

# Package ‘eye’

June 25, 2020

**Title** Analysis of Eye Data

**Version** 0.1.0

**Description** A tool to facilitate common tasks in ophthalmic research:

Conversion between different visual acuity notations (Snellen, logMAR and ETDRS), counting of patients, recode right and left eyes and reshape eye side specific variables between wide and long format. The 'eye' package also contains a real life data set of people with intravitreal injections with anti-vascular endothelial growth factor (anti-VEGF), made available by Fasler et al. (2019) <doi:10.1136/bmjopen-2018-027441>. Visual acuity conversion is based on Schulze-Bonsel et al. (2006) <doi:10.1167/iovs.05-0981>, Gregori et al. (2010) <doi:10.1097/iae.0b013e3181d87e04>, Beck et al. (2003) <doi:10.1016/s0002-9394(02)01825-1> and Bach (2007) <http:michaelbach.de/sci/acuity.html>.

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age . . . . .	2
amd . . . . .	3
blink . . . . .	4
clean_va . . . . .	6
eyes . . . . .	7
hyperop . . . . .	9
myop . . . . .	10
print_methods . . . . .	11
recodeye . . . . .	12
reveal . . . . .	13
reveal_methods . . . . .	14
snellen_steps . . . . .	15
va . . . . .	16
va_chart . . . . .	19
va_methods . . . . .	20

<b>Index</b>	<b>22</b>
--------------	-----------

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age	<i>age</i>
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**Description**

calculates age in years, as durations or periods

**Usage**

```
age(from_date, to_date = lubridate::now(), period = FALSE, dec = 1)
```

**Arguments**

from_date	start date
to_date	end date
period	Calculating period (TRUE) or duration (FALSE- default)
dec	How many decimals are displayed

**Value**

Numeric vector

**Author(s)**

Antoine Fabri and Tjebo Heeren

**See Also**

[OP on stackoverflow](#) from which this function was inspired. [Read about periods and durations](#)

## Examples

```
age("1984-10-16")

dob <- c("1984-10-16", "2000-01-01")
test_date <- as.Date(dob) + c(15000, 20000)
age(dob, test_date)
```

---

amd

*Real life data of patients with neovascular AMD*

---

## Description

A dataset containing anonymized real life human subjects data on eyes with treatment naive neovascular age-related macular degeneration (AMD), which underwent intravitreal anti-VEGF therapy with ranibizumab and/or aflibercept.

## Usage

```
data("amd")
```

## Format

A data frame with 40764 rows and 7 variables:

**Id** Anonymized patient identifier

**Eye** Left or right eye of patient (0 = right, 1 = left)

**FollowupDays** Days after date of first appointment (0 = first appointment)

**BaselineAge** Age (years) at day of first appointment

**Gender** Gender of patient (0 = male, 1 = female)

**VA\_ETDRS\_Letters** Visual acuity in Early Treatment Diabetic Retinopathy Study letters

**InjectionNumber** Current number of injection at appointment date

## Details

The data was collected in Moorfields Eye Hospital, London, UK. (Information governance sign off Moorfields Eye Hospital 19/07/2018)

Data was accessed on the 25th May 2020

## Spurious data entries

Note there are erroneous visual acuity entries in this data set which I noticed during the work on this package. The data set curator has been contacted and it was concluded that these were erroneous entries in the original medical health records. I decided to keep the values in the data set and wait for the final decision how to proceed from the data set curator (if they are going to replace it with missing values or not). I believe this is a great example for the challenges of real life data and a reminder to remain vigilant when doing data analysis.

**Source**

<https://datadryad.org/stash/dataset/doi:10.5061/dryad.97r9289>

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blink

*Your data in a blink of an eye*

---

**Description**

blink summarizes your data tailored to the need of ophthalmic research: It looks for VA and IOP columns and summarises those with common statistics. In order to make it work, it requires specific column naming - please see section "column names" and "data coding". For more details how blink works, see `vignette("eye")`

