

# Package ‘edm1’

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**Title** edm

**Version** 1.0

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**Description** What the package does (one paragraph).

**License** GPL-3

**description** Set of tools in R

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**Maintainer** First Last <first.last@example.com>

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---

append_row	<i>append_row</i>
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---

**Description**

Append the last row from dataframe to the another or same dataframe  
Append the last row from dataframe to the another or same dataframe

**Usage**

```
append_row(df_in, df, hmn = 1, na_col = c(), unique_do_not_know = NA)  
append_row(df_in, df, hmn = 1, na_col = c(), unique_do_not_know = NA)
```

**Arguments**

- df\_in is the dataframe from which the row will append to another or the same dataframe
- df is the dataframe to which the row will append
- hmn is how many time the last row will be appended
- na\_col is a vector containing the columns that won't append and will be replaced by another value (unique\_do\_not\_know)
- unique\_do\_not\_know is the value of the non appending column in the appending row

---

can_be_num	<i>can_be_num</i>
------------	-------------------

---

**Description**

Return TRUE if a variable can be converted to a number and FALSE if not

Return TRUE if a variable can be converted to a number and FALSE if not

**Usage**

```
can_be_num(x)
```

```
can_be_num(x)
```

**Arguments**

x	is the input value
---	--------------------

---

change_date	<i>change_date</i>
-------------	--------------------

---

**Description**

Allow to add to a date second-minute-hour-day-month-year

Allow to add to a date second-minute-hour-day-month-year

**Usage**

```
change_date (
  date_,
  sep_,
  day_ = NA,
  month_ = NA,
  year_ = NA,
  hour_ = NA,
  min_ = NA,
  second_ = NA,
  frmt = "snhdmy"
)
```

```
change_date (
  date_,
  sep_,
  day_ = NA,
  month_ = NA,
  year_ = NA,
  hour_ = NA,
  min_ = NA,
  second_ = NA,
  frmt = "snhdmy"
)
```

**Arguments**

date_	is the input date
sep_	is the date separator
day_	is the day to add (can be negative)
month_	is the month to add (can be negative)
year_	is the year to add (can be negative)
hour_	is the hour to add (can be negative)
min_	is the minute to add (can be negative)
second_	is the second to add (can be negative)
frmt	is the format of the input date, (default set to "snhdmy" (second, minute, hour, day, month, year), so all variable are taken in count), if you only want to work with standard date for example change this variable to "dmy"

---

closest_date	<i>closest_date</i>
--------------	---------------------

---

**Description**

return the closest dates from a vector compared to the input date

return the closest dates from a vector compared to the input date

**Usage**

```
closest_date(
  vec,
  date_,
  frmt,
  sep_ = "/",
  sep_vec = "/",
  only_ = "both",
  head = NA
)

closest_date(
  vec,
  date_,
  frmt,
  sep_ = "/",
  sep_vec = "/",
  only_ = "both",
  head = NA
)
```

**Arguments**

vec	is a vector containing the dates to be compared to the input date
date_	is the input date
frmt	is the format of the input date, (default set to "snhdmy" (second, minute, hour, day, month, year), so all variable are taken in count), if you only want to work with standard date for example change this variable to "dmy"
sep_	is the separator for the input date
sep_vec	is the separator for the dates contained in vec
head	is the number of dates that will be returned (default set to NA so all dates in vec will be returned)
only	is can be changed to "+" or "-" to repectively only return the higher dates and the lower dates (default set to "both")

---

cost_and_taxes	<i>cost and taxes</i>
----------------	-----------------------

---

**Description**

Allow to calculate basic variables related to cost and taxes from a bunch of products (elements) So put every variable you know in the following order:

Allow to calculate basic variables related to cost and taxes from a bunch of products (elements) So put every variable you know in the following order:

