 <- tribble(
  ~STUDYID, ~USUBJID, ~AVISIT, ~BASETYPE, ~PARAMCD, ~CHG,
  "XXX001", "P01", "BASELINE", "LAST", "SBCVA", 0,
  "XXX001", "P01", "WEEK 2", "LAST", "FBCVA", 2,
  "XXX001", "P02", "BASELINE", "LAST", "SBCVA", -13,
  "XXX001", "P02", "WEEK 2", "LAST", "FBCVA", 5,
  "XXX001", "P03", "BASELINE", "LAST", "SBCVA", NA,
  "XXX001", "P03", "WEEK 2", "LAST", "FBCVA", 17
)

derive_var_bcvacritxfl(
  dataset = adbcva1,
  crit_var = exprs(CHG),
  bcva_ranges = list(c(0, 5), c(-5, -1), c(10, 15)),
  bcva_uplims = list(5, 10),
  bcva_lowlims = list(8),
  additional_text = ""
)

adbcva2 <- tribble(
  ~STUDYID, ~USUBJID, ~AVISIT, ~BASETYPE, ~PARAMCD, ~AVAL, ~CHG,
  "XXX001", "P01", "BASELINE", "LAST", "SBCVA", 4, NA,
  "XXX001", "P01", "BASELINE", "LAST", "SBCVA", 6, NA,
  "XXX001", "P01", "AVERAGE BASELINE", "AVERAGE", "SBCVA", 5, NA,
  "XXX001", "P01", "WEEK 2", "LAST", "SBCVA", -3, NA,
  "XXX001", "P01", "WEEK 4", "LAST", "SBCVA", -10, NA,
  "XXX001", "P01", "WEEK 6", "LAST", "SBCVA", 12, NA,
  "XXX001", "P01", "WEEK 2", "AVERAGE", "SBCVA", -2, -7,
  "XXX001", "P01", "WEEK 4", "AVERAGE", "SBCVA", 6, 1,
  "XXX001", "P01", "WEEK 6", "AVERAGE", "SBCVA", 3, -2
)

restrict_derivation(
  adbcva2,
  derivation = derive_var_bcvacritxfl,
  args = params(
    crit_var = exprs(CHG),
    bcva_ranges = list(c(0, 5), c(-10, 0)),
    bcva_lowlims = list(5),
    additional_text = " (AVERAGE)"
  ),
  filter = PARAMCD %in% c("SBCVA", "FBCVA") & BASETYPE == "AVERAGE"
)
```

`derive_var_studyeye` *Derive Study Eye*

## Description

Derive Study Eye (`STUDYEYE`) in the ADSL dataset

## Usage

```
derive_var_studyeye(dataset_adsl, dataset_sc, sctestcd_value = "FOCID")
```

## Arguments

| | |
|---|---|
| `dataset_adsl` | ADSL input dataset |
| `dataset_sc` | SC input dataset |
| `sctestcd_value` | `SCTESTCD` value flagging Study Eye selection records. Default: "FOCID". |

## Details

Study Eye is derived in ADSL using the "Study Eye selection" records in the SC SDTM dataset.

## Value

The input ADSL dataset with an additional column named `STUDYEYE`

## Author(s)

Edoardo Mancini

## Examples

```
library(tibble)
library(admiral)

adsl <- tribble(
  ~STUDYID, ~USUBJID,
  "XXX001", "P01",
  "XXX001", "P02",
  "XXX001", "P03",
  "XXX001", "P04",
  "XXX001", "P05"
)

sc <- tribble(
  ~STUDYID, ~USUBJID, ~SCTESTCD, ~SCSTRESC,
  "XXX001", "P01", "FOCID", "OS",
  "XXX001", "P01", "ACOHORT", "COHORT1",
  "XXX001", "P02", "FOCID", "OD",
  "XXX001", "P02", "ACOHORT", "COHORT3",
```

```
  "XXX001", "P04", "FOCID", "OU",
  "XXX001", "P05", "FOCID", "OD",
  "XXX001", "P06", "FOCID", "OS"
)

derive_var_studyeye(adsl, sc)
```

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