**Usage**

```
blink(x, va_to = "logmar", va_cols = NULL, iop_cols = NULL, fct_level = 0:4)
```

**Arguments**

<code>x</code>	data frame
<code>va_to</code>	to which VA notation (passed to <code>va()</code> )
<code>va_cols</code>	if specified, overruling automatic VA columns selection. tidyselection supported
<code>iop_cols</code>	if specified, overruling automatic IOP columns selection. tidyselection supported
<code>fct_level</code>	Remove columns for Summarizing when all unique values fall into range. character or numeric vector, default 1:4

**Details**

blink is basically a wrapper around [myop](#), [eyes](#) and [reveal](#):

- Duplicate rows are always removed
- Column names are prepared for myopization (see [myop](#))
- VA will always be converted to logmar

**Value**

object of class `blink` and `list`. Class `blink` contains the myopized data, count of patients and eyes, and summaries for visual acuities and intraocular pressure.

**Data coding**

- Only common codes supported:
- **eyes**: "r", "re", "od", "right" - or numeric coding `r:l = 0:1` or `1:2`
- **Visual acuity**: "VA", "BCVA", "Acuity"
- **Intraocular pressure**: "IOP", "GAT", "NCT", "pressure"

### Column name rules

- No spaces!
- Do not use numeric coding for eyes in column names
- Separate eye and VA and IOP codes with **underscores** ("bcva\_l\_preop", "VA\_r", "left\_va", "IOP\_re")
- Avoid separate VA or IOP codes if this is not actually containing VA/ IOP data (e.g. "stableVA" instead of "stable\_va", ChangeIOP instead of "change\_IOP")
- Keep names short
- Don't use underscores when you don't have to. Consider each section divided by an underscore as a relevant characteristic of your variable. ("preop" instead of "pre\_op", "VA" instead of "VA\_ETDRS\_Letters")
- Use common codes for your patient column (see [eyes](#), section Guessing) (e.g., "pat", "patient" or "ID", ideally both: "patientID" or "patID")
- **Don't be too creative with your names!**

### Names examples

#### Good names:

```
-c("patid", "surgery_right", "iop_r_preop", "va_r_preop", "iop_r", "iop_l")
```

#### OK names

`-c("Id", "Eye", "BaselineAge", "VA_ETDRS_Letters", "InjectionNumber")`: Names are long and there are two unnecessary underscore in the VA column. Better just "VA" `-c("id", "r", "l")`: All names are commonly used (good!), but which dimension of "r"/"l" are we exactly looking at?

#### Bad names (eye will fail)

- `c("id", "iopr", "iop_l", "VA_r", "VA_l")`: eye won't be able to recognize IOP and VA columns
- `c("id", "iop_r", "iop_l", "stable_iop_r", "stable_iop_l")`: eye *may* wrongly identify the (probably logical) columns "stable\_iop" as columns containing IOP data. Better maybe: "stableIOP\_l"
- `c("person", "goldmann", "vision")`: eye will not recognize that at all

### tidy data

**blink and myop work more reliably with clean data** (any package will, really!). [clean data](#).

### column removal

Done with [remCols](#): Removes columns that only contain values defined in `fmt_levels` or logicals from selected columns (currently for both automatically and manually selected columns). `fmt_levels` are removed because they are likely categorical codes.

**See Also**

[About tidyselection.](#)

How to rename your columns (two threads on stackoverflow.com):

- [Rename columns 1](#)
- [Rename columns 2](#)

**Examples**

```
blink(amd)

messy_df <- data.frame( id = letters[1:3],
  iop_r_preop = sample(21:23), iop_r_postop = sample(11:13),
  iop_l_postop = sample(11:13), iop_l_preop = sample(31:33),
  va_r_preop = sample(41:43), va_l_preop = sample(41:43),
  va_r_postop = sample(51:53), va_l_postop = sample(45:47)
)
blink(messy_df)
```

---

clean\_va

*Visual acuity entry cleaner*

---

**Description**

VA cleaning:

1. `isNAstring()`: Replacing empty placeholders (".", "", "(any number of empty space)", "NULL", "NA", "N/A" ) - any cases - with NA
2. `convert_NLP()` Simplifying the notation for qualitative VA notation (NPL becomes NLP, PL becomes LP)
3. Removing non-Snellen character strings