**Usage**

```
cost_and_taxes (
  qte = NA,
  pu = NA,
  prix_ht = NA,
  tva = NA,
  prix_ttc = NA,
  prix_tva = NA,
  pu_ttc = NA,
  adjust = NA,
  prix_d_ht = NA,
  prix_d_ttc = NA,
  pu_d = NA,
  pu_d_ttc = NA
)
```

```
cost_and_taxes (
  qte = NA,
  pu = NA,
  prix_ht = NA,
  tva = NA,
  prix_ttc = NA,
  prix_tva = NA,
  pu_ttc = NA,
```

```

adjust = NA,
prix_d_ht = NA,
prix_d_ttc = NA,
pu_d = NA,
pu_d_ttc = NA
)

```

### Arguments

qte	is the quantity of elements
pu	is the price of a single elements without taxes
prix_ht	is the duty-free price of the whole set of elements
tva	is the percentage of all taxes
prix_ttc	is the price of all the elements with taxes
prix_tva	is the cost of all the taxes
pu_ttc	is the price of a single element taxes included
adjust	is the discount percentage
prix_d_ht	is the free-duty price of an element after discount
prix_d_ttc	is the price with taxes of an element after discount
pu_d	is the price of a single element after discount and without taxes
pu_d_ttc	is the free-duty price of a single element after discount
	the function return a vector with the previous variables in the same order those that could not be calculated will be represented with NA value

---

data\_gen

*data\_gen*


---

### Description

Allo to generate in a csv all kind of data you can imagine according to what you provide

Allo to generate in a csv all kind of data you can imagine according to what you provide

### Usage

```

data_gen(
  type_ = c("number", "mixed", "string"),
  strt_l = c(0, 0, 10),
  nb_r = c(50, 10, 40),
  output = "gened.csv",
  properties = c("1-5", "1-5", "1-5"),
  type_distri = c("random", "random", "random"),
  str_source = c("a", "b", "c", "d", "e", "f", "g", "h", "i", "j", "k", "l", "m",
    "o", "p", "q", "r", "s", "t", "u", "w", "x", "y", "z"),
  round_l = c(0, 0, 0),
  sep_ = ",",
)

```

```

data_gen(
  type_ = c("number", "mixed", "string"),
  strt_l = c(0, 0, 10),
  nb_r = c(50, 10, 40),
  output = "gened.csv",
  properties = c("1-5", "1-5", "1-5"),
  type_distri = c("random", "random", "random"),
  str_source = c("a", "b", "c", "d", "e", "f", "g", "h", "i", "j", "k", "l", "m",
    "o", "p", "q", "r", "s", "t", "u", "w", "x", "y", "z"),
  round_l = c(0, 0, 0),
  sep_ = ",",
)

```

### Arguments

type_	is a vector for wich argument is a column, a column can be made of numbers ("number"), string ("string") or both ("mixed")
strt_l	is a vector containing for each column the row from which the data will begin to be generated
nb_r	is a vector containing for each column, the number of row full from generated data
output	is the name of the output csv file
properties	is linked to type_distri because it is the parameters ("min_val-max_val") for "random type", ("u-x") for the poisson distribution, ("u-d") for gaussian distribution
type_distri	is a vector which, for each column, associate a type of distribution ("random", "poisson", "gaussian"), it meas that non only the number but also the length of the string will be randomly generated according to these distribution laws
str_source	is the source (vector) from which the character creating random string are (default set to the occidental alphabet)
round_l	is a vector which, for each column containing number, associate a round value
sep_	is the separator used to write data in the csv

### Value

new generated data in addition to saving it in the output

new generated data in addition to saving it in the output

---

data\_mesdup

*data\_mesdup*

---

### Description

Allow to automatically arrange 1 dimensional data according to vector and parameters

Allow to automatically arrange 1 dimensional data according to vector and parameters

Usage

```
data_meshup(  
  data,  
  cols = NA,  
  file_ = NA,  
  sep_ = ";",  
  organisation = c(2, 1, 0),  
  unic_sep1 = "_",  
  unic_sep2 = "-"  
)  
  
data_meshup(  
  data,  
  cols = NA,  
  file_ = NA,  
  sep_ = ";",  
  organisation = c(2, 1, 0),  
  unic_sep1 = "_",  
  unic_sep2 = "-"  
)
```

