

# Package ‘unheadr’

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**Type** Package

**Title** Handle Data with Messy Header Rows and Broken Values

**Version** 0.3.1

**Depends** R (>= 3.1.0)

**Description** Verb-like functions to work with messy data, often derived from spreadsheets or parsed PDF tables. Includes functions for unwrapping values broken up across rows, relocating embedded grouping values, and to annotate meaningful formatting in spreadsheet files.

**License** MIT + file LICENSE

**Encoding** UTF-8

**LazyData** true

**Imports** dplyr (>= 0.8.4), rlang (>= 0.2.1), forcats, stringr, tidyr, magrittr, tidyxl, readxl, tibble

**RoxygenNote** 7.1.1

**Suggests** knitr, rmarkdown, testthat (>= 2.1.0), covr

**VignetteBuilder** knitr

**URL** <https://github.com/luisDVA/unheadr>, <https://unheadr.liomys.mx/>

**BugReports** <https://github.com/luisDVA/unheadr/issues>

**NeedsCompilation** no

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**Repository** CRAN

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## R topics documented:

annotate_mf . . . . .	2
annotate_mf_all . . . . .	3

AOEunits . . . . .	4
AOEunits_raw . . . . .	5
boutiques.xlsx . . . . .	5
dog_test.xlsx . . . . .	6
mash_colnames . . . . .	6
primates2017 . . . . .	8
primates2017_broken . . . . .	9
primates2017_wrapped . . . . .	9
regex_valign . . . . .	10
unbreak_rows . . . . .	11
unbreak_vals . . . . .	12
untangle2 . . . . .	13
unwrap_cols . . . . .	14
%>% . . . . .	14

<b>Index</b>	<b>15</b>
--------------	-----------

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annotate_mf	<i>Annotate meaningful formatting</i>
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## Description

Turns cell formatting into annotations for values in the target row.

## Usage

```
annotate_mf(xlfilepath, orig, new)
```

## Arguments

xlfilepath	Path to a single-sheet spreadsheet file (xls or.xlsx).
orig	Variable to annotate formatting in.
new	Name of new variable with cell formatting pasted as a string.

## Details

At this point, seven popular approaches for meaningful formatting (bold, colored text, italic, strikethrough, underline, double underline, and cell highlighting) are hardcoded in the function. `sheets`, `skip`, and `range` arguments for spreadsheet input are not supported. The hex8 code of the fill color used for text color and cell highlighting is also appended in the output. Ensure the data in the spreadsheet are rectangular before running.

## Value

A tibble with a new column with meaningful formatting embedded as text.

## Examples

```
example_spreadsheet <- system.file("extdata/dog_test.xlsx", package = "unheadr")
annotate_mf(example_spreadsheet, orig = Task, new = Task_annotated)
```

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annotate_mf_all	<i>Annotate meaningful formatting for all cells</i>
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## Description

Turns cell formatting into annotations for all values across all variables.

## Usage

```
annotate_mf_all(xlfilepath)
```

## Arguments

xlfilepath      Path to a single-sheet spreadsheet file (xls or xlsx).

## Details

At this point, seven popular approaches for meaningful formatting (bold, colored text, italic, strikethrough, underline, double underline, and cell highlighting) are hardcoded in the function. `sheets`, `skip`, and `range` arguments for spreadsheet input are not supported. The hex8 code of the fill color used for text color and cell highlighting is also appended in the output. Ensure the data in the spreadsheet are rectangular before running.

## Value

A tibble with meaningful formatting embedded as text for all rows and columns.

## Examples

```
example_spreadsheet <- system.file("extdata/boutiques.xlsx", package = "unheadr")
annotate_mf_all(example_spreadsheet)
```

## Description

A dataset with the numerical values that determine the behavior and performance of selected military units available in AoE2:DE (July 2020 Game Update).

## Usage

AOEunits

## Format

A data frame with 128 observations of 19 variables:

**unit** Unit name  
**building** Building in which each unit is trained  
**type** Unit class  
**age** Age at which the unit becomes trainable  
**cost\_wood** Unit cost in Wood  
**cost\_food** Unit cost in Food  
**cost\_gold** Unit cost in Gold  
**build\_time** Training time in seconds  
**rate\_of\_fire** Attack speed  
**attack\_delay** Retasking time  
**movement\_speed** Travel speed on land  
**line\_of\_sight** Vision over the surrounding area  
**hit\_points** Unit health  
**min\_range** Minimum attacking range for ranged units  
**range** Maximum attacking range for ranged units  
**damage** Damage inflicted per attack  
**accuracy** Chance that an attack will be on target  
**melee\_armor** Armor against melee attacks  
**pierce\_armor** Armor against projectiles

## Source

Age of Empires II. Copyright Microsoft Corporation. This dataset was created under Microsoft's "Game Content Usage Rules" <https://www.xbox.com/en-us/developers/rules> using assets from Age of Empires II, and it is not endorsed by or affiliated with Microsoft. All information shown is an interpretation of data collected in-game with no guarantee on the accuracy of any of the data presented.

