

# Package ‘jwutil’

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**Title** Tools for Data Manipulation and Testing

**Version** 1.2.3

**Description** This is a set of simple utilities for various data manipulation and testing tasks. The goal is to use core R tools well, without bringing in many dependencies. Main areas of interest are semi-automated data frame manipulation, such as converting factors in multiple binary indicator columns. There are testing functions which provide 'testthat' expectations to permute arguments to function calls. There are functions and data to test extreme numbers, dates, and bad input of various kinds which should allow testing failure and corner cases, which can be used for fuzzing your functions. The test suite has many examples of usage.

**License** GPL-3

**URL** <https://github.com/jackwasey/jwutil>

**BugReports** <https://github.com/jackwasey/jwutil/issues>

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jwutil-package

*Tools for testing and data manipulation not found elsewhere***Description**

This is a set of simple utilities for various data manipulation and testing tasks. The goal is to use core R tools well, without bringing in many dependencies. Main areas of interest are semi-automated data frame manipulation, such as converting factors in multiple binary indicator columns. There are testing functions which provide 'testthat' expectations to permute arguments to function calls. There are functions and data to test extreme numbers, dates, and bad input of various kinds which

should allow testing failure and corner cases, which can be used for fuzzing your functions. The test suite has many examples of usage.

### Author(s)

**Maintainer:** Jack O. Wasey <jack@jackwasey.com> (0000-0003-3738-4637) [copyright holder]

### See Also

Useful links:

- <https://github.com/jackwasey/jwutil>
- Report bugs at <https://github.com/jackwasey/jwutil/issues>

---

add_time_to_date	<i>convert separate lists of dates and times to POSIXlt objects</i>
------------------	---

---

### Description

Some datetime data is presented as a separate dates and times. This function restores the full date-time.

### Usage

```
add_time_to_date(tms, dts, verbose = FALSE)
```

### Arguments

tms	vector of times, i.e. number in range 0 to 2400, as string or integer, with or without trailing zeros
dts	vector of dates, in string format %Y-%m-%d or simple R Date objects
verbose	single logical value, if TRUE then produce verbose messages

### Value

vector of POSIXlt date-times

---

affixFields	<i>update a set of data frame field names</i>
-------------	---

---

**Description**

prefix or suffix

**Usage**

```
affixFields(fields, affix, skip = NULL, renameHow = c("suffix",  
"prefix"), sep = ".")
```

**Arguments**

fields	char vector
affix	character
skip	char vector, defaults to include all fields
renameHow	should be "suffix" or "prefix", default is suffix
sep	default "."

**Value**

character vector, same length as fields

---

as_char_no_warn	<i>convert to character vector without warning</i>
-----------------	--

---

**Description**

convert to character vector without warning

**Usage**

```
as_char_no_warn(x)
```

**Arguments**

x	vector, typically numeric or a factor
---	---------------------------------------

**Value**

character vector

---

as\_numeric\_nowarn      *convert factor or vector to numeric without warnings*

---

### Description

correctly converts factors to vectors, and then converts to numeric or integer, which may silently introduce NAs. Invisible rounding errors can be a problem going from numeric to integer, so consider adding tolerance to this conversion. `asIntegerNoWarn` silently `floors`.

### Usage

```
as_numeric_nowarn(x)
```

```
as_integer_nowarn(x)
```

```
is_integerish(x, tol = 1e-09, na.ignore = FALSE)
```

```
areIntegers(x, tol = 1e-09, na.ignore = FALSE)
```

### Arguments

<code>x</code>	is a vector, probably of numbers or characters
<code>tol</code>	tolerance when considering if two numbers are integers, default 1e-9
<code>na.ignore</code>	logical, if TRUE will pass through NA values, otherwise, they are marked FALSE.

### Details

"are" functions return a value for each input, where "allIs" functions return a single logical.

### Value

numeric vector, may have NA values

logical vector

### Functions

- `areIntegers`: Deprecated

### Examples

```
stopifnot(is_integerish("1"))
```

---

bad_input	<i>bad input data for tests</i>
-----------	---------------------------------

---

**Description**

a variety of horrible data

**Usage**

```
bad_input
```

**Format**

An object of class list of length 43.

---

binary_col_names	<i>names of fields which are numeric, binary or combinations thereof</i>
------------------	--

---

**Description**

Doesn't make any allowance for factors.

**Usage**

```
binary_col_names(x, invert = FALSE)
```

```
two_cat_col_names(x, invert = FALSE, ignore_na = FALSE, trim = TRUE)
```

```
binary_cols(x, invert = FALSE)
```

```
two_cat_cols(x, invert = FALSE)
```

**Arguments**

x	data frame
invert	single logical, if true, will return non-binary columns
ignore_na	If TRUE, then return columns with two distinct values in addition to NA. Default is FALSE, i.e. NA is counted as a distinct item.
trim	If character column found, then trim white space before assessing

**Value**

vector of column names

**Functions**

- `two_cat_col_names`: Get the columns which have exactly two categories therein, not including NA values. This would catch 0,1 "Yes", "No", etc.
- `binary_cols`: Get the data frame containing just the binary columns.
- `two_cat_cols`: Get the data frame containing only columns of input which have two categories

**Examples**

```

dat <- data.frame(
  c("a", "b"), c(TRUE, FALSE), c(1, 0), c(1L, 0L),
  c(1L, 2L), c(0.1, 0.2), c("9", "8")
)
names(dat) <- c(
  "char", "bin", "binfloat", "binint",
  "int", "float", "charint"
)
binary_cols(dat)
binary_col_names(dat)
binary_col_names(dat, invert = TRUE)
df <- data.frame(
  x = c("A", "B", "A", "B"),
  y = letters[1:4],
  z = c("y", NA, "y", NA),
  stringsAsFactors = FALSE
)
two_cat_col_names(df)
df[1, 1] <- NA
df[2, 2] <- NA
df
stopifnot(two_cat_col_names(df) == "z")
stopifnot(two_cat_col_names(df, ignore_na = TRUE) == "x")

```

---

 build\_formula

*build simple linear formula from variable names*


---

**Description**

build simple linear formula from variable names given by two character vectors. TODO: allow unquoted names.