Arguments

data	is the data provided (vector) each column is separated by a unic separator and each dataset from the same column is separated by another unic separator (ex: c("", c("d", "-", "e", "-", "f"), "", c("a", "a1", "-", "b", "-", "c", "c1")"_"))
cols	is the colnames of the data generated in a csv
file_	is the file to which the data will be outputed
sep_	is the separator of the csv outputed
organisation	is the way variables include themselves, for instance ,resuming precedent example, if organisation=c(1, 0) so the data output will be: d, a d, a1 e, c f, c f, c1
unic_sep1	is the unic separator between variables (default is "_")
unic_sep2	is the unic separator between datasets (default is "-")

---

date_sort	<i>date_sort</i>
-----------	------------------

---

Description

Allow to ascendely or desendely sort dates in a vector.

Usage

```
date_sort(vec, asc = F, sep = "-")
```



**Arguments**

<code>vec</code>	is the vector containing the dates.
<code>asc</code>	is a boolean variable, that if set to TRUE will sort the dates ascendely and descendely if set to FALSE
<code>sep</code>	is the separator of the date strings ex: "11-12-1998" the separator is "-"

---

<code>days_from_month</code>	<i>days_from_month</i>
------------------------------	------------------------

---

**Description**

Allow to find the number of days month from a month date, take in count leap year

Allow to find the number of days month from a month date, take in count leap year

**Usage**

```
days_from_month(date_, sep_)
```

```
days_from_month(date_, sep_)
```

**Arguments**

<code>date_</code>	is the input date
<code>sep_</code>	is the separator of the input date

---

<code>df_tuned</code>	<i>df_tuned</i>
-----------------------	-----------------

---

**Description**

Allow to return a list from a dataframe following these rules: First situation, I want the vectors from the returned list be composed of values that are separated by special values contained in a vector ex: `data.frame(c(1, 1, 2, 1), c(1, 1, 2, 1), c(1, 1, 1, 2))` will return `list(c(1, 1), c(1, 1, 1), c(1, 1, 1, 1))` or `list(c(1, 1, 2), c(1, 1, 1, 2), c(1, 1, 1, 1, 2))` if i have chosen to take in count the 2. As you noticed here the value to stop is 2 but it can be several contained in a vector Second situation: I want to return a list for every jump of 3. If i take this dataframe `data.frame(c(1, 1, 2, 1, 4, 4), c(1, 1, 2, 1, 3, 3), c(1, 1, 1, 2, 3, 3))` it will return `list(c(1, 1, 2), c(1, 4, 4), c(1, 1, 2), c(1, 3, 3), c(1, 1, 1), c(2, 3, 3))`

Allow to return a list from a dataframe following these rules: First situation, I want the vectors from the returned list be composed of values that are separated by special values contained in a vector ex: `data.frame(c(1, 1, 2, 1), c(1, 1, 2, 1), c(1, 1, 1, 2))` will return `list(c(1, 1), c(1, 1, 1), c(1, 1, 1, 1))` or `list(c(1, 1, 2), c(1, 1, 1, 2), c(1, 1, 1, 1, 2))` if i have chosen to take in count the 2. As you noticed here the value to stop is 2 but it can be several contained in a vector Second situation: I want to return a list for every jump of 3. If i take this dataframe `data.frame(c(1, 1, 2, 1, 4, 4), c(1, 1, 2, 1, 3, 3), c(1, 1, 1, 2, 3, 3))` it will return `list(c(1, 1, 2), c(1, 4, 4), c(1, 1, 2), c(1, 3, 3), c(1, 1, 1), c(2, 3, 3))`

**Usage**

```
df_tuned(df, val_to_stop, index_rc = NA, included = "yes")

df_tuned(df, val_to_stop, index_rc = NA, included = "yes")
```

