2657 Functions

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Part I

Function Descriptions and Examples

concat.split

The concat.split function takes a column with multiple values, splits the values into a list or into separate columns, and returns a new data.frame.

Arguments

- data: the source data.frame.
- split.col: the variable that needs to be split; can be specified either by the column number or the variable name.
- to.list: logical; should the split column be returned as a single variable list (named "original-variable_list") or multiple new variables? If to.list is TRUE, the mode argument is ignored and a list of the original values are returned.
- mode: can be either binary or value (where binary is default and it recodes values to 1 or NA, like Boolean, but without assuming 0 when data is not available).
- sep: the character separating each value (defaults to ",").
- drop.col: logical (whether to remove the original variable from the output or not; defaults to TRUE).

Examples

First load some data from a CSV stored at github. The URL is an HTTPS, so we need to use getURL from RCurl.

```
require(RCurl)
baseURL = c("https://raw.github.com/mrdwab/2657-R-Functions/master/")
temp = getURL(paste0(baseURL, "data/concatenated-cells.csv"))
concat.test = read.csv(textConnection(temp))
rm(temp)

# How big is the dataset?
dim(concat.test)

[1] 48 4

# Just show me the first few rows
head(concat.test)
```

```
Hates
   Name
           Likes
                                   Siblings
   Boyd 1,2,4,5,6 Reynolds , Albert , Ortega
1
2 Rufus 1,2,4,5,6 Cohen , Bert , Montgomery 1;2;3;4;
  Dana 1,2,4,5,6
                                    Pierce
4 Carole 1,2,4,5,6 Colon , Michelle , Ballard
                                                1:4:
5 Ramona 1,2,5,6
                           Snyder , Joann ,
                                              1;2;3;
6 Kelley 1,2,5,6
                           James , Roxanne ,
                                              1;4;
Notice that the data have been entered in a very silly manner. Let's split it up!
# Load the function!
# require(RCurl)
# baseURL = c("https://raw.github.com/mrdwab/2657-R-Functions/master/")
source(textConnection(getURL(paste0(baseURL, "scripts/concat.split.R"))))
# Split up the second column, selecting by column number
head(concat.split(concat.test, 2))
   Name
            Likes
                                   Siblings
                                              Hates Likes_1 Likes_2 Likes_3
   Boyd 1,2,4,5,6 Reynolds , Albert , Ortega
                                              2;4;
                                                      1
                                                                  1
  Rufus 1,2,4,5,6 Cohen , Bert , Montgomery 1;2;3;4;
                                                          1
                                     Pierce
  Dana 1,2,4,5,6
                                               2;
                                                          1
                                                                 1
                                                                         NA
4 Carole 1,2,4,5,6 Colon , Michelle , Ballard
                                                         1
                                                                        NA
                                               1;4;
                                                                 1
5 Ramona 1,2,5,6
                          Snyder , Joann ,
                                            1;2;3;
                                                         1
                                                                        NA
                          James , Roxanne ,
6 Kelley 1,2,5,6
                                              1;4;
 Likes_4 Likes_5 Likes_6
1
       1
            1
                       1
2
       1
               1
3
       1
               1
4
               1
                      1
       1
5
      NA
               1
                       1
6
      NA
               1
# ... or by name, and drop the offensive first column
head(concat.split(concat.test, "Likes", drop.col=TRUE))
   Name
                                     Hates Likes_1 Likes_2 Likes_3 Likes_4
                          Siblings
   Boyd Reynolds , Albert , Ortega
                                                               NA
                                                                        1
                                    2;4;
                                           1
                                                        1
2 Rufus Cohen, Bert, Montgomery 1;2;3;4;
                                                1
                                                        1
                                                               NA
                                                1
                                                        1
                                                               NA
   Dana
                           Pierce
                                      2;
                                                       1
4 Carole Colon , Michelle , Ballard
                                      1;4;
                                               1
                                                               NA
                                                                        1
                                               1
                                                       1
5 Ramona
                Snyder , Joann , 1;2;3;
                                                               NA
                                                                       NΑ
6 Kelley
                 James , Roxanne ,
                                    1;4;
                                               1
                                                       1
                                                              NA
                                                                       NA