**Usage**

```
build_formula(left, right)
```

```
buildLinearFormula(left, right)
```



**Arguments**

left            character vector  
right           character vector

**Value**

formula

**Examples**

```
print(f <- build_formula(left = "A", right = c("B", "C")))
class(f)
build_formula(left = "Species", right = names(iris)[1:4])
```

---

combn\_subset            *all unique combinations of a vector and all its non-zero subsets*

---

**Description**

all unique combinations of a vector and all its non-zero subsets

**Usage**

```
combn_subset(x)
```

**Arguments**

x                    vector to be subsetted and combined

**Value**

list of vectors with all combinations of x and its subsets

**Examples**

```
combn_subset(c("a", "b"))
combn_subset(c(10, 20, 30))
combn_subset(NULL)
```

---

countIsNa	<i>count NA in vector</i>
-----------	---------------------------

---

**Description**

count the number of NAs in a vector. also consider 'base::anyNA'

**Usage**

```
countIsNa(x)
```

**Arguments**

x                    vector

**Value**

integer

---

countNonNaCumulative	<i>running totals of number of non-NA values in consecutive fields</i>
----------------------	--

---

**Description**

counts non-NA fields in first field, then progreses through fields, OR new field and saves running total for each field TODO: tests

**Usage**

```
countNonNaCumulative(d)
```

**Arguments**

d                    data.frame

**Value**

vector of cumulative non-NA counts with names corresponding to the given data frame

---

countNotNumeric	<i>count non-numeric elements</i>
-----------------	-----------------------------------

---

**Description**

counts the number of non-numeric elements in a vector, without throwing warnings

**Usage**

```
countNotNumeric(x)
```

**Arguments**

x is usually a character vector

**Details**

did have extras = c(".", "NA")

**Value**

integer

---

countNumeric	<i>count numeric elements</i>
--------------	-------------------------------

---

**Description**

counts the number of numeric elements in a vector, without throwing warnings

**Usage**

```
countNumeric(x)
```

**Arguments**

x is usually a character vector

**Value**

integer

---

dput\_expect\_equal      *dput a testthat test*

---

### Description

Generate an R expression containing a test that expectation for the given expression and its result. This is useful when you know that a certain output is correct, and wish to generate a test case to reflect this.

### Usage

```
dput_expect_equal(...)
```

### Arguments

...                    expressions

### Value

character vector with each element containing an R expression with expect\_equal test case corresponding to the evaluated input expressions.

### Examples

```
dput_expect_equal("a" %nin% c("b", "c", "d"))
```

---

drop\_duplicate\_fields      *Drop fields with duplicate data*

---

### Description

Compares all data in each field to every other field, and drops the latter match. Will find multiple matches. Doesn't do any type conversions yet. This is purely by content, not by field name.

### Usage

```
drop_duplicate_fields(df, verbose = FALSE)
```

```
dropDuplicateFields(df, verbose = FALSE)
```

### Arguments

df                    data.frame

verbose              single logical value, if TRUE then produce verbose messages

**Value**

data frame without duplicate fields

**Functions**

- dropDuplicateFields: Deprecated

**Examples**

```
d <- data.frame(LETTERS, letters, letters)[1:10, ]
drop_duplicate_fields(d)
```

---

drop_rows_with_na	<i>drops rows with NA values in specified fields</i>
-------------------	--

---

**Description**

employs [complete.cases](#) which is fast internal C code. Returns a data frame with unused factor levels dropped (these may have been introduced by dropping rows with some NA values)

**Usage**

```
drop_rows_with_na(x, fld = names(x), verbose = FALSE)
```

```
dropRowsWithNAField(x, fld = names(x), verbose = FALSE)
```

**Arguments**

x	data frame
fld	vector with names of fields which must have no NA values
verbose	single logical value, if TRUE then produce verbose messages

**Value**

data frame without rows containing NA in the specified data fields. There may be NA values in the resulting data frame in fields which are not listed in fld.

**Functions**

- dropRowsWithNAField: Deprecated, use drop\_rows\_with\_na

---

expect\_that\_combine\_all\_args

*alternative expect\_that from testthat which permutes all the inputs to a function which should give the same result where  $n \text{ args} \geq 2$  and the function is commutative.*

---

### Description

This makes a lot of assumptions, needs more testing. It can't handle mixed error/no error outcomes after permutation, which is an important feature to consider. The command following this function attaches this function to the testthat namespace. This means that it can call internal testthat functions, but does not mean it appears as testthat::expect\_that\_combine

### Usage

```
expect_that_combine_all_args(object, condition, info = NULL,
  label = NULL)
```

```
expect_that_combine_first_arg(object, condition, info = NULL,
  label = NULL)
```

### Arguments

object	See <a href="#">expect_that</a> .
condition	See <a href="#">expect_that</a> .
info	See <a href="#">expect_that</a> .
label	See <a href="#">expect_that</a> .

### Value

testthat result

### Examples

```
expect_that_combine_all_args(
  sum(1, 2, 3),
  testthat::equals(6)
)
## Not run:
expect_that_combine_all_args(stop("a", "b"), testthat::throws_error())
expect_that_combine_all_args(sum(1, 2), testthat::equals(3))
expect_that_combine_first_arg(sum(c(1, 2)), testthat::equals(3))

## End(Not run)
```

---

extreme_numbers	<i>extreme numbers</i>
-----------------	------------------------

---

**Description**

very biggest and smallest non-zero numbers the current machine can handle, positive and negative.

**Usage**

```
extreme_numbers
```

**Format**

An object of class numeric of length 8.

---

factor_nosort	<i>Fast Factor Generation</i>
---------------	-------------------------------

---

**Description**

This function generates factors more quickly, without leveraging fastmatch. The speed increase with fastmatch for ICD-9 codes was about 33 using Rcpp, and a hashed matching algorithm.

**Usage**

```
factor_nosort(x, levels = NULL, labels = levels)
```

**Arguments**

x	An object of atomic type integer, numeric, character or logical.
levels	An optional character vector of levels. Is coerced to the same type as x. By default, we compute the levels as <code>sort(unique.default(x))</code> .
labels	A set of labels used to rename the levels, if desired.