**Arguments**

<code>df</code>	is the input data.frame
<code>val_to_stop</code>	is the vector containing the values to stop
<code>index_rc</code>	is the value for the jump (default set to NA so default will be first case)
<code>included</code>	is if the values to stop has to be also returned in the vectors (defaultn set to "yes")

---

<code>diff_xlsx</code>	<i>diff_xlsx</i>
------------------------	------------------

---

**Description**

Allow to see the difference between two datasets and output it into an xlsx file. If the dimensions of the new datasets are bigger than the old one, only the matching cells will be compared, if the dimensions of the new one are lower than the old one, there will be an error.

Allow to see the difference between two datasets and output it into an xlsx file

**Usage**

```
diff_xlsx(
  file_,
  sht,
  v_old_begin,
  v_old_end,
  v_new_begin,
  v_new_end,
  df2 = NA,
  overwrite = T,
  color_ = "red",
  pattern = "",
  output = "out.xlsx",
  new_val = T,
  pattern_only = T
)
```

```
diff_xlsx(
  file_,
  sht,
  v_old_begin,
  v_old_end,
  v_new_begin,
  v_new_end,
  df2 = NA,
  overwrite = T,
```

```

    color_ = "red",
    pattern = "",
    output = "out.xlsx",
    new_val = T,
    pattern_only = T
  )

```

### Arguments

file_	is the file where the data is
sht	is the sheet where the data is
v_old_begin	is the corrdinate (row, column) where the data to be compared starts
v_old_end	is the same but for its end
v_new_begin	is the coordinates where the comparator data starts
v_new_end	is the same but for its end
df2	is optional, if the comparator dataset is directly a dataframe
overwrite	allow to overwrite differences is (set to T by default)
color_	is the color the differences will be outputed
pattern	is the pattern that will be added to the differences if overwritten is set to TRUE
output	is the name of the outputed xlsx (can be set to NA if no output)
new_val	if overwrite is TRUE, then the differences will be overwritten by the comparator data
pattern_only	will cover differences by pattern if overwritten is set to TRUE

---

file\_rec

*file\_rec*


---

### Description

Allow to get all the files recursively from a path according to an end and start depth value. If you want to have an other version of this function that uses a more sophisticated algorith (which can be faster), check file\_rec2. Depth example: if i have dir/dir2/dir3, dir/dir2b/dir3b, i have a depth equal to 3

### Usage

```
file_rec(xmax, xmin = 1, pathc = ".")
```

### Arguments

xmax	is the end depth value
xmin	is the start depth value
pathc	is the reference path

---

file\_rec2

*file\_rec2*


---

### Description

Allow to find the directories and the subdirectories with a specified end and start depth value from a path. This function might be more powerfull than file\_rec because it uses a custom algorythm that does not nee to perform a full recursive search before tuning it to only find the directories with a good value of depth. Depth example: if i have dir/dir2/dir3, dir/dir2b/dir3b, i have a depth equal to 3

### Usage

```
file_rec2(xmax, xmin = 1, pathc = ".")
```

### Arguments

xmax	is the depth value
xmin	is the minimum value of depth
pathc	is the reference path, from which depth value is equal to 1

---

format\_date

*formate\_date*


---

### Description

Allow to convert xx-month-xxxx date type to xx-xx-xxxx

Allow to convert xx-month-xxxx date type to xx-xx-xxxx

### Usage

```
format_date(f_dialect, sentc, sep_in = "-", sep_out = "-")
```

```
format_date(f_dialect, sentc, sep_in = "-", sep_out = "-")
```

### Arguments

f_dialect	are the months from the language of which the month come
sentc	is the date to convert
sep_in	is the separator of the dat input (default is "-")
sep_out	is the separator of the converted date (default is "-")

---

get\_rec

get\_rec

---

### Description

Allow to get the value of directorie depth from a path.