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AOEunits_raw	<i>Statistics for game units in Age of Empires II: Definitive Edition in a messy presentation</i>
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### Description

A messy version of the [AOEunits](#) dataset, meant for demonstrating data cleaning functions.

### Usage

AOEunits\_raw

### Format

A data frame with 139 observations of 15 variables. See [AOEunits](#) for variable descriptions.

### Source

Age of Empires II. Copyright Microsoft Corporation. This dataset was created under Microsoft's "Game Content Usage Rules" <https://www.xbox.com/en-us/developers/rules> using assets from Age of Empires II, and it is not endorsed by or affiliated with Microsoft. All information shown is an interpretation of data collected in-game with no guarantee on the accuracy of any of the data presented.

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boutiques.xlsx	<i>boutiques.xlsx spreadsheet</i>
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### Description

Open XML Format Spreadsheet with 1 sheet, 6 columns, and 8 rows. Toy dataset with Q1 profits for different store locations. Additional information is encoded as meaningful formatting. Bold indicates losses (negative values), colors indicate continent, and italic indicates a second location in the same city.

### Details

This data is used in the example for `annotate_mf_all()`.

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dog_test.xlsx	<i>dog_test.xlsx spreadsheet</i>
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---

### Description

Open XML Format Spreadsheet with 1 sheet, 2 columns, and 12 rows. Items describe various tasks or behaviors that dogs can be evaluated on, assigned into three categories which appear along with their average scores as embedded subheaders with meaningful formatting.

### Details

This data is used in the example for `annotate_mf()`.

### Source

Items are modified from the checklist written by Junior Watson.

### References

<http://www.dogtrainingbasics.com/checklist-well-behaved-dog/>

---

mash_colnames	<i>Make many header rows into column names</i>
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### Description

Make many header rows into column names

### Usage

```
mash_colnames(
  df,
  n_name_rows,
  keep_names = TRUE,
  sliding_headers = FALSE,
  sep = "_"
)
```

### Arguments

df	A data.frame or tibble object in which the names are broken up across the top <i>n</i> rows.
n_name_rows	Number of rows at the top of the data to be used to create the new variable (column) names. Must be $\geq 1$ .

keep_names	If TRUE, existing names will be included in building the new variable names. Defaults to TRUE.
sliding_headers	If TRUE, empty values in the first (topmost) header header row be filled column-wise. Defaults to FALSE. See details.
sep	Character string to separate the unified values (default is underscore).

### Details

Tables are often shared with the column names broken up across the first few rows. This function takes the number of rows at the top of a table that hold the broken up names and whether or not to include the names, and mashes the values column-wise into a single string for each column. The `keep_names` argument can be helpful for tables we imported using a `skip` argument. If `keep_names` is set to FALSE, adjust the value of `n_name_rows` accordingly.

This function will throw a warning when possible NA values end up in the variable names. `sliding_headers` can be used for tables with ragged names in which not every column has a value in the very first row. In these cases attribution by adjacency is assumed, and when `sliding_headers` is set to TRUE the names in the topmost row are filled row-wise. This can be useful for tables reporting survey data or experimental designs in an untidy manner.

### Value

The original data frame, but with new column names and without the top `n` rows that held the broken up names.

### Author(s)

This function was originally contributed by Jarrett Byrnes through a GitHub issue.

### Examples

```
babies <-
  data.frame(
    stringsAsFactors = FALSE,
    Baby = c(NA, NA, "Angie", "Yean", "Pierre"),
    Age = c("in", "months", "11", "9", "7"),
    Weight = c("kg", NA, "2", "3", "4"),
    Ward = c(NA, NA, "A", "B", "C")
  )
# Including the object names
mash_colnames(babies, n_name_rows = 2, keep_names = TRUE)

babies_skip <-
  data.frame(
    stringsAsFactors = FALSE,
    X1 = c("Baby", NA, NA, "Jennie", "Yean", "Pierre"),
    X2 = c("Age", "in", "months", "11", "9", "7"),
    X3 = c("Hospital", NA, NA, "A", "B", "A")
  )
#' # Discarding the automatically-generated names (X1, X2, etc...)
```

```

mash_colnames(babies_skip, n_name_rows = 3, keep_names = FALSE)

fish_experiment <-
  data.frame(
    stringsAsFactors = FALSE,
    X1 = c("Sample", NA, "Pacific", "Atlantic", "Freshwater"),
    X2 = c("Larvae", "Control", "12", "11", "10"),
    X3 = c(NA, "Low Dose", "11", "12", "8"),
    X4 = c(NA, "High Dose", "8", "7", "9"),
    X5 = c("Adult", "Control", "13", "13", "8"),
    X6 = c(NA, "Low Dose", "13", "12", "7"),
    X7 = c(NA, "High Dose", "10", "10", "9")
  )
# Ragged names
mash_colnames(fish_experiment,
  n_name_rows = 2,
  keep_names = FALSE, sliding_headers = TRUE
)

```

---

primates2017

*Comparative data for 54 species of primates*

---

### Description

A dataset with embedded subheaders.