**Details**

NaNs are converted to NA when used on numeric values. Extracted from <https://github.com/kevinushey/Kmisc.git>

These feature from base R are missing: `exclude = NA`, `ordered = is.ordered(x)`, `nmax = NA`

I don't think there is any requirement for factor levels to be sorted in advance, especially not for ICD-9 codes where a simple alphanumeric sorting will likely be completely wrong.

**Author(s)**

Kevin Ushey, adapted by Jack Wasey

**Examples**

```
## Not run:
pts <- icd::random_unordered_patients(1e7)
u <- unique.default(pts$code)
# this shows that stringr (which uses stringi) sort takes 50% longer than
# built-in R sort.
microbenchmark::microbenchmark(sort(u), str_sort(u))

# this shows that \code{factor_} is about 50% faster than \code{factor} for
# big vectors of strings

# without sorting is much faster:
microbenchmark::microbenchmark(factor(pts$code),
  # factor_(pts$code),
  factor_nosort(pts$code),
  times = 25
)

## End(Not run)
```

---

factor\_to\_df

---

*Convert factor into a data.frame of logicals*


---

**Description**

Convert a single factor into a data.frame with multiple true or false fields, one for each factor. The ‘vtreat’ package may be a better choice for more comprehensive data preparation.

**Usage**

```
factor_to_df(fctr, prefix = deparse(substitute(fctr)), sep = "",
  drop_empty = TRUE, na_as_col = TRUE, verbose = FALSE)

factorToDataframeLogical(fctr, prefix = deparse(substitute(fctr)),
  sep = "", drop_empty = TRUE, na_as_col = TRUE, verbose = FALSE)
```

**Arguments**

fctr	factor
prefix	defaults to "f" to pre-pend the factor level when constructing the data frame columns names
sep	scalar character, introduced between factor names and levels when forming new data frame column names
drop_empty	logical, if ‘TRUE’ (the default) factor levels with no associated values are dropped.
na_as_col	logical scalar: if NA data and/or NA levels, then covert to NA strings and expand these as for any other factor
verbose	single logical value, if TRUE then produce verbose messages



**Value**

data.frame with columns of logicals

**Examples**

```
n <- 10
m <- 20
l <- LETTERS[seq_len(n)]
set.seed(1441)
f <- factor(sample(l, m, replace = TRUE), levels = l)
g <- factor_to_df(f, drop_empty = FALSE)
print(g)
stopifnot(nrow(g) == m, ncol(g) == n)
factor_to_df(
  shuffle(factor(shuffle(LETTERS[1:10]))),
  prefix = ""
)
factor_to_df(factor(c(NA, 1, 2, 3)))
factor_to_df(factor(c(NA, 1, 2, 3)), na_as_col = FALSE)
```

---

fillMissingCombs      *fill out missing combinations of factors with NA*

---

**Description**

fill out missing combinations of factors with NA

**Usage**

```
fillMissingCombs(df)
```

**Arguments**

df                    data frame

**Details**

Adapted from [http://www.cookbook-r.com/Manipulating\\_data/Summarizing\\_data/#using-aggregate](http://www.cookbook-r.com/Manipulating_data/Summarizing_data/#using-aggregate)

---

filter_better	<i>filter data with diagnostics</i>
---------------	-------------------------------------

---

**Description**

applies an expression to a data frame, and gives information about the numbers of dropped rows.

**Usage**

```
filter_better(x, expr, verbose = TRUE)
```

```
filterBetter(x, expr, verbose = TRUE)
```

**Arguments**

x	data frame
expr	expression in the context of the data frame, i.e. the terms should be column names.
verbose	logical default is TRUE

**Value**

filtered data frame

**Functions**

- filterBetter: Deprecated

---

fix_na_ish	<i>Fix NA-like strings to be NA (or other value of choice)</i>
------------	--

---

**Description**

Fix NA-like strings to be NA (or other value of choice)

**Usage**

```
fix_na_ish(x, extra_na = NULL, new_val = NA)
```

**Arguments**

x	data frame
extra_na	Additional values to consider equivalent to NA
new_val	New value to be used instead of NA-ish values, default is NA

**Examples**

```
df <- data.frame(
  a = c("NA", "n/a", 1, NA),
  b = c("three", "na", NaN, " N/A "),
  stringsAsFactors = FALSE
)
df
fix_na-ish(df)
fix_na-ish(df, extra_na = "three", new_val = "0")
```

---

flattenList	<i>flatten a list</i>
-------------	-----------------------

---

**Description**

unlike unlist, this function returns a list of objects of different data types, but removes any depth

**Usage**

```
flattenList(..., na_rm = FALSE)
```

**Arguments**

...	list or any set of objects which will be made into a list, may include lists and nested lists
na_rm	will drop NA values if TRUE

**Value**

list without nested lists, objects with preserved data types

**Source**

<https://stackoverflow.com/questions/8139677/how-to-flatten-a-list-to-a-list-without-coercion>

---

getDropped	<i>get items or numerics that would be dropped in a merge</i>
------------	---

---

**Description**

converts both vectors to numeric. This simulates merging when one key is character (but contains integer numbers), and another key is stored as integer.

**Usage**

```
getDropped(x, y)
```

**Arguments**

x                    vector or factor  
y                    vector or factor

**Value**

list of two vectors

---

get\_factor\_fields      *get names of the factor fields in a data frame*

---

**Description**

Get the names of those fields in a data frame which are factors.

**Usage**

```
get_factor_fields(x, consider = names(x))  
get_non_factor_fields(x, consider = names(x))  
getFactorNames(x, consider = names(x))  
getNonFactorNames(x, consider = names(x))
```

**Arguments**

x                    data frame  
consider            character vector of field names of the data frame to test, default is to use all of them.

**Value**

vector

**Functions**

- get\_non\_factor\_fields: Get the fields which are not factors, instead.
- getFactorNames: Deprecated
- getNonFactorNames: Deprecated

---

get_na_fields	<i>get NA field names from data frame</i>
---------------	---

---

**Description**

Get the names of any columns in a data frame which have NA values.