**Usage**

```
clean_va(x, quali = c("nlp", "lp", "hm", "cf"))

convert_NLP(x, replace_PL = c(pl = "lp", npl = "nlp"), tolower = TRUE)

isNAstring(
  x,
  full = c("\\.+", "", "\\s+", "n/a", "na", "null"),
  tolower = TRUE
)
```

**Arguments**

x	Vector with VA entries
quali	strings for qualitative visual acuity entries
replace_PL	named vector how to rename qualitative VA
tolower	if TRUE, x will be converted to lower first
full	vector of full strings to be replaced by NA

**Value**

character vector

**See Also**

Other VA cleaner: [va\(\)](#)

Other VA cleaner: [va\(\)](#)

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eyes	<i>Count patients and eyes</i>
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**Description**

Counts number of patients and eyes (right and left).

eyestr: identical to eyes(x, report = TRUE, ...)

**Usage**

```
eyes(x, id = NULL, eye = NULL, report = FALSE, ...)
```

```
eyestr(x, id = NULL, eye = NULL, small_num = TRUE, para = FALSE, UK = FALSE)
```

**Arguments**

x	required. (data frame)
id	Patient identifying column
eye	Eye identifying column.
report	if TRUE, text returned for report
...	passed to <a href="#">eyes_to_string</a>
small_num	If TRUE: writing numbers <= 12 as words
para	If TRUE: Adding "A total of" to comply with most journal standards and to avoid awkward long numbers.
UK	Logical, Use UK (English) style (TRUE) or USA (American) style (FALSE).

## Details

eyes guesses columns that identify patients and eyes.

## Value

eyes: Named integer vector with count of patients and eyes

eyestr: Character string - can be directly pasted into reports

## Guessing

For any below, **cases are always ignored** (you can write in upper or lower case, as you please)

**id** and **eye** arguments overrule the name guessing for the respective columns.

### patient ID columns:

- First, eyes is looking for names that contain both strings "pat" and "id" (the order doesn't matter)
- Next, it will look for columns that are plainly called "ID"
- Last, it will search for all names that contain either "pat" or "id"

### eye variable column:

- eyes looks for columns called either "eye" or "eyes"

## Eye coding

- eyes recognizes integer coding 0:1 and 1:2, with right being the lower number. For strings coding it recognizes right eyes: c("r", "re", "od", "right") and left eyes: c("l", "le", "os", "left")

## Report

Using [eyes\\_to\\_string](#) to parse the output of eyes into a text which you can use for reports. Arguments to `eyes_to_string` are passed via ...:

- **small\_num** If TRUE (default): numbers  $\leq 12$  as words
- **para** If TRUE (not default): Adding "A total of" to comply with most journal standards and to avoid awkward long numbers.
- **UK** TRUE: UK style (English) or FALSE (default): US style (American).

## Examples

```
eyes(amd)
eyestr(amd, para = TRUE)
```



---

hyperop	<i>Hyperopic eye data</i>
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---

## Description

Pivot eye-related variables to two columns

## Usage

```
hyperop(x, cols, eye = NULL)
```

## Arguments

x	data frame
cols	columns which should be made "wide". Tidyselection supported
eye	eye column (default looking for "eye" or "eyes", all cases)

## Details

Basically the opposite of `myop()` - a slightly intelligent wrapper around `tidyr::pivot_longer()` and `tidyr::pivot_wider()` Will find the eye column, unify the codes for the eyes (all to "r" and "l") and pivot the columns wide, that have been specified in "cols".

### Good names and tidy data always help!

For more information about shaping data and good names, see `vignette("eye")`, or `?blink` or `?myop`

## Value

A tibble, see also [tibble::tibble](#)

## See Also

[About tidyselection](#)

## Examples

```
# Example to clean a bit messy data frame

iopva <- data.frame(
  id = c("a", "e", "j", "h"),
  va_r = c(37L, 36L, 33L, 38L),
  iop_r = c(38L, 40L, 33L, 34L),
  va_l = c(30L, 39L, 37L, 40L),
  iop_l = c(31L, 34L, 33L, 31L)
)
myop_iop <- myop(iopva)
hyperop(myop_iop, cols = matches("va|iop"))
```

---

myop	<i>Myopic eye data</i>
------	------------------------

---

## Description

Pivot "eye" variable to one column

## Usage

```
myop(x, var = "value")
```

```
myopic(x, var = "value")
```

## Arguments

x	data frame
var	Character vector of length 1 specifying the variable if there is only one column per eye with no further info on the variable (default "value")

## Details

Out of convenience, data is often entered in a very "wide" format: there will be two columns for the same variable, one column for each eye. `myop` will pivot the eye variable to one column and keep all other variables wide. E.g., eight columns that store data of four variables for two eyes will be pivoted to 5 columns (one eye and four further variable columns, see also *examples*).

