

# Package ‘REDCapCAST’

July 4, 2023

**Title** REDCap Castellated Data Handling

**Version** 23.6.2

**Description** Forked from 'REDCapRITS' by Paul Egeler and Spectrum Health.

See <<https://github.com/SpectrumHealthResearch/REDCapRITS>>.

Handles castellated datasets from 'REDCap' projects with repeating instruments.

Assists in casting tidy tables from raw 'REDCap' data exports for each repeated instrument. Keeps a focused data export approach, by allowing to only export required data from the database.

'REDCap' (Research Electronic Data Capture) is a secure, web-based software platform designed to support data capture for research studies, providing

1) an intuitive interface for validated data capture; 2) audit trails for tracking data manipulation and export procedures; 3) automated export procedures for seamless data downloads to common statistical packages; and 4) procedures for data integration and interoperability with external sources (Harris et al (2009) <[doi:10.1016/j.jbi.2008.08.010](https://doi.org/10.1016/j.jbi.2008.08.010)>; Harris et al (2019) <[doi:10.1016/j.jbi.2019.103208](https://doi.org/10.1016/j.jbi.2019.103208)>).

**Depends** R (>= 3.4.0)

**Suggests** spelling, RCurl, httr, jsonlite, testthat, Hmisc, readr, covr, knitr, rmarkdown, gt, keyring

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.2.3

**URL** <https://github.com/agdamsbo/REDCapCAST>,  
<https://agdamsbo.github.io/REDCapCAST/>

**BugReports** <https://github.com/agdamsbo/REDCapCAST/issues>

**Imports** dplyr, REDCapR, tidyr, tidyselct

**Collate** 'utils.r' 'process\_user\_input.r' 'REDCap\_split.r' 'ds2dd.R'  
'read\_redcap\_tables.R' 'redcap\_wider.R' 'redcapcast\_data.R'  
'redcapcast\_meta.R'

**Language** en-US

**VignetteBuilder** knitr**NeedsCompilation** no**Author** Andreas Gammelgaard Damsbo [aut, cre]  
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clean_redcap_name	<i>clean_redcap_name</i>
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### Description

Stepwise removal on non-alphanumeric characters, trailing white space, substitutes spaces for underscores and converts to lower case. Trying to make up for different naming conventions.

### Usage

```
clean_redcap_name(x)
```

### Arguments

x                   vector or data frame for cleaning

### Value

vector or data frame, same format as input

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d2w *Convert single digits to words*

---

**Description**

Convert single digits to words

**Usage**

```
d2w(x, lang = "en", neutrum = FALSE, everything = FALSE)
```

**Arguments**

x	data. Handle vectors, data.frames and lists
lang	language. Danish (da) and English (en), Default is "en"
neutrum	for numbers depending on counted word
everything	flag to also split numbers >9 to single digits

**Value**

returns characters in same format as input

**Examples**

```
d2w(c(2:8,21))
d2w(data.frame(2:7,3:8,1),lang="da",neutrum=TRUE)

## If everything=T, also larger numbers are reduced.
## Elements in the list are same length as input
d2w(list(2:8,c(2,6,4,23),2), everything=TRUE)
```

---

ds2dd *Data set to data dictionary function*

---

**Description**

Migrated from stRoke ds2dd(). Fits better with the functionality of 'REDCapCAST'

**Usage**

```
ds2dd(
  ds,
  record.id = "record_id",
  form.name = "basis",
  field.type = "text",
  field.label = NULL,
  include.column.names = FALSE,
  metadata = names(redcapcast_meta)
)
```

**Arguments**

ds	data set
record.id	name or column number of id variable, moved to first row of data dictionary, character of integer. Default is "record_id".
form.name	vector of form names, character string, length 1 or length equal to number of variables. Default is "basis".
field.type	vector of field types, character string, length 1 or length equal to number of variables. Default is "text".
field.label	vector of form names, character string, length 1 or length equal to number of variables. Default is NULL and is then identical to field names.
include.column.names	Flag to give detailed output including new column names for original data set for upload.
metadata	Metadata column names. Default is the included REDCapCAST::redcapcast_data.