**Usage**

```
get_na_fields(x, na_ish = FALSE, extra_na = NULL)
```

```
getNAFields(x, na_ish = FALSE, extra_na = NULL)
```

```
get_non_na_fields(x)
```

```
getNonNAFields(x)
```

**Arguments**

x	data.frame
na_ish	Logical, if 'TRUE', also consider NA-like strings, using 'is_na_ish'
extra_na	passed on to 'is_na_ish'

**Value**

vector of names of fields which contain any NA values, length zero if no matches

**Functions**

- getNAFields: Deprecated
- getNonNAFields: Deprecated

---

get_numeric_char_field_names	<i>Find columns which are numeric</i>
------------------------------	---------------------------------------

---

**Description**

Get field names, or the data itself, of fields in a data frame which are numeric, or numeric-like characters.

**Usage**

```

get_numeric_char_field_names(x, invert = FALSE, attrition = 0.05)

get_numeric_field_names(x, invert = FALSE)

get_numeric_fields(x, invert = FALSE)

```

**Arguments**

x	Data frame
invert	Logical, if FALSE – the default – the numeric fields are returned, otherwise, non-numeric fields are returned.
attrition	If less than this proportion of rows become NA on conversion to numeric, then accept this is a numeric column after all.

---

 invwhich

*inverse which*


---

**Description**

for a given vector of ordinals which would reference items in a vector, list etc, invwhich returns a logical vector with TRUE for the cited positions. If length is not provided, the maximum index is used.

**Usage**

```
invwhich(which, len = max(which))
```

**Arguments**

which	integer vector of indices, as would be produced by which
len	integer scalar: length of return vector, defaults to max(which)

**Value**

logical vector of length length

---

is.Date	<i>is the object a Date</i>
---------	-----------------------------

---

**Description**

copied from lubridate

**Usage**

```
is.Date(x)
```

**Arguments**

x	object to test
---	----------------

**Value**

logical

---

isFlat	<i>determine whether a list is nested</i>
--------	---

---

**Description**

Returns TRUE if the given list is not nested.

**Usage**

```
isFlat(x)
```

**Arguments**

x	list
---	------

**Value**

single logical

---

isRowSorted	<i>is every row sorted?</i>
-------------	-----------------------------

---

**Description**

Quickly run through rows of a matrix looking for any non-ascending rows in C++

**Usage**

```
isRowSorted(x)
```

**Arguments**

x                    matrix, each row containing ordered or disordered numerics

---

isValidTime	<i>check if a time is valid in 24h clock</i>
-------------	--

---

**Description**

allow leading and trailing space, optional colon in middle, 2400 is not allowed. TODO: can lubri-date do this better?

**Usage**

```
isValidTime(tms, na.rm = FALSE)
```

**Arguments**

tms                    is a vector of characters which may represent times  
na.rm                   logical if true, will ignore NA values, otherwise these will test as invalid.

**Value**

logical vector, with NA out if NA given



---

is_na_ish	<i>Determine whether a value is, or should be, 'NA'</i>
-----------	---

---

**Description**

Determine whether a value is, or should be, 'NA'

**Usage**

```
is_na_ish(x, extra_na = NULL)
```

**Arguments**

x	vector to test
extra_na	Additional values to consider equivalent to NA

**Examples**

```
is_na_ish(c(NA, "1"))  
is_na_ish(c("NA", "N/A", "NaN"))  
is_na_ish(c(NA))  
is_na_ish(c(NA))
```

---

is_numeric_str	<i>Which elements of a character vector are numeric</i>
----------------	---

---

**Description**

Takes a character vector and returns a logical vector of the same length, indicating which values are numeric. NA is considered non-numeric. NA is never returned from this function.

**Usage**

```
is_numeric_str(x, extras = c(".", "NA", NA))  
  
areNumeric(x, extras = c(".", "NA", NA))
```

**Arguments**

x	character vector
extras	character vector containing acceptable alternatives to numeric values which will result in returning TRUE for that element. Default is c(".", "NA", NA).

**Value**

logical vector of same length as input

**Functions**

- areNumeric: Deprecated

**Examples**

```
areNumeric(c("1", "2", "3"))
areNumeric(c("1L", "2.2"))
areNumeric(c("NA", NA, ".", "", "-1.9"))
```

---

jw_df_basics	<i>minimal basic pre-processing metrics</i>
--------------	---

---

**Description**

minimal basic pre-processing metrics

**Usage**

```
jw_df_basics(x, df_list)
```

**Arguments**

x	data.frame input
df_list	list of data frames

---

jw_scan_build	<i>Build with current Makevars, but with clang scan-build static analysis</i>
---------------	---

---

**Description**

C and C++ compilers are replaced by ‘scan-build clang’, and restored afterwards. Other flags and anything else in ‘~/R/Makevars’ is left alone.

**Usage**

```
jw_scan_build(path = ".", clang = "clang-8",
  scan_build = "scan-build")
```

**Arguments**

path	Path to package root, default is ".".
clang	Path or name of clang compiler executable. Currently ‘clang-8’ which is what MacOS homebrew currently (early 2019) provides.
scan_build	Path or name of scan-build executable. Current ‘scan-build’, also from MacOS Homebrew. On linux, this has the LLVM version suffix, e.g., ‘scan-build-8’.

---

listTrim	<i>trim null or empty values from a list</i>
----------	--

---

**Description**

delele null/empty entries in a list. Recursively looks through list if nested.

**Usage**

```
listTrim(x)
```

**Arguments**

x	list
---	------

**Value**

trimmed list

---

listTrimFlat	<i>trim null or empty values from a list</i>
--------------	--

---

**Description**

Trim NULL or empty values from a flat list.