### **myop requires a specific data format**

If there is a column called "eye" or "eyes", `myop` will not make any changes - because the data is then already assumed to be in long format. If you *also* have columns with eye-specific values, then you have messy data. Maybe, you could remove or rename the "eye" column and then let `myop` do the work.

`myop` will only recognize meaningful coding for eyes:

- Right eyes: "r", "re", "od", "right"
- Left eyes: "l", "le", "os", "left"
- for other codes see also [set\\_codes](#) The strings for eyes need to be **separated by period or underscores**. (Periods will be replaced by underscores). Any order is allowed.
- **Will work:** "va\_r", "right\_morningpressure", "night\_iop.le", "gat\_os\_postop"
- **Will fail:** "VAr", "rightmorningPressure", "night\_IOPl", "gatOSpostop"

An exception is when there is only one column for each eye. Then the column names can consist of eye strings (see above) only. In this case, `var` will be used to "name" the resulting variable.

If there are only eye columns in your data (should actually not happen), `myop` will create identifiers by row position.

**Please always check the result for plausibility.** Depending a lot on how the data was entered, the results could become quite surprising. There is basically a nearly infinite amount of possible combinations of how to enter data, and it is likely that `myop` will not be able to deal with all of them

**Value**

A tibble, see also [tibble::tibble](#)

**internal preparation**

- Rename data names with [myop\\_rename](#), replacing "." with "\_"
- Use of [sort\\_substr\(\)](#) - sorting eye strings first, then strings coding for methods (IOP/VA), then the rest.

**myopization**

The actual work is done with [myopizer](#) and [myop\\_pivot](#)

**Examples**

```
# Example to clean a bit messy data frame
iopva <- data.frame(
  id = c("a", "e", "j", "h"),
  va_r = c(37L, 36L, 33L, 38L),
  iop_r = c(38L, 40L, 33L, 34L),
  va_l = c(30L, 39L, 37L, 40L),
  iop_l = c(31L, 34L, 33L, 31L)
)
myop(iopva)

iop_wide <- data.frame(id = letters[1:3], r = 11:13 , l = 14:16)
# the variable has not been exactly named, so you can specify
# it with the var argument
myop(iop_wide, var = "iop")
```

---

print\_methods

*print eye classes*

---

**Description**

S3 methods for VA classes "snellen", "logmar" and "etdrs". **snellen** is always also a character class because it is more categorical than continuous. **logmar** and **etdrs** are both numerics (logMAR is double, etdrs is integer).

S3 methods for blink class

**Usage**

```
## S3 method for class 'snellen'
print(x, ...)

## S3 method for class 'logmar'
print(x, ...)
```

```
## S3 method for class 'etdrs'
print(x, ...)
```

```
## S3 method for class 'blink'
print(x, ...)
```

### Arguments

x                    object of class "blink"  
 ...                  arguments passed to [print.default](#)

### Value

No return value, called for side effects (printing)

---

recodeye

*Recode eyes*

---

### Description

recoding eyes to "r" and "l"

### Usage

```
recodeye(
  x,
  to = c("r", "l"),
  eyecodes = list(c("r", "re", "od", "right"), c("l", "le", "os", "left")),
  numcode = NULL
)
```

### Arguments

x                    vector of strings  
 to                    to which codes. Right coded first.  
 eyecodes            list of substrings which should be converted to right and left eyes - first vector for right, then for left eyes  
 numcode            if you have numeric coding which is not 0:1 or 1:2 for right:left, specify it here.