### Usage

```
get_rec(pathc = ".")
```

### Arguments

pathc	is the reference path example: if i have dir/dir2/dir3, dir/dir2b/dir3b, i have a depth equal to 3
-------	--

---

insert\_df

insert\_df

---

### Description

Allow to insert dataframe into another dataframe according to coordinates (row, column) from the dataframe that will be inserted

Allow to insert dataframe into another dataframe according to coordinates (row, column) from the dataframe that will be inserted

### Usage

```
insert_df(df_in, df_ins, ins_loc)
```

```
insert_df(df_in, df_ins, ins_loc)
```

### Arguments

df_in	is the dataframe that will be inserted
df_ins	is the dataset to be inserted ins_loc is a vector containg two parameters (row, column) of the begining for the insertion
ins_loc	is a vector containg two parameters (row, column) of the begining for the insertion

---

letter_to_nb	<i>letter_to_nb</i>
--------------	---------------------

---

**Description**

Allow to get the number of a spreadsheet based column by the letter ex: AAA = 703

Allow to get the number of a spreadsheet based column by the letter ex: AAA = 703

**Usage**

```
letter_to_nb(letter)
```

```
letter_to_nb(letter)
```

**Arguments**

letter            is the letter (name of the column)

---

list_files	<i>list_files</i>
------------	-------------------

---

**Description**

A list.files() based function addressing the need of listing the files with extension a or or extension b ...

**Usage**

```
list_files(patternc, pathc = ".")
```

**Arguments**

patternc            is a vector containing all the extensions you want

pathc                is the path, can be a vector of multiple path because list.files() supports it.

---

match\_n

match\_n

---

### Description

Allow to get the indexes for the nth occurrence of a value in a vector. Example: `c(1, 2, 3, 1, 2)`, the first occurrence of 1 and 2 is at index 1 and 2 respectively, but the second occurrence is respectively at the 4th and 5th index.

### Usage

```
match_n(vec, mc, n = 1, wnb = "#####")
```

### Arguments

<code>vec</code>	is the input vector <code>mc</code> is a vector containing the values you want to get the index for the nth occurrence in <code>vec</code>
<code>wnb</code>	is a string you are sure is not in <code>mc</code>
<code>n</code>	the value of the occurrence

---

match\_n2

match\_n2

---

### Description

Allow to get the indexes for the nth occurrence of a value in a vector. Example: `c(1, 2, 3, 1, 2)`, the first occurrence of 1 and 2 is at index 1 and 2 respectively, but the second occurrence is respectively at the 4th and 5th index.

### Usage

```
match_n2(vec, mc, n, wnb = "#####")
```

### Arguments

<code>vec</code>	is the input vector <code>mc</code> is a vector containing the values you want to get the index for the nth occurrence in <code>vec</code>
<code>n</code>	is a vector containing the occurrences for each value in <code>mc</code> so if I have <code>mc &lt;- c(3, 27)</code> and <code>n &lt;- c(1, 2)</code> , I want the first occurrence for 3 and the second for 27 in <code>vec</code> . If the length of <code>n</code> is inferior of the length of <code>mc</code> , <code>n</code> will extend with its last value as new arguments. It means that if <code>mc &lt;- c(3, 27)</code> but <code>n &lt;- c(1)</code> so <code>n</code> will extend to <code>c(1, 1)</code> , so we will get the first occurrence of 3 and 27 in <code>vec</code> .
<code>wnb</code>	is a string you are sure is not in <code>mc</code>

---

multitud

*multitud*


---

### Description

From a list containing vectors allow to generate a vector following this rule: `list(c("a", "b"), c("1", "2"), c("A", "Z", "E")) -> c("a1A", "a2A", "b1A", "b2A", "a1Z", ...)`

From a list containing vectors allow to generate a vector following this rule:

### Usage

```
multitud(l, sep_ = "")
```

```
multitud(l, sep_ = "")
```

### Arguments

`l` is the list

`sep_` is the separator between elements (default is set to "" as you see in the example)