**Usage**

```
listTrimFlat(x)
```

**Arguments**

x	list
---	------

**Value**

trimmed list

---

list_named	<i>Make a list using input argument names as names</i>
------------	--

---

**Description**

Make a list using input argument names as names

**Usage**

```
list_named(...)
```

**Arguments**

... arguments whose names become list item names, and whose values become the values in the list

**Examples**

```
a <- c(1, 2)
b <- c("c", "d")
stopifnot(
  identical(
    list_named(a, b),
    list(a = a, b = b)
  )
)
```

---

logical_to_binary	<i>Convert logical columns of data frame to 0s and 1s</i>
-------------------	---

---

**Description**

Encode TRUE as 1, and FALSE as 0 (integers)

**Usage**

```
logical_to_binary(x)
```

```
logicalToBinary(x)
```

**Arguments**

x data frame which may contain logical fields

**Value**

data frame without logical fields

**Examples**

```
d <- data.frame(
  a = c(TRUE, FALSE, TRUE),
  b = c(FALSE, TRUE, FALSE),
  c = c(-1, 0, 1),
  d = c("not", "logical", "values")
)
logical_to_binary(d)
```

ls.objects

*Summarize objects***Description**

Get type, size (bytes) and dimensions of objects

**Usage**

```
ls.objects(env = parent.frame(), pattern, order.by, decreasing = FALSE,
  head = FALSE, n = 5)
```

**Arguments**

env	Environment to search, default is the parent frame
pattern	regex pattern to match objects of interest
order.by	which column to order by
decreasing	default is TRUE
head	default is FALSE but if true, just show top n
n	number to show if limiting to head

lsf

*list all functions in a package***Description**

List functions in a package

**Usage**

```
lsf(pkg)
```

**Arguments**

pkg	character string containing package name
-----	--

**Value**

character vector of functions in given package

---

lsos	<i>show largest objects</i>
------	-----------------------------

---

**Description**

<https://gist.github.com/1187166.git> Taken from <http://stackoverflow.com/questions/1358003/tricks-to-manage-the-available-memory-in-an-r-session>

**Usage**

```
lsos(..., n = 10)
```

**Arguments**

...	arguments passed on to <code>.ls.objects</code>
n	scalar integer, number of objects to show

---

lsp	<i>List all items in a package</i>
-----	------------------------------------

---

**Description**

By default includes names beginning with '.'

**Usage**

```
lsp(package, all.names = TRUE, pattern)
```

**Arguments**

package	character scalar: name of the package
all.names	= TRUE, set to FALSE to ignore items beginning with a period
pattern	= optional pattern to match

**Value**

character vector of package contents

**Examples**

```
lsp("jwutil")
tail(lsp("base"), 30L)
```

---

match_multi	<i>Match across columns for multiple lookup values</i>
-------------	--

---

**Description**

This provides a succinct way to query a data frame for conditions, which is otherwise very verbose in base R or dplyr

**Usage**

```
match_multi(x, cols, table, incomparables = NULL)
```

**Arguments**

x	data.frame
cols	character vector of column names to be found in x
table	vector of items to find
incomparables	passed on to the base function match

**Value**

matrix with same number of rows as x, and a column for each of cols

**Examples**

```
j <- cars[1:10, ]
match_multi(j, "speed", 7)
match_multi(j, "dist", 22)
match_multi(j, c("speed", "dist"), 10)
match_multi(j, c("speed", "dist"), c(7, 17))
```

---

mergeLists	<i>merge lists by names</i>
------------	-----------------------------

---

**Description**

merge lists by vector combining all the vector elements of the list items with the matching names. Unnamed vectors in the list will be dropped silently.

**Usage**

```
mergeLists(x, y)
```

**Arguments**

x	unnested list with named elements, each of which is a vector
y	unnested list with named elements, each of which is a vector

**Value**

list of vectors

---

merge_better	<i>Merge better</i>
--------------	---------------------

---

**Description**

Apply built-in R [merge](#) but with additional features for safety and information.

**Usage**

```
merge_better(x, y, by.x, by.y, all.x = FALSE, all.y = FALSE,
  affix = NULL, renameConflict = c("suffix", "prefix"),
  renameAll = c("no", "suffix", "prefix"), convert_factors = TRUE,
  verbose = FALSE)
```

```
mergeBetter(x, y, by.x, by.y, all.x = FALSE, all.y = FALSE,
  affix = NULL, renameConflict = c("suffix", "prefix"),
  renameAll = c("no", "suffix", "prefix"), convert_factors = TRUE,
  verbose = FALSE)
```

**Arguments**

x	data frame
y	data frame
by.x	field in x to merge on. Unlike merge, this is compulsory.
by.y	field in y to merge on. Unlike merge, this is compulsory.
all.x	outer join to keep all x values
all.y	outer join to keep all y values
affix	either prefix or suffix to disambiguate files. By default, this is the name of the table specified in y. In all other respects in this function, x and y are symmetric.
renameConflict	- determines whether prefix or suffix is added to disambiguate conflicting column names. Value can be "suffix", "prefix". Suffix is the default.
renameAll	- regardless of column name clashes, "prefix" or "suffix" with every field with original table name, or "no" for neither
convert_factors	Default is TRUE which causes factors to be converted to character before merge. This is almost certainly safer.
verbose	logical or numbers 0, 1 or 2. 1 or TRUE will give moderate verbosity, 2 will give full verbosity. 0 or FALSE turns off all messages.



**Value**

merged data frame

**Examples**

```
df <- data.frame(a = c("1", "2"), b = 1:2, stringsAsFactors = FALSE)
eg <- data.frame(a = c("1", "3"), b = 3:4, stringsAsFactors = FALSE)
mergeBetter(x = df, y = eg, by.x = "a", by.y = "a", verbose = TRUE)
```

---

min\_r\_version

*Find minimum R version required for package*

---

**Description**

Recursively search dependencies for R version, and find the highest stated R version requirement.

**Usage**

```
min_r_version(pkg)
```

**Arguments**

pkg                      string with name of package to check

**Source**

Based on ideas from <http://stackoverflow.com/questions/38686427/determine-minimum-r-version-for-all-package-dependencies>

**Examples**

```
base <- c(
  "base", "compiler", "datasets", "grDevices", "graphics",
  "grid", "methods", "parallel", "profile", "splines", "stats",
  "stats4", "tcltk", "tools", "translations"
)
## Not run:
base_reqs <- lapply(base, min_r_version)
contrib <- c(
  "KernSmooth", "MASS", "Matrix", "boot",
  "class", "cluster", "codetools", "foreign", "lattice",
  "mgcv", "nlme", "nnet", "rpart", "spatial", "survival"
)
contrib_reqs <- lapply(contrib, min_r_version)
min_r_version("icd")

## End(Not run)
```

---

npc *Print integers with percentage of total rounded to integer*

---

**Description**

Intended for succinctly printing summary data in a scientific publication.