### Usage

```
primates2017
```

### Format

A data frame with 69 rows and 4 variables:

**scientific\_name** scientific names, with geographic region and taxonomic family embedded as sub-headers.

**common\_name** vernacular name

**red\_list\_status** IUCN Red List Status in January 2017

**mass\_kg** mean body mass in kilograms

### Source

Estrada, Alejandro, et al. "Impending extinction crisis of the world's primates: Why primates matter." *Science Advances* 3.1 (2017): e1600946. doi: [10.1126/sciadv.1600946](https://doi.org/10.1126/sciadv.1600946)



---

primates2017\_broken    *Comparative data for 16 species of primates with some broken values*

---

**Description**

A dataset with embedded subheaders and some values (T. obscurus, T. leucocephalus and N. bengalensis) in the scientific\_names variable broken up across two rows (typically done to fit the content in a table).

**Usage**

primates2017\_broken

**Format**

A data frame with 19 rows and 4 variables:

**scientific\_name** scientific names, with embedded subheaders for geographic region and taxonomic family and broken values

**common\_name** vernacular name

**red\_list\_status** IUCN Red List Status in January 2017

**mass\_kg** mean body mass in kilograms

**Source**

Estrada, Alejandro, et al. "Impending extinction crisis of the world's primates: Why primates matter." Science Advances 3.1 (2017): e1600946. doi: [10.1126/sciadv.1600946](https://doi.org/10.1126/sciadv.1600946)

---

primates2017\_wrapped    *Comparative data for two species of primates*

---

**Description**

A dataset in which the elements for some of the values are in separate rows'

**Usage**

primates2017\_wrapped

**Format**

A data frame with 9 rows and 6 variables:

**scientific\_name** scientific names, see reference

**common\_name** vernacular name

**habitat** habitat types listed in the IUCN Red List assessments

**red\_list\_status** IUCN Red List Status in January 2017

**mass\_kg** mean body mass in kilograms

**country** Countries where the species is present, from IUCN Red List assessments

**Source**

Estrada, Alejandro, et al. "Impending extinction crisis of the world's primates: Why primates matter." *Science Advances* 3.1 (2017): e1600946. doi: [10.1126/sciadv.1600946](https://doi.org/10.1126/sciadv.1600946)

---

regex_valign	<i>Vertical character string alignment through regular expressions</i>
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---

**Description**

Aligning strings with regex.

**Usage**

```
regex_valign(stringvec, regex_ai, sep_str = "")
```

**Arguments**

**stringvec** A character vector with one element for each line.

**regex\_ai** A regular expression matching the position for alignment.

**sep\_str** Optional character vector that will be inserted at the positions matched by the regular expression.

**Details**

Written mainly for reading fixed width files, text, or tables parsed from PDFs.

**Value**

A character vector with one element for each line, with padding inserted at the matched positions so that elements are vertically aligned across lines.

**See Also**

This function is based loosely on `textutils::valign()`.

**Examples**

```

guests <-
  unlist(strsplit(c("6          COAHUILA          20/03/2020
7          COAHUILA          20/03/2020
18 BAJA CALIFORNIA      16/03/2020
109       CDMX          12/03/2020
1230     QUERETARO      21/03/2020"), "\n"))

# align at first uppercase word boundary , inserting a separator
regex_valign(guests, "\\b(?=[A-Z])", " - ")
# align dates at end of string
regex_valign(guests, "\\b(?=[0-9]{2}[\\/]]{1}[0-9]{2}[\\/]]{1}[0-9]{4}$)")

```

---

unbreak_rows	<i>Merge rows up</i>
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---

**Description**

Merge rows up

**Usage**

```
unbreak_rows(df, regex, ogcol, sep = " ")
```

**Arguments**

df	A data frame with at least two contiguous rows to be merged.
regex	A regular expression to identify sets of rows to be merged, meant for the leading of the two contiguous rows.
ogcol	Variable with the text strings to match.
sep	Character string to separate the unified values (default is space).

**Details**

This function recodes empty strings ("") to NA for smoother pattern matching.

**Value**

A tibble or data frame with merged rows. Values of the lagging rows are pasted onto the values in the leading row, whitespace is squished, and the lagging row is dropped.