### Value

Character vector

### See Also

Other string matching functions: [getElem](#), [set\\_codes\(\)](#), [sort\\_substr\(\)](#), [str\\_func\\_facs](#)

**Examples**

```
x <- c("r", "re", "od", "right", "l", "le", "os", "left")
recodeye(x)
## chose the resulting codes
recodeye(x, to = c("right", "left"))
x <- 1:2
recodeye(x)
## or, if right is coded with 2)
recodeye(x, numcode = 2:1)
```

---

 reveal

*reveal*


---

**Description**

Shows commonly used summary statistics

**Usage**

```
reveal(x, by = NULL, dec = 1, funs = NULL)
```

**Arguments**

x	data frame, numeric vector, or list of numeric vectors
by	character vector with the names of the columns. Can be several variables!
dec	how many decimals are displayed
funs	not really meant to be used at the moment - change the Summarizing functions with a named(!) list of functions

**Details**

Character vectors (or character columns) will be removed.

**Value**

data frame

**See Also**

Other revealers: [reveal\\_methods](#), [reveal\\_split\(\)](#)

**Examples**

```
x = y = z = c(rnorm(20), NA)
mylist <- list(x = x, y = y, z = z)
## vectors
reveal(x)
reveal(1:10)
## named or unnamed list
reveal(mylist)
set.seed(42)
mydf <- cbind(group = rep(letters[1:3], 4),
  setNames(as.data.frame(replicate(c(rnorm(11), NA), n = 3)), letters[24:26]))
## data frames
reveal(mydf)
## data frames by group
reveal(mydf, by = "group")
```

---

reveal_methods	<i>reveals little helper</i>
----------------	------------------------------

---

**Description**

S3 generic and methods

**Usage**

```
revealEye(x, ...)

## S3 method for class 'list'
revealEye(x, by, dec, funs, ...)

## S3 method for class 'numeric'
revealEye(x, dec, funs, ...)

## S3 method for class 'data.frame'
revealEye(x, dec, funs, ...)

## Default S3 method:
revealEye(x, dec, funs, ...)
```

**Arguments**

x	data frame, numeric vector, or list of numeric vectors
...	further arguments passed to methods
by	character vector with the names of the columns. Can be several variables!
dec	how many decimals are displayed
funs	not really meant to be used at the moment - change the Summarizing functions with a named(!) list of functions

**Value**

data frame

**See Also**

Other revealer: [reveal\\_split\(\)](#), [reveal\(\)](#)

---

snellen_steps	<i>Convert plus minus entries</i>
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---

**Description**

used in conversion method for class snellen

- Removing "plus" and "minus" from snellen notation
  - if entry -1 to +3 : take same Snellen value
  - if <= -2 : take Snellen value one line below
  - if >+3 (unlikely, but unfortunately not impossible): Snellen value one line above

**Usage**

```
snellensteps(y)
```

**Arguments**

`y` Vector with VA entries of class snellen - needs to be in format xx/yy

**Value**

character vector of Snellen entries

**snellen\_steps**

Snellen are unfortunately often entered with "+/-", which is a violation of a psychophysical method designed to assign one unambiguous value to visual acuity, with non-arbitrary thresholds based on psychometric functions. Therefore, transforming "+/-" notation to actual results is in itself problematic and the below suggestion to convert it will remain an approximation to the most likely "true" result. Even more so, as the given conditions should work for charts with 4 or 5 optotypes in a line, and visual acuity is not always tested on such charts. Yet, I believe that the approach is still better than just omitting the letters or (worse) assigning a missing value to those entries.

**See Also**

[https://en.wikipedia.org/wiki/Psychometric\\_function](https://en.wikipedia.org/wiki/Psychometric_function)

Other VA converter: [va\\_dissect\(\)](#), [va\\_methods](#), [va\(\)](#), [which\\_va\(\)](#)

---

va *Visual acuity notation conversion*

---

### Description

Cleans and converts visual acuity notations (classes) between Snellen (decimal, meter and feet), ETDRS, and logMAR. `va` detects the VA class and will convert to logMAR as default.

