

# Package ‘admiralophtha’

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**Type** Package

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

**Version** 1.4.0

**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 (>= 4.1)

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

**Suggests** devtools, diffdf, knitr, lintr, methods, miniUI,  
pharmaversesdtm (>= 1.3.0), pkgdown, rmarkdown, roxygen2,  
spelling, testthat (>= 3.0.0), tibble, usethis, withr

**VignetteBuilder** knitr

**Config/testthat/edition** 3

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**RoxygenNote** 7.3.3

**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 the [ADBCVA template](#).

**See Also**

Other datasets: [admiralophtha\\_adoe](#), [admiralophtha\\_advfq](#)

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*admiralophtha\_adoe*      *Ophthalmology Exam Analysis Dataset*

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**Description**

An example Ophthalmology Exam Analysis dataset

**Usage**

`admiralophtha_adoe`

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 19136 rows and 103 columns.

**Source**

Derived from the OE and ADSL datasets using `{admiral}`, `{admiralophtha}` and the [ADOE template](#).

**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 972 rows and 93 columns.

### Source

Derived from the ADSL and QS datasets using `{admiral}`, `{admiralophtha}` and the [ADVFAQ template](#). The full, open-source VFQ questionnaire can be accessed [here](#).

### 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.  
**Permitted values** a numeric value, e.g. 2, -5, 1.4  
**Default value** none

**Details**

ETDRS value converted to logMAR as:

$$\log MAR = -0.02 * ETDRS + 1.7$$

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](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))
```

---

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.

**Permitted values** a numeric value, e.g. 2, -5, 1.4

**Default value** none

**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](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	<i>Derive Affected Eye</i>
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### Description

Derive Affected Eye (AFEYE) in occurrence datasets

### Usage

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

### Arguments

dataset	Input dataset. <b>Permitted values</b> a dataset, i.e., a data.frame or tibble <b>Default value</b> none
loc_var	Location variable, usually XXLOC. <b>Permitted values</b> an unquoted symbol, e.g., AVAL <b>Default value</b> none
lat_var	Laterality variable, usually XXLAT. <b>Permitted values</b> an unquoted symbol, e.g., AVAL <b>Default value</b> none
loc_vals	xxLOC values for which AFEYE is derived. <b>Permitted values</b> a character vector, e.g. c("EYE", "RETINA") <b>Default value</b> "EYE"

### 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**

**[Deprecated]** The `derive_var_bcvacritxfl()` function has been deprecated in favor of `admiral::derive_vars_crit_fl` - please see the [criterion flag section of the ADBCVA vignette](#) for more details.

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). <b>Permitted values</b> a dataset, i.e., a data.frame or tibble <b>Default value</b> none
crit_var	Variable with respect to which CRITx/CRITxFL are derived (usually CHG or AVAL). <b>Permitted values</b> an unquoted symbol, e.g., AVAL <b>Default value</b> none
bcva_ranges	List of numeric vectors. 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. <b>Permitted values</b> a list containing one or more numeric vectors, each of length two. E.g. list(c(1, 2), c(3, 4)) <b>Default value</b> 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. <b>Permitted values</b> a list containing one or more numeric scalars. E.g. list(2, -4) <b>Default value</b> 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. <b>Permitted values</b> a list containing one or more numeric scalars. E.g. list(2, -4) <b>Default value</b> NULL
additional_text	string containing additional text to append to CRITx. <b>Permitted values</b> a character scalar, i.e., a character vector of length one <b>Default value</b> ""
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. <b>Permitted values</b> a positive integer, e.g. 2 or 5 <b>Default value</b> NULL

**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 pairs `CRITx`, `CRITxFL`.

**Author(s)**

Edoardo Mancini

**See Also**

Other deprecated: [derive\\_var\\_bcvacritxfl\\_util\(\)](#)

**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. <b>Permitted values</b> a dataset, i.e., a <code>data.frame</code> or <code>tibble</code> <b>Default value</b> none
dataset_sc	SC input dataset. <b>Permitted values</b> a dataset, i.e., a <code>data.frame</code> or <code>tibble</code> <b>Default value</b> none
sctestcd_value	SCTESTCD value flagging Study Eye selection records. <b>Permitted values</b> a character scalar, i.e., a character vector of length one <b>Default value</b> "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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