**Examples**

```

bball <-
  data.frame(
    stringsAsFactors = FALSE,
    v1 = c(
      "Player", NA, "Steve McDichael", "Dean Wesrey",
      "Karl Dandleton"
    ),
    v2 = c("Most points", "in a game", "55", "43", "41"),
    v3 = c("Season", "(year ending)", "2001", "2000", "2010")
  )
unbreak_rows(bball, "Most", v2)

```

---

unbreak_vals	<i>Unbreak values using regex to match the lagging half of the broken value</i>
--------------	---

---

**Description**

Unbreak values using regex to match the lagging half of the broken value

**Usage**

```
unbreak_vals(df, regex, ogcol, newcol, sep = " ", slice_groups)
```

**Arguments**

df	A data frame with one or more values within a variable broken up across two rows.
regex	Regular expression for matching the lagging half of the broken values.
ogcol	Variable to unbreak.
newcol	Name of the new variable with the unified values.
sep	Character string to separate the unified values (default is space).
slice_groups	Deprecated. See details and Package News.

**Details**

This function is limited to quite specific cases, but useful when dealing with tables that contain, for example, scientific names broken across two rows. For unwrapping values, see [unwrap\\_cols](#).

**Value**

A tibble with 'unbroken' values. The variable that originally contained the broken values gets dropped, and the new variable with the unified values is placed as the first column. The `slice_groups` argument is now deprecated; the extra rows and the variable with broken values will be dropped.

**Examples**

```
data(primates2017_broken)
# regex matches strings starting in lowercase (broken species epithets)
unbreak_vals(primates2017_broken, "^[a-z]", scientific_name, sciname_new)
```

---

untangle2

*Rectangling embedded subheaders*


---

**Description**

Rectangling embedded subheaders

**Usage**

```
untangle2(df, regex, orig, new)
```

**Arguments**

df	A data frame with embedded subheaders.
regex	Regular expression to match the subheaders.
orig	Variable containing the extraneous subheaders.
new	Name of variable that will contain the group values.

**Details**

Special thanks to Jenny Bryan for fixing the initial tidyeval code and overall function structure.

**Value**

A tibble without the matched subheaders and a new variable containing the grouping data.

**Examples**

```
data(primates2017)
# put taxonomic family in its own variable (matches the suffix "DAE")
untangle2(primates2017, "DAE$", scientific_name, family)
# put geographic regions in their own variable (matching them all by name)
untangle2(
  primates2017, "Asia|Madagascar|Mainland Africa|Neotropics",
  scientific_name, family
)
# with magrittr pipes (re-exported in this package)
primates2017 %>%
  untangle2("DAE$", scientific_name, family) %>%
  untangle2(
    "Asia|Madagascar|Mainland Africa|Neotropics",
    scientific_name, region
  )
```

---

unwrap_cols	<i>Unwrap values and clean up NAs used as padding</i>
-------------	---

---

**Description**

Unwrap values and clean up NAs used as padding

**Usage**

```
unwrap_cols(df, groupingVar, separator)
```

**Arguments**

df	A data frame with wrapped values and an inconsistent number of NA values used to as within-group padding.
groupingVar	Name of the variable describing the observational units.
separator	Character string defining the separator that will delimit the elements of the unwrapped value.

**Details**

This is roughly the opposite of `tidyr::separate_rows()`.

**Value**

A summarized tibble. Order is preserved in the grouping variable by making it a factor.

**Examples**

```
data(primates2017_wrapped)
# using commas to separate elements
unwrap_cols(primates2017_wrapped, scientific_name, ", ")

# separating with semicolons
df <- data.frame(
  ounits = c("A", NA, "B", "C", "D", NA),
  vals = c(1, 2, 2, 3, 1, 3)
)
unwrap_cols(df, ounits, ";")
```

---

%>%	<i>re-export magrittr pipe operator</i>
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**Description**

re-export magrittr pipe operator

# Index

## \* datasets

- AOEunits, [4](#)
- AOEunits\_raw, [5](#)
- primates2017, [8](#)
- primates2017\_broken, [9](#)
- primates2017\_wrapped, [9](#)

[%>%](#), [14](#)

[annotate\\_mf](#), [2](#)

[annotate\\_mf\\_all](#), [3](#)

[AOEunits](#), [4](#), [5](#)

[AOEunits\\_raw](#), [5](#)

[boutiques.xlsx](#), [5](#)

[dog\\_test.xlsx](#), [6](#)

[mash\\_colnames](#), [6](#)

[primates2017](#), [8](#)

[primates2017\\_broken](#), [9](#)

[primates2017\\_wrapped](#), [9](#)

[regex\\_valign](#), [10](#)

[unbreak\\_rows](#), [11](#)

[unbreak\\_vals](#), [12](#)

[untangle2](#), [13](#)

[unwrap\\_cols](#), [12](#), [14](#)