**Usage**

```
npc(x, n, fmt = "%d (%s)")
```

**Arguments**

x	numeric number
n	numeric total
fmt	sprintf format, default being %d (%s)

**Examples**

```
npc(1, 100)
npc(1, 1)
npc(2, 1)
npc(1.321, 7.7432)
npc(7239, 234897)
npc(-10, 1000)
```

---

numbers\_to\_long\_and\_float  
*convert numbers to long and float types*

---

**Description**

intended for generating values for stress testing functions

**Usage**

```
numbers_to_long_and_float(..., na.rm = TRUE)
```

**Arguments**

...	list of values to convert to long and double
na.rm	logical, defaults to TRUE, so output contains only long and float values.

**Value**

list of long and double versions of convertible values from the input

---

opt\_binary\_brute      *selects columns from a data frame using an optimization function*

---

### Description

The optimization function is called with the data frame `x` and the names of each combination of the names of `x`'s columns. An example of real-world usage is to automate selection of columns according to the optimization function.

### Usage

```
opt_binary_brute(x, fun = opt_binary_fun, verbose = FALSE)
```

### Arguments

<code>x</code>	data frame
<code>fun</code>	function which takes parameters <code>x = data.frame</code> , <code>n = columns</code>
<code>verbose</code>	single logical value, if TRUE then produce verbose messages

### Examples

```
j <- data.frame(a = 1:5, b = 6:2, c = c(0, 2, 4, 6, 8))
opt_binary_brute(j)
j[1, 1] <- NA
j[1:4, 2] <- NA
my_opt_fun <- function(x, n) sum(!unlist(lapply(x, is.na)))
opt_binary_brute(j, fun = my_opt_fun)
```

---

percentize      *Convert a number into rounded integer percentage string*

---

### Description

The number is converted into a percentage, then rounded.

### Usage

```
percentize(x)
```

### Arguments

<code>x</code>	numeric
----------------	---------

**Examples**

```
percentize(-1)
percentize(1L)
percentize(7.7)
percentize(0.01)
percentize(0.001)
```

---

percent_signif	<i>Return percentage string to given significant figures</i>
----------------	--

---

**Description**

From jwutil development version

**Usage**

```
percent_signif(x, figures = 3, sep = "")
```

**Arguments**

x	numeric or integer values
figures	integer number of significant figures to format
sep	character used to separate number from percent symbol, default is empty string

---

permute	<i>Generate all permutations of input</i>
---------	---

---

**Description**

Systematically permute the input vector or list, which is very slow for long x. Am amazed something this simple isn't either in base R, or in a straightforward form in a package.

**Usage**

```
permute(x)
```

**Arguments**

x	list or vector
---	----------------

**Details**

TODO: limit to a certain cut-off, after which we randomly sample

**Value**

data frame, each row being one permutation

**Examples**

```
ltr <- c("a", "b", "c", "d")
x <- permute(ltr)
print(x)
stopifnot(nrow(x) == factorial(length(ltr)))
ltr <- c("a", "b", "b")
x <- permute(ltr)
print(x)
stopifnot(nrow(x) == factorial(length(ltr)))
```

---

permuteWithRepeats      *Generate all permutations of input, reusing values in each result row*

---

**Description**

Expand the given vector into all possible values in each location, with or without duplicates.

**Usage**

```
permuteWithRepeats(x, unique = TRUE)
```

**Arguments**

x                      list or vector  
unique                  logical, if TRUE, the default, only unique results are returned

**Value**

data frame, each row being one permutation

**Examples**

```
ltr <- c("a", "b", "c")
x <- permuteWithRepeats(ltr, unique = FALSE)
print(x)
stopifnot(nrow(x) == length(ltr)^length(ltr))
# duplicate results are dropped
y <- permuteWithRepeats(c("X", "Y", "Y"))
print(y)
stopifnot(nrow(y) == 2^3)
z <- permuteWithRepeats(c("X", "Y", "Y", "Y"))
stopifnot(nrow(z) == 2^4)
a <- permuteWithRepeats(c(1, 2, 3, 1))
stopifnot(nrow(a) == 3^4)
```

---

platformIsLinux	<i>Are we running on Linux, Mac or Windows?</i>
-----------------	---

---

**Description**

Are we running on Linux, Mac or Windows?

**Usage**

```
platformIsLinux()
```

```
platformIsWindows()
```

```
platformIsMac()
```

**Value**

logical

---

propIsNa	<i>Proportion of NA values in a vector</i>
----------	--

---

**Description**

get fraction of NA in a vector

**Usage**

```
propIsNa(x)
```

**Arguments**

x is a vector which may have NA values

**Value**

numeric proportion of NAs in the supplied vector

---

`propNaPerField`      *return proportion of NA values per field*

---

**Description**

Return proportion of values which are NA in each field of the given data frame.

**Usage**

```
propNaPerField(x)
```

**Arguments**

`x`                    is a data frame

**Value**

numeric vector

---

`propRowSorted`      *proportion of non-descending rows in matrix*

---

**Description**

first performs `isRowSorted` to get a logical vector, then sums TRUE values and takes fraction of total

**Usage**

```
propRowSorted(x)
```

**Arguments**

`x`                    matrix, each row containing ordered or disordered numerics

**Value**

double, the proportion from 0 to 1

---

random\_test\_dates      *generate random Dates or POSIXlt test datetimes*

---

**Description**

generate random Dates and POSIXlt test datetimes

**Usage**

```
random_test_dates(n = n_rnd, origin = as.Date("2000-01-01"),
  dayspread = 365 * 150)
```

```
random_test_posixlt_datetimes(n = n_rnd,
  origin = as.Date("2000-01-01"), dayspread = 365 * 150)
```

**Arguments**

n	integer number to generate
origin	Date defaults to Jan 1, 2000.
dayspread	integer number of days either side of origin to pick random dates from, defaults to 150 years.

**Value**

vector of POSIXlt datetimes or Dates

---

random\_test\_numbers      *create extreme random numbers*

---

**Description**

create random Dates, POSIX dates, letters and numbers. The numbers explore limits of R precision and floating point and integer ranges. Zero, negatives, positives.