**Value**

data.frame or list of data.frame and vector

**Examples**

```
redcapcast_data$record_id <- seq_len(nrow(redcapcast_data))
ds2dd(redcapcast_data, include.column.names=TRUE)
```

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focused\_metadata      *focused\_metadata*

---

**Description**

Extracts limited metadata for variables in a dataset

**Usage**

```
focused_metadata(metadata, vars_in_data)
```

**Arguments**

metadata      A dataframe containing metadata  
vars\_in\_data    Vector of variable names in the dataset

**Value**

A dataframe containing metadata for the variables in the dataset

---

match\_fields\_to\_form    *Match fields to forms*

---

**Description**

Match fields to forms

**Usage**

```
match_fields_to_form(metadata, vars_in_data)
```

**Arguments**

metadata      A data frame containing field names and form names  
vars\_in\_data    A character vector of variable names

**Value**

A data frame containing field names and form names

---

read\_redcap\_tables      *Download REDCap data*

---

**Description**

Implementation of REDCap\_split with a focused data acquisition approach using REDCapR::redcap\_read nad only downloading specified fields, forms and/or events using the built-in focused\_metadata including some clean-up. Works with longitudinal projects with repeating instruments.

**Usage**

```
read_redcap_tables(
  uri,
  token,
  records = NULL,
  fields = NULL,
  events = NULL,
  forms = NULL,
  raw_or_label = "label",
  split_forms = "all",
  generics = c("record_id", "redcap_event_name", "redcap_repeat_instrument",
    "redcap_repeat_instance")
)
```

**Arguments**

uri	REDCap database uri
token	API token
records	records to download
fields	fields to download
events	events to download
forms	forms to download
raw_or_label	raw or label tags
split_forms	Whether to split "repeating" or "all" forms, default is all.
generics	vector of auto-generated generic variable names to ignore when discarding empty rows

**Value**

list of instruments

**Examples**

```
# Examples will be provided later
```

---

redcapcast_data	<i>Data set for demonstration</i>
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**Description**

This is a small dataset from a REDCap database for demonstrational purposes. Contains only synthetic data.

**Usage**

```
data(redcapcast_data)
```

**Format**

A data frame with 22 variables:

**record\_id** ID, numeric  
**redcap\_event\_name** Event name, character  
**redcap\_repeat\_instrument** Repeat instrument, character  
**redcap\_repeat\_instance** Repeat instance, numeric  
**cpr** CPR number, character  
**inclusion** Inclusion date, date  
**dob** Date of birth, date  
**age** Age decimal, numeric  
**age\_integer** Age integer, numeric  
**sex** Legal sex, character  
**cohabitation** Cohabitation status, character  
**hypertension** Hypertension, character  
**diabetes** diabetes, character  
**region** region, character  
**baseline\_data\_start\_complete** Completed, character  
**mrs\_assessed** mRS Assessed, character  
**mrs\_date** Assesment date, date  
**mrs\_score** Score, numeric  
**mrs\_complete** Complete, numeric  
**event\_date** Event date, date  
**event\_type** Event type, character  
**new\_event\_complete** Completed, character

---

redcapcast\_meta

*REDCap metadata from data base*

---

**Description**

This metadata dataset from a REDCap database is for demonstrational purposes.

**Usage**

```
data(redcapcast_meta)
```

**Format**

A data frame with 22 variables:

**field\_name** field\_name, character

**form\_name** form\_name, character

**section\_header** section\_header, character

**field\_type** field\_type, character

**field\_label** field\_label, character

**select\_choices\_or\_calculations** select\_choices\_or\_calculations, character

**field\_note** field\_note, character

**text\_validation\_type\_or\_show\_slider\_number** text\_validation\_type\_or\_show\_slider\_number, character

**text\_validation\_min** text\_validation\_min, character

**text\_validation\_max** text\_validation\_max, character

**identifier** identifier, character

**branching\_logic** branching\_logic, character

**required\_field** required\_field, character

**custom\_alignment** custom\_alignment, character

**question\_number** question\_number, character

**matrix\_group\_name** matrix\_group\_name, character

**matrix\_ranking** matrix\_ranking, character

**field\_annotation** field\_annotation, character

---

REDCap\_split

*Split REDCap repeating instruments table into multiple tables*

---

**Description**

This will take output from a REDCap export and split it into a base table and child tables for each repeating instrument. Metadata is used to determine which fields should be included in each resultant table.

**Usage**

```
REDCap_split(
  records,
  metadata,
  primary_table_name = "",
  forms = c("repeating", "all")
)
```



**Arguments**

records	Exported project records. May be a data.frame, response, or character vector containing JSON from an API call.
metadata	Project metadata (the data dictionary). May be a data.frame, response, or character vector containing JSON from an API call.
primary_table_name	Name given to the list element for the primary output table (as described in <i>README.md</i> ). Ignored if forms = 'all'.
forms	Indicate whether to create separate tables for repeating instruments only or for all forms.

**Value**

A list of "data.frame"s. The number of tables will differ depending on the forms option selected.

- 'repeating': one base table and one or more tables for each repeating instrument.
- 'all': a data.frame for each instrument, regardless of whether it is a repeating instrument or not.