### Usage

```
va(
  x,
  to = "logmar",
  type = NULL,
  from_logmar = TRUE,
  logmarstep = FALSE,
  mixed = FALSE
)
```

### Arguments

<code>x</code>	Vector with visual acuity entries. Must be atomic. Snellen fractions need to be entered with "/"
<code>to</code>	To which class to convert. "etdrs", "logmar" or "snellen" - any case allowed
<code>type</code>	To which Snellen notation to convert: "m", "dec" or "ft"
<code>from_logmar</code>	chose logmar when guessing between two notations (logmar vs. snellen decimal or logmar vs. etdrs)
<code>logmarstep</code>	how +/- entries are evaluated. FALSE: increase/decrease Snellen fractions by lines. TRUE: plus/minus entries equivalent to 0.02 logmar or 1 ETDRS letter
<code>mixed</code>	TRUE Elements will be converted one by one. Most plausibility checks will be overruled!

### Details

Each class can be converted from one to another, and `va()` converts to logMAR by default. In case of ambiguous detection, logMAR is selected as default, or the other alternative is selected with `from_logmar = FALSE`.

### Value

vector of `va` class. See also "VA classes"



### VA conversion

- **logMAR to ETDRS:** logMAR rounded to the first digit and converted with the chart.
- **Snellen to logMAR:**  $\text{logMAR} = -1 * \log_{10}(\text{snellen\_frac})$
- **Snellen to ETDRS:**  $\text{ETDRS} = 85 + 50 * \log_{10}(\text{snellen\_frac})$  [Gregori et al.](#)
- **ETDRS to logMAR:**  $\text{logMAR} = -0.02 * \text{etdrs} + 1.7$  [Beck et al.](#)
- **Hand movements and counting fingers** are converted following [Schulze-Bonsel et al.](#)
- **(No) light perception** are converted following the suggestions by [Michael Bach](#)
- **To Snellen:** Although there seems to be no good statistical reason to convert back to Snellen, it is a very natural thing to eye specialists to think in Snellen. A conversion to snellen gives a good gauge of how the visual acuity for the patients are. However, back-conversion should not be considered an exact science and any attempt to use formulas will result in very weird Snellen values that have no correspondence to common charts. Therefore, Snellen matching the nearest ETDRS and logMAR value in the [va\\_chart](#) are used.

### Accepted VA formats

- Snellen fractions (meter/ feet) need to be entered as fraction with "/".
- **when converting to ETDRS or logMAR:** any fraction is allowed , e.g. 3/60 and 2/200 will also be recognized.
- **When converting between Snellen fractions:** *has to be either 6/ or 20/*. Other fractions will not be recognized - see "Examples"
- ETDRS must be integer-equivalent between 0 and 100 (integer equivalent means, it can also be a character vector)
- logMAR must be between -0.3 and 3.0
- Qualitative must be either of PL, LP, NLP, NPL, HM, CF (any case allowed)
- Any element which is not recognized will be converted to NA
- Vectors containing several notations ("mixed") are guessed and converted element by element with [which\\_va\\_dissect](#) and [va\\_dissect](#)

### VA detection

- Internally done with [which\\_va\(\)](#) based on the following rules
- if x integer and  $3 < x \leq 100$ : etdrs
- if x integer and  $0 \leq x \leq 3$ : logmar, but you can choose etdrs
- if x numeric and  $-0.3 \leq x \leq 3$ : logmar
- if x numeric and all x in  $\text{intersection}(\text{va\_chart}\$\text{logMAR}, \text{va\_chart}\$\text{snellen\_dec})$ : logmar, but you can choose snellen
- *non-mixed class*: if all x in  $\text{va\_chart}\$\text{snellen\_dec}$ : snellen
- *mixed class* ([which\\_va\\_dissect](#)): snellen\_dec not supported.
- if character and format x/y: snellen (fraction)
- if one of "CF", "HM", "LP", "PL", "NLP", or "NPL": quali
- if numeric x beyond the ranges from above: NA

- Any other string or NA: NA

Detection and conversion is on a vector as a whole by [which\\_va\(\)](#). If a "mixed" VA notation is found, [which\\_va\\_dissect\(\)](#) and [va\\_dissect\(\)](#) will be called instead for each VA vector element individually.

### Problematic cases

There can be ambiguous cases for detection (detection defaults to logmar): x is one of 0,1,2,3 - This can be ETDRS and logMAR. x is one of c(1.5, 1, 0.8, 0.5, 0.4, 0.3, 0.2, 0.1, 0) - This can be snellen decimal or logMAR.