**Usage**

```
random_test_numbers(n = n_rnd, min = NULL, max = NULL, hole = NULL)
```

```
random_test_integers(n = n_rnd, min = -.Machine$integer.max,
  max = .Machine$integer.max, hole = NULL)
```

```
random_test_letters(n = n_rnd, max_str_len = 257)
```



**Arguments**

n	integer number of each group to generate
min	optional minimum number
max	optional maximum number
hole	is a closed range of numbers not to include, e.g. c(1,2) would discard 1, 1.1 pi/2 and 2
max_str_len	integer scalar, maximum length of possible strings created, as distinct from number of strings given by n

**Value**

vector length  $5n+1$  containing variety of difficult numbers for testing purposes

---

read\_xlsx\_linux      *read .xlsx file, interpret as CSV, and return a data frame*

---

**Description**

currently relies on Linux xlsx2csv command, but could potentially be done with VB script in Windows. This offers a different backend to other Excel parsing functions in R,

**Usage**

```
read_xlsx_linux(file)
```

**Arguments**

file            is the path to the .xlsx file

**Value**

data frame

**See Also**

readxl package by Hadley Wickham

---

read_zip_url	<i>read file from zip at URL</i>
--------------	----------------------------------

---

### Description

downloads zip file, and opens named file filename, or the single file in zip if filename is not specified. FUN is a function, with additional arguments to FUN given by .... @details TODO: update from icd package

### Usage

```
read_zip_url(url, filename = NULL, FUN = readLines, ...)
```

```
read.zip.url(url, filename = NULL, FUN = readLines, ...)
```

### Arguments

url	character vector of length one containing URL of zip file.
filename	character vector of length one containing name of file to extract from zip. If not specified, and the zip contains a single file, then this single file will be used.
FUN	function used to process the file in the zip, defaults to readLines. The first argument to FUN will be the path of the extracted filename
...	further arguments to FUN

### Functions

- read.zip.url: Deprecated

---

reqinst	<i>Load packages with library, installing any which are missing</i>
---------	---

---

### Description

Load packages with library, installing any which are missing

### Usage

```
reqinst(pkgs)
```

### Arguments

pkgs	character vector of packages to load and attach, with installation if necessary
------	---

---

rm_r	<i>recursive remove</i>
------	-------------------------

---

**Description**

search through environments until the variables in the list `x` are all gone. This doesn't delete functions.

**Usage**

```
rm_r(x, envir = parent.frame())
```

**Arguments**

<code>x</code>	variables to annihilate
<code>envir</code>	environment to start at, defaults to calling frame.

---

save_in_data_dir	<i>Save given variable in package data directory</i>
------------------	--

---

**Description**

File is named `varname.RData` with an optional suffix before `.RData`

**Usage**

```
save_in_data_dir(var_name, suffix = "", data_path = "data",
  package_dir = getwd(), envir = parent.frame())
```

**Arguments**

<code>var_name</code>	character or symbol, e.g. "myvar" or myvar, either of which would find myvar in the parent environment, and save it as myvar.RData in package_root/data.
<code>suffix</code>	character scalar
<code>data_path</code>	path to data directory, default is data in current directory.
<code>package_dir</code>	character containing the directory root of the package tree in which to save the data. Default is the current working directory.
<code>envir</code>	environment in which to look for the variable to save

**Value**

invisibly returns the data

---

shuffle	<i>Shuffle a vector</i>
---------	-------------------------

---

**Description**

Randomly shuffle the order of a vector or list. This is to improve quality of bad data to throw at functions when testing.

**Usage**

```
shuffle(x)
```

**Arguments**

x	list or vector
---	----------------

**Value**

list or vector of same length as input, (probably) in a different order

**Examples**

```
set.seed(1441)
shuffle(LETTERS)
```

---

sort_clip_char	<i>Take clipboard contents, and write sorted character vector back</i>
----------------	--

---

**Description**

Take clipboard contents, and write sorted character vector back

**Usage**

```
sort_clip_char(c1 = NULL)
```

**Arguments**

c1	Name of class to give to data before sorting, default is NULL.
----	--

---

source_purl	<i>Extract code from knitr vignette and source it</i>
-------------	---

---

**Description**

Extract code from knitr vignette and source it.

**Usage**

```
source_purl(input, documentation = 1L, ...)
```

**Arguments**

input	path to file as single character string
documentation	single integer value passed on to <code>knitr::purl</code> . An integer specifying the level of documentation to go the tangled script: 0 means pure code (discard all text chunks); 1 (default) means add the chunk headers to code; 2 means add all text chunks to code as roxygen comments
...	further parameters passed to source

---

strip	<i>strip all whitespace</i>
-------	-----------------------------

---

**Description**

could do this with regular expression, but slow, and this function is called frequently. My only use case works with removal of all space character whitespace, and I don't expect <TAB>. This uses non-unicode aware matching for speed. This can be changed by setting `useBytes` to `FALSE`.

**Usage**

```
strip(x, pattern = " ", useBytes = TRUE)
```

**Arguments**

x	is a character vector to strip
pattern	is the non-regex of the character to strip, default " "
useBytes	logical scalar. Unlike <code>gsub</code> , this will default to <code>TRUE</code> here, therefore breaking unicode.

**Details**

`gsub` is probably quicker than `stringr/stringi`. For comorbidity processing, this package prefers the faster [base](#) functions, whereas `stringr` is used for tasks which are not time critical, e.g. parsing source data to be included in the distributed `icd` package.

**Value**

character vector

**Examples**

```
## Not run:
requireNamespace("microbenchmark")
requireNamespace("stringr")
x <- random_string(25000)
microbenchmark::microbenchmark(
  gsub(x = x, pattern = "A", replacement = "", fixed = TRUE, useBytes = TRUE),
  gsub(x = x, pattern = "A", replacement = "", fixed = TRUE, useBytes = TRUE, perl = TRUE),
  gsub(x = x, pattern = "A", replacement = ""),
  stringr::str_replace_all(x, "A", "")
)

## End(Not run)
```

---

strip\_for\_formula      *strip a string so that it can be used as a variable name in a formula.*

---

**Description**

This excludes many symbols, so just strip all symbols leaving alphanumeric, and no whitespace.