**Author(s)**

Paul W. Egeler, M.S., GStat

**Examples**

```
## Not run:
# Using an API call -----

library(RCurl)

# Get the records
records <- postForm(
  uri = api_url,      # Supply your site-specific URI
  token = api_token, # Supply your own API token
  content = 'record',
  format = 'json',
  returnFormat = 'json'
)

# Get the metadata
metadata <- postForm(
  uri = api_url,      # Supply your site-specific URI
  token = api_token, # Supply your own API token
  content = 'metadata',
  format = 'json'
)

# Convert exported JSON strings into a list of data.frames
REDCapRITS::REDCap_split(records, metadata)
```

```

# Using a raw data export -----
# Get the records
records <- read.csv("/path/to/data/ExampleProject_DATA_2018-06-03_1700.csv")

# Get the metadata
metadata <- read.csv(
"/path/to/data/ExampleProject_DataDictionary_2018-06-03.csv")

# Split the tables
REDCapRITS::REDCap_split(records, metadata)

# In conjunction with the R export script -----
# You must set the working directory first since the REDCap data export
# script contains relative file references.
old <- getwd()
setwd("/path/to/data/")

# Run the data export script supplied by REDCap.
# This will create a data.frame of your records called 'data'
source("ExampleProject_R_2018-06-03_1700.r")

# Get the metadata
metadata <- read.csv("ExampleProject_DataDictionary_2018-06-03.csv")

# Split the tables
REDCapRITS::REDCap_split(data, metadata)
setwd(old)

## End(Not run)

```

---

redcap\_wider

*Redcap Wider*


---

## Description

Converts a list of REDCap data frames from long to wide format. Handles longitudinal projects, but not yet repeated instruments.

## Usage

```

redcap_wider(
  list,
  event.glue = "{.value}_{redcap_event_name}",
  inst.glue = "{.value}_{redcap_repeat_instance}"
)

```

**Arguments**

`list` A list of data frames.  
`event.glue` A dplyr::glue string for repeated events naming  
`inst.glue` A dplyr::glue string for repeated instruments naming

**Value**

The list of data frames in wide format.

**Examples**

```
list <- list(data.frame(record_id = c(1,2,1,2),
  redcap_event_name = c("baseline", "baseline", "followup", "followup"),
  age = c(25,26,27,28)),
  data.frame(record_id = c(1,2),
  redcap_event_name = c("baseline", "baseline"),
  gender = c("male", "female")))
redcap_wider(list)
```

---

sanitize_split	<i>Sanitize list of data frames</i>
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---

**Description**

Removing empty rows

**Usage**

```
sanitize_split(
  l,
  generic.names = c("record_id", "redcap_event_name", "redcap_repeat_instrument",
    "redcap_repeat_instance")
)
```

**Arguments**

`l` A list of data frames.  
`generic.names` A vector of generic names to be excluded.

**Value**

A list of data frames with generic names excluded.

---

`split_non_repeating_forms`*Split a data frame into separate tables for each form*

---

**Description**

Split a data frame into separate tables for each form

**Usage**

```
split_non_repeating_forms(table, universal_fields, fields)
```

**Arguments**

<code>table</code>	A data frame
<code>universal_fields</code>	A character vector of fields that should be included in every table
<code>fields</code>	A two-column matrix containing the names of fields that should be included in each form

**Value**

A list of data frames, one for each non-repeating form

**Examples**

```
# Create a table
table <- data.frame(
  id = c(1, 2, 3, 4, 5),
  form_a_name = c("John", "Alice", "Bob", "Eve", "Mallory"),
  form_a_age = c(25, 30, 25, 15, 20),
  form_b_name = c("John", "Alice", "Bob", "Eve", "Mallory"),
  form_b_gender = c("M", "F", "M", "F", "F")
)

# Create the universal fields
universal_fields <- c("id")

# Create the fields
fields <- matrix(
  c("form_a_name", "form_a",
    "form_a_age", "form_a",
    "form_b_name", "form_b",
    "form_b_gender", "form_b"),
  ncol = 2, byrow = TRUE
)

# Split the table
split_non_repeating_forms(table, universal_fields, fields)
```

---

strsplitx	<i>Extended string splitting</i>
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---

**Description**

Can be used as a substitute of the base function. Main claim to fame is easing the split around the defined delimiter, see example.

**Usage**

```
strsplitx(x, split, type = "classic", perl = FALSE, ...)
```

**Arguments**

x	data
split	delimiter
type	Split type. Can be c("classic", "before", "after", "around")
perl	perl param from strsplit()
...	additional parameters are passed to base strsplit handling splits

**Value**

list

**Examples**

```
test <- c("12 months follow-up", "3 steps", "mRS 6 weeks", "Counting to 231 now")
strsplitx(test, "[0-9]", type="around")
```

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