### Details

`list(c("a", "b"), c("1", "2"), c("A", "Z", "E")) -> c("a1A", "a2A", "b1A", "b2A", "a1Z", ...)`

---

nb\_to\_letter

*nb\_to\_letter*


---

### Description

Allow to get the letter of a spreadsheet based column by the number ex: 703 = AAA

Allow to get the letter of a spreadsheet based column by the number ex: 703 = AAA

### Usage

```
nb_to_letter(x)
```

```
nb_to_letter(x)
```

### Arguments

`x` is the number of the column



---

pattern\_generator    *pattern\_generator*

---

### Description

Allow to create patterns which have a part that is varying randomly each time.

Allow to create patterns which have a part that is varying aeach time randomly

### Usage

```
pattern_generator(base_, from_, lngth, hmn = 1, after = 1)
```

```
pattern_generator(base_, from_, lngth, hmn = 1, after = 1)
```

### Arguments

base_	is the pattern that will be kept
from_	is the vector from which the element of the varying part will be generated
hmn	is how many of varying pattern from the same base will be created
after	is set to 1 by default, it means that the varying part will be after the fixed part, set to 0 if you want the varying part to be before

---

pattern\_gettr    *pattern\_gettr*

---

### Description

Search for pattern(s) contained in a vector in another vector and return a list containing matched one (first index) and their position (second index) according to these rules: First case: Search for patterns strictly, it means that the searched pattern(s) will be matched only if the patterns contained in the vector that is beeing explored by the function are present like this c("pattern\_searched", "other", ..., "pattern\_searched") and not as c("other\_thing pattern\_searched other\_thing", "other", ..., "pattern\_searched other\_thing") Second case: It is the opposite to the first case, it means that if the pattern is partially present like in the first position and the last, it will be considered like a matched pattern

Search for pattern(s) contained in a vector in another vector and return a list containing matched one (first index) and their position (second index) according to these rules:

### Usage

```
pattern_gettr(
  word_,
  vct,
  occ = c(1),
  strict,
  btwn,
  all_in_word = "yes",
  notatall = "###"
```

```

)

pattern_gettr(
  word_,
  vct,
  occ = c(1),
  strict,
  btwn,
  all_in_word = "yes",
  notatall = "###"
)

```

### Arguments

<code>word_</code>	is the vector containing the patterns
<code>vct</code>	is the vector being searched for patterns
<code>occ</code>	a vector containing the occurrence of the pattern in <code>word_</code> to be matched in the vector being searched, if the occurrence is 2 for the <code>nth</code> pattern in <code>word_</code> and only one occurrence is found in <code>vct</code> so no pattern will be matched, put "forever" to no longer depend on the occurrence for the associated pattern
<code>strict</code>	a vector containing the "strict" condition for each <code>nth</code> vector in <code>word_</code> ("strict" is the string to activate this option)
<code>btwn</code>	is a vector containing the condition ("yes" to activate this option) meaning that if "yes", all elements between two matched pattern in <code>vct</code> will be returned, so the patterns you enter in <code>word_</code> have to be in the order you think it will appear in <code>vct</code>
<code>all_in_word</code>	is a value (default set to "yes", "no" to activate this option) that, if activated, won't authorize a previous matched pattern to be matched again
<code>notatall</code>	is a string that you are sure is not present in <code>vct</code> REGEX can also be used as pattern

### Details

First case: Search for patterns strictly, it means that the searched pattern(s) will be matched only if the patterns contained in the vector that is being explored by the function are present like this `c("pattern_searched", "other", ..., "pattern_searched")` and not as `c("other_thing pattern_searched other_thing", "other", ..., "pattern_searched other_thing")` Second case: It is the opposite to the first case, it means that if the pattern is partially present like in the first position and the last, it will be considered like a matched pattern

---

<code>pattern_tuning</code>	<i>pattern_tuning</i>
-----------------------------	-----------------------

---

### Description

Allow to tune a pattern very precisely and output a vector containing its variations `n` times.