**Usage**

```
strip_for_formula(x)
```

**Arguments**

x                      character vector of potential formula variables

**Value**

character vector of length x

---

str_multi_match	<i>return the actual matches from a bracketed regex</i>
-----------------	---

---

**Description**

Be careful: this may throw funny results for exotic regex, but so far, it seems okay. it also drops the first result which always seems to be a duplicate or whole-string match

**Usage**

```
str_multi_match(pattern, text, dropEmpty = FALSE, ...)
```

```
strMultiMatch(pattern, text, dropEmpty = FALSE, ...)
```

**Arguments**

pattern	regular expression: if it has bracketed sections, these submatches are returned
text	is the string to match against. This vector should be the same length as the pattern vector, or the pattern vector should be length one.
dropEmpty	logical whether to drop rows with no matches
...	are additional parameters passed to regexexec and regmatches. I haven't tried this: it may need two separate variables containing lists of params, since this will send everything to both functions.

**Value**

list of character vectors, list length being the length of the input text vector.

**Functions**

- strMultiMatch: Deprecated

---

trim	<i>strip whitespace from ends of each string in given character vector</i>
------	--

---

**Description**

slower than strip.

**Usage**

```
trim(x)
```

**Arguments**

x	is a character vector to trim
---	-------------------------------

**Value**

character vector

---

two_cat_to_logical	<i>Take dataframe, and convert any columns with just two categories into logical</i>
--------------------	--

---

**Description**

E.g. "Yes" would be converted to TRUE, "0" to FALSE, etc. If heuristics fail, then the function stops with an error message. NA values are counted, unless ignore\_na is TRUE. When they are considered, na\_val indicates whether they are attributed TRUE or FALSE.

**Usage**

```
two_cat_to_logical(x, ignore_na = FALSE, na_val = FALSE)
```

**Arguments**

x	input data frame
ignore_na	logical
na_val	Single value to use in place of NA``, default is FALSE`

**Value**

data frame with two categories columns replaced by logical columns

**Examples**

```
df <- data.frame(
  a = c("y", "n", "y", "y", "n"),
  b = c(FALSE, TRUE, FALSE, TRUE, TRUE),
  c = c(NA, NA, NA, NA, NA),
  d = c(NA, "yes", NA, NA, "yes"),
  e = c("y ", "n ", NA, "y ", "n "),
  f = c("YES ", "NO ", "NO ", " YES", " NO "),
  stringsAsFactors = FALSE
)
df
res <- two_cat_to_logical(df)
stopifnot(identical(res$a, c(TRUE, FALSE, TRUE, TRUE, FALSE)))
stopifnot(identical(res$b, c(FALSE, TRUE, FALSE, TRUE, TRUE)))
two_cat_to_logical(df, ignore_na = TRUE)
```



---

unzip_single	<i>unzip a single file from URL</i>
--------------	-------------------------------------

---

**Description**

take a single file from zip located at a given URL, unzip into temporary directory, and copy to the given save\_path

**Usage**

```
unzip_single(url, file_name, save_path)
```

**Arguments**

url	URL of a zip file
file_name	file name of the resource within the zip file
save_path	file path to save the first file from the zip

---

unzip_to_data_raw	<i>Unzip file to data-raw</i>
-------------------	-------------------------------

---

**Description**

Get a zip file from a URL, extract contents, and save file in data-raw. If the file already exists there, it is only retrieved if force is set to TRUE. If offline is FALSE, then NULL is returned if the file isn't already downloaded.

**Usage**

```
unzip_to_data_raw(url, file_name, force = FALSE, verbose = FALSE,
  offline = TRUE, data_raw_path = "data-raw")
```

```
download_to_data_raw(url, file_name = regmatches(url, regexpr("[^/]*$",
  url)), offline = TRUE, data_raw_path = "data-raw")
```

**Arguments**

url	URL of a zip file
file_name	file name of a single file in that zip
force	logical, if TRUE, then download even if already in data-raw
verbose	single logical value, if TRUE then produce verbose messages
offline	single logical, if TRUE then don't pull the file from internet, only return path and file name if the file already exists in data-raw. This is helpful for testing without using the internet.
data_raw_path	path where the data-raw directory is.

**Details**

The file name is changed to a conservative cross platform name using `make.names`

**Value**

path of unzipped file in `data-raw`

---

<code>update_github_pkgs</code>	<i>Update github_install packages</i>
---------------------------------	---------------------------------------

---

**Description**

Update `github_install` packages

**Usage**

```
update_github_pkgs()
```

**Value**

Returns invisibly the names of packages which need updating. The function outputs the commands to run to actually update them (by reinstalling from github). Doesn't do this automatically because it would mean bringing in a lot of dependencies.

---

<code>zeroes</code>	<i>zeroes</i>
---------------------	---------------

---

**Description**

long, float and complex types

**Usage**

```
zeroes
```

**Format**

An object of class `list` of length 3.

---

zero_na	<i>Zero NA values in a data.frame</i>
---------	---------------------------------------

---

**Description**

Zero NA values in a data.frame, including cols and excluding ignore. Also does not replace Date or POSIXt fields.

**Usage**

```
zero_na(x, cols = names(x), ignore = character(), verbose = FALSE,
        na_ish = TRUE, new_val = 0)
```

**Arguments**

x	data.frame
cols	names of columns to work on, default is all columns
ignore	character vector of columns names to ignore
verbose	TRUE or FALSE
na_ish	Logical, default 'TRUE' which will convert NA-like strings, too
new_val	'0'

**Examples**

```
d <- data.frame(1:5, 6:10, 11:15)
d[2, 3] <- NA
d[5, 2] <- NA
d[1, 1] <- NA
print(d)
zero_na(d)
d[1, 1] <- "NA"
zero_na(d, na_ish = TRUE)
```

---

%nin%	<i>inverse of %in%</i>
-------	------------------------

---

**Description**

borrowed from Hmisc. See `nomatch = 0L) > 0L`

**Usage**

```
x %nin% table
```

**Arguments**

x is the vector of values to be matched  
table is actually a vector, to be matched against

**Value**

logical vector of length of x

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