**snellen decimals** are a particular challenge and va may wrongly assign logMAR - this could happen if there are unusual snellen decimal values in the data which are not part of [va\\_chart](#). E.g., check the values with `unique(x)`.

### Snellen "+/-" entries

By default, plus/minus entries are evaluated as intended by the test design: Snellen fractions increase/decrease only by lines.

- if entry -1 to +3 : take same Snellen value
- if <= -2 : take Snellen value one line below
- if >+3 (unlikely, but unfortunately not impossible):

If `logmarstep = TRUE`, each snellen optotype will be considered equivalent to 0.02 logmar or 1 ETDRS letter (assuming 5 letters in a row in a chart)

### VA cleaning

For more details see [clean\\_va\(\)](#)

1. NA is assigned to strings such as ".", "", "n/a" or " "
2. notation for qualitative entries is simplified.

### VA classes

`convert_VA` returns a vector of three classes:

1. va
2. One of snellen, logmar, etdrs or quali.
3. Either of character (for Snellen and qualitative), numeric (for logMAR), or integer (for ETDRS).

### See Also

Other VA converter: [snellen\\_steps](#), [va\\_dissect\(\)](#), [va\\_methods](#), [which\\_va\(\)](#)

Other VA cleaner: [clean\\_va\(\)](#)

**Examples**

```
## will automatically detect VA class and convert to logMAR by default
## ETDRS letters
x <- c(23, 56, 74, 58)
va(x)

## ... or convert to snellen
va(x, to = "snellen")

## snellen, mixed with categories. Also dealing with those "plus/minus" entries
va(c("NLP", "NPL", "PL", "LP", "HM", "CF", "6/60", "20/200", "6/9",
     "20/40", "20/40+3", "20/50-2"))

## A mix of notations is also possible
x <- c("NLP", "0.8", "34", "3/60", "2/200", "20/40+3", "20/50-2")
va(x)

## Any fraction is possible, and empty values
x <- c("CF", "3/60", "2/200", "", "20/40+3", ".", " ")
va(x)

## but this not any fraction when converting from one class to the other
x <- c("3/60", "2/200", "6/60", "20/200", "6/9")
va(x, to="snellen", type = "m")
```

---

va\_chart

*Visual acuity conversion chart*


---

**Description**

Conversion between snellen, logMAR and ETDRS. Snellen feet, meter and decimal supported. Three qualitative common vision measures included (light perception, hand movement and counting fingers). Further details for conversion used can be found in [va](#) and [va\\_methods](#)

**Usage**

```
data("va_chart")
```

**Format**

A data frame with 29 rows and 5 variables:

**snellen\_ft** snellen VA in feet  
**snellen\_m** snellen VA in meter  
**snellen\_dec** decimal snellen VA  
**logmar** logMAR VA  
**etdrs** VA in ETDRS letters  
**quali** VA categories

**See Also**

- This chart and VA conversion formulas are based on charts in [Holladay et al.](#), [Beck et al.](#), and [Gregori et al.](#).

Categories **(no) light perception, counting fingers** and **hand movements** are converted following [Schulze-Bonsel et al.](#) and [Michael Bach's suggestions](#)

---

va\_methods

*VA conversion methods*


---

**Description**

S3 methods for VA conversion

**Usage**

```
convertVA(x, to, ...)

## S3 method for class 'quali'
convertVA(x, to, snellnot, ...)

## S3 method for class 'snellen'
convertVA(x, to, snellnot, logmarstep, ...)

## S3 method for class 'logmar'
convertVA(x, to, snellnot, ...)

## S3 method for class 'etdrs'
convertVA(x, to, snellnot, ...)

## Default S3 method:
convertVA(x, to, snellnot, ...)
```

**Arguments**

x	vector of visual acuities
to	to which VA class to convert
...	further arguments passed to methods
snellnot	which snellen notation. One of "ft", "m" or "dec"
logmarstep	how plus/minus entries are evaluated. Default to increase/decrease snellen fractions by lines. If TRUE, each snellen optotype will be considered equivalent to 0.02 logmar or 1 ETDRS letter (assuming 5 letters in a row in a chart)

**Details**

VA can be snellen feet/meter/decimal, logMAR, ETDRS, or "qualitative" (Counting fingers, etc.)