Allow to tune a pattern very precisely

**Usage**

```
pattern_tuning(pattnr, spe_nb, spe_l, exclude_type, hmn = 1, rg = c(0, 0))

pattern_tuning(pattnr, spe_nb, spe_l, exclude_type, hmn = 1, rg = c(0, 0))
```

**Arguments**

pattnr	is the character that will be tuned
spe_nb	is the number of new character that will be replaced
spe_l	is the source vector from which the new characters will be replace old ones
exclude_type	is character that won't be replaced
hmh	is how many output the function will return
rg	is a vector with two parameters (index of the first letter that will be replaced, index of the last letter that will be replaced) default is set to all the letters from the source pattern

---

see\_df

---

see\_df

---

**Description**

Allow to return a dataframe with TRUE cells where the condition entered are respected and FALSE where these are not

Allow to return a dataframe with TRUE cells where the condition entered are respected and FALSE where these are not

**Usage**

```
see_df(df, condition_l, val_l)

see_df(df, condition_l, val_l)
```

**Arguments**

df	is the input dataframe
condition_l	is the vector of the possible conditions ("==", ">", "<", "!=", "%")
val_l	is a list with the vector of values related to condition_l (so the values has to be placed in the same order)
conjunction_l	contains the   or & conjunctions, so if the length of condition_l is equal to 3, there will be 2 conjunctions. If the length of conjunction_l is inferior to the length of condition_l minus 1, conjunction_l will match its goal length value with its last argument as the last arguments. For example, c("&", " ", "&") with a goal length value of 5 -> c("&", " ", "&", "&", "&")

**Examples**

```
see_df(df, c("%%", "=="), list(c(2, 11), c(3)), list("|")) will return all the values that
```

---

see_file	<i>see_file</i>
----------	-----------------

---

**Description**

Allow to get the filename or its extension

Allow to get the filename or its extension

**Usage**

```
see_file(string_, index_ext = 1, ext = T)
```

```
see_file(string_, index_ext = 1, ext = T)
```

**Arguments**

string_	is the input string
index_ext	is the occurrence of the dot that separates the filename and its extension
ext	is a boolean that if set to TRUE, will return the file extension and if set to FALSE, will return filename

---

see_idx	<i>see_idx</i>
---------	----------------

---

**Description**

Allow to find the indexes of the elements of the first vector in the second. If the element(s) is not found, the element returned at the same index will be "FALSE".

**Usage**

```
see_idx(v1, v2, exclude_val = "#####", no_more = F)
```

**Arguments**

v1	is the first vector
v2	is the second vector

---

see\_inside

see\_inside

---

## Description

Return a list containing all the column of the files in the current directory with a chosen file extension and its associated file and sheet if xlsx. For example if i have 2 files "out.csv" with 2 columns and "out.xlsx" with 1 column for its first sheet and 2 for its second one, the return will look like this: c(column\_1, column\_2, column\_3, column\_4, column\_5, unique\_separator, "1-2-out.csv", "3-3-sheet\_1-out.xlsx", 4-5-sheet\_2-out.xlsx)

Return a list containing all the column of the files in the current directory with a chosen file extension and its associated file and sheet if xlsx. For example if i have 2 files "out.csv" with 2 columns and "out.xlsx" with 1 column for its first sheet and 2 for its second one, the return will look like this: c(column\_1, column\_2, column\_3, column\_4, column\_5, unique\_separator, "1-2-out.csv", "3-3-sheet\_1-out.xlsx", 4-5-sheet\_2-out.xlsx)

## Usage

```
see_inside(pattern_, path_ = ".", sep_ = c(", "), unique_sep = "#####", rec = F)
```

```
see_inside(pattern_, path_ = ".", sep_ = c(", "), unique_sep = "#####", rec = F)
```