 Likes_5 Likes_6
1
      1
             1
2
       1
               1
3
       1
               1
4
       1
               1
5
       1
               1
       1
               1
# The "Hates" column uses a different separator:
head(concat.split(concat.test, "Hates", sep=";", drop.col=TRUE))
                                   Siblings Hates_1 Hates_2 Hates_3 Hates_4
   Boyd 1,2,4,5,6 Reynolds , Albert , Ortega
                                                NA
                                                         1
                                                                NΑ
                                                                         1
```

```
1 1
NA 1
2 Rufus 1,2,4,5,6 Cohen , Bert , Montgomery
                                                            1
                                                                   1
                                  Pierce
3 Dana 1,2,4,5,6
                                                            NΑ
                                                                  NA
4 Carole 1,2,4,5,6 Colon , Michelle , Ballard
                                            1
                                                   NA
                                                            NA
                                                                   1
5 Ramona 1,2,5,6 Snyder, Joann,
                                                    1
                                                            1
                                                                   NA
                        James , Roxanne ,
6 Kelley 1,2,5,6
                                             1
                                                     NA
                                                            NA
                                                                   1
# Retain the original values
head(concat.split(concat.test, 2, mode="value", drop.col=TRUE))
                                  Hates Likes 1 Likes 2 Likes 3 Likes 4
                        Siblings
1 Boyd Reynolds , Albert , Ortega
                                            1
                                                     2
                                                           NA
                                  2;4;
2 Rufus Cohen, Bert, Montgomery 1;2;3;4;
                                                     2
                                                           NA
                                             1
  Dana
                          Pierce
                                             1
                                                           NA
                                                                   4
                                   2;
4 Carole Colon , Michelle , Ballard
                                   1;4;
                                            1
                                                     2
                                                           NA
                                                                   4
5 Ramona
                                            1
                                                     2
                                                           NA
               Snyder , Joann , 1;2;3;
                                                                  NA
6 Kelley
               James , Roxanne ,
                                            1
                                                   2
                                                          NA
                                                                  NA
                                1;4;
 Likes_5 Likes_6
     5
              6
1
      5
2
              6
      5
              6
3
4
       5
              6
5
       5
              6
       5
              6
6
# Let,s try splitting some strings... Same syntax
head(concat.split(concat.test, 3, drop.col=TRUE))
                 Hates Siblings_1 Siblings_2 Siblings_3
   Name
           Likes
  Boyd 1,2,4,5,6 2;4; Reynolds Albert
                                                Ortega
                                      Bert Montgomery
2 Rufus 1,2,4,5,6 1;2;3;4; Cohen
3 Dana 1,2,4,5,6 2;
                           Pierce
                                       <NA>
                                                <NA>
                            Colon Michelle
4 Carole 1,2,4,5,6
                   1;4;
                                               Ballard
                          Snyder
5 Ramona 1,2,5,6
                  1;2;3;
                                    Joann
                                               <NA>
                            James
6 Kelley 1,2,5,6 1;4;
                                     Roxanne
                                                  <NA>
# Split up the "Likes column" into a list variable; retain original column
head(concat.split(concat.test, 2, to.list=TRUE, drop.col=FALSE))
           Likes
                                 Siblings
                                            Hates
                                                    Likes_list
  Boyd 1,2,4,5,6 Reynolds , Albert , Ortega
                                           2;4; 1, 2, 4, 5, 6
2 Rufus 1,2,4,5,6 Cohen , Bert , Montgomery 1;2;3;4; 1, 2, 4, 5, 6
3 Dana 1,2,4,5,6
                                  Pierce
                                             2; 1, 2, 4, 5, 6
4 Carole 1,2,4,5,6 Colon , Michelle , Ballard
                                           1;4; 1, 2, 4, 5, 6
                                         1;2;3; 1, 2, 5, 6
5 Ramona 1,2,5,6
                         Snyder , Joann ,
6 Kelley 1,2,5,6
                        James , Roxanne ,
                                                    1, 2, 5, 6
                                           1;4;
# View the structure of the output for the first 10 rows to verify
# that the new column is a list; note the difference between "Likes"
# and "Likes_list".
str(concat.split(concat.test, 2, to.list=TRUE, drop.col=FALSE)[1:10, c(2, 5)])
'data.frame': 10 obs