- Snellen fractions need to be either form 6/x or 20/x
- ETDRS must be between 0 and 100
- logMAR must be between -0.3 and 3.0
- Qualitative must be PL, LP, NLP, NPL, HM, CF (any case allowed)

Any element which is not recognized will be converted to NA

For other conversion rules see [va](#)

**Value**

vector with visual acuity of class va. See also "VA classes"

**Conversion**

Although there seems to be no good statistical reason to convert back to Snellen, it is a very natural thing to eye specialists to think in Snellen. A conversion to snellen gives a good gauge of how the visual acuity for the patients are. However, back-conversion should not be considered an exact science and any attempt to use formulas will result in very weird Snellen values that have no correspondence to common charts. Therefore, Snellen matching the nearest ETDRS and logMAR value in the [va\\_chart](#) are used.

Further:

- logMAR to ETDRS: logMAR rounded to the first digit and converted with the chart.
- Snellen to logMAR:  $\text{logMAR} = -1 * \log_{10}(\text{snellen\_frac})$
- Snellen to ETDRS:  $\text{ETDRS} = 85 + 50 * \log_{10}(\text{snellen\_frac})$  [Gregori et al.](#)
- ETDRS to logMAR:  $\text{logMAR} = -0.02 * \text{etdrs} + 1.7$  [Beck et al.](#)

**VA classes**

convert\_VA returns a vector of three classes:

1. va
2. One of snellen, logmar, etdrs or quali.
3. Either of character (for Snellen and qualitative), numeric (for logMAR), or integer (for ETDRS).

**See Also**

Other VA converter: [snellen\\_steps](#), [va\\_dissect\(\)](#), [va\(\)](#), [which\\_va\(\)](#)

# Index

## \*Topic **datasets**

- amd, [3](#)
- va\_chart, [19](#)
  
- age, [2](#)
- amd, [3](#)
  
- blink, [4](#)
  
- clean\_va, [6](#), [18](#)
- clean\_va(), [18](#)
- convert\_NLP (clean\_va), [6](#)
- convert\_NLP(), [6](#)
- convertVA (va\_methods), [20](#)
  
- eyes, [4](#), [5](#), [7](#)
- eyes\_to\_string, [7](#), [8](#)
- eyestr (eyes), [7](#)
  
- getElement, [12](#)
  
- hyperop, [9](#)
  
- isNAstring (clean\_va), [6](#)
- isNAstring(), [6](#)
  
- myop, [4](#), [10](#)
- myop(), [9](#)
- myop\_pivot, [11](#)
- myop\_rename, [11](#)
- myopic (myop), [10](#)
- myopizer, [11](#)
  
- print.blink (print\_methods), [11](#)
- print.default, [12](#)
- print.etdrs (print\_methods), [11](#)
- print.logmar (print\_methods), [11](#)
- print.snellen (print\_methods), [11](#)
- print\_methods, [11](#)
  
- recodeye, [12](#)
  
- remCols, [5](#)
- reveal, [4](#), [13](#), [15](#)
- reveal\_methods, [13](#), [14](#)
- reveal\_split, [13](#), [15](#)
- revealEye (reveal\_methods), [14](#)
  
- set\_codes, [10](#), [12](#)
- snellen\_steps, [15](#), [18](#), [21](#)
- snellensteps (snellen\_steps), [15](#)
- sort\_substr, [12](#)
- sort\_substr(), [11](#)
- str\_func\_facs, [12](#)
  
- tibble::tibble, [9](#), [11](#)
- tidyr::pivot\_longer(), [9](#)
- tidyr::pivot\_wider(), [9](#)
  
- va, [7](#), [15](#), [16](#), [19](#), [21](#)
- va(), [4](#)
- va\_chart, [17](#), [18](#), [19](#), [21](#)
- va\_dissect, [15](#), [17](#), [18](#), [21](#)
- va\_dissect(), [18](#)
- va\_methods, [15](#), [18](#), [19](#), [20](#)
  
- which\_va, [15](#), [18](#), [21](#)
- which\_va(), [17](#), [18](#)
- which\_va\_dissect, [17](#)
- which\_va\_dissect(), [18](#)