## Arguments

- |            |   |
|------------|---|
| sep_       | is a vector containing the separator for each csv type file in order following the operating system file order, if the vector does not match the number of the csv files found, it will assume the separator for the rest of the files is the same as the last csv file found. It means that if you know the separator is the same for all the csv type files, you just have to put the separator once in the vector. |
| unique_sep | is a pattern that you know will never be in your input files  |
| rec        | allow to get files recursively  |
|            | If x is the return value, to see all the files name, position of the columns and possible sheet name associated with, do the following: <code>print(x[(grep(unique_sep, x)+1):length(x)])</code> If you just want to see the columns do the following: <code>print(x[1:(grep(unique_sep, x) - 1)])</code>   |
| pattern    | is a vector containin the file extension of the spreadsheets ("xlsx", "csv"...)   |
| path       | is the path where are located the files   |

## Examples

```
print(x[(grep(unique_sep, x)[1]+1):length(x)])
If you just want to see the columns do the following:
print(x[1:(grep(unique_sep, x) - 1)])
```

---

unique_pos	<i>unique_pos</i> Allow to find indexes of the unique values from a vector.
------------	---

---

### Description

Return the indexes of the first unique values from a vector

### Usage

```
unique_pos(vec)
```

```
unique_pos(vec)
```

### Arguments

vec	is the input vector
-----	---------------------

---

until_stnl	<i>until_stnl</i>
------------	-------------------

---

### Description

Maxes a vector to a chosen length ex: if i want my vector c(1, 2) to be 5 of length this function will return me: c(1, 2, 1, 2, 1)

Maxes a vector to a chosen length

### Usage

```
until_stnl(vec1, goal)
```

```
until_stnl(vec1, goal)
```

### Arguments

vec1	is the input vector
goal	is the length to reach

### Details

ex: if i want my vector c(1, 2) to be 5 of length this function will return me: c(1, 2, 1, 2, 1)

---

val_replacer	<i>val_replacer</i>
--------------	---------------------

---

**Description**

Allow to replace value from dataframe to another one.

Allow to replace value from dataframe to another one.

**Usage**

```
val_replacer(df, val_replaced = F, val_replacor = T, df_rpt = NA)
```

```
val_replacer(df, val_replaced = F, val_replacor = T, df_rpt = NA)
```

**Arguments**

df is the input dataframe

val\_replaced is a vector of the value(s) to be replaced

val\_replacor is the value that will replace val\_replaced

df\_rpt is the replacement matrix and has to be the same dimension as df. Only the indexes that are equal to TRUE will be authorized indexes for the values to be replaced in the input matrix

---

vec_in_df	<i>vec_in_df</i>
-----------	------------------

---

**Description**

Allow to see if vectors are present in a dataframe ex: 1, 2, 1 3, 4, 1 1, 5, 8 the vector c(4, 1) with the coefficient 1 and the start position at the second column is contained in the dataframe

Allow to see if vectors are present in a dataframe

**Usage**

```
vec_in_df(df_, vec_l, coeff_, strt_l, distinct = "NA")
```

```
vec_in_df(df_, vec_l, coeff_, strt_l, distinct = "NA")
```

**Arguments**

df\_ is the input dataframe

vec\_l is a list the vectors

coeff\_ is the related coefficient of the vector

strt\_l is a vector containing the start position for each vector

**Details**

ex: 1, 2, 1 3, 4, 1 1, 5, 8

the vector c(4, 1) with the coefficient 1 and the start position at the second column is contained in the dataframe

---

vlookup_df	<i>vlookup_df</i>
------------	-------------------

---

**Description**

Allow to perform a vlookup on a dataframe

Allow to perform a vlookup on a dataframe

**Usage**

```
vlookup_df(df, v_id, col_id = 1, included_col_id = "yes")
```

```
vlookup_df(df, v_id, col_id = 1, included_col_id = "yes")
```

**Arguments**

df is the input dataframe

v\_id is a vector containing the ids

col\_id is the column that contains the ids (default is equal to 1)

included\_col\_id  
is if the result should return the col\_id (default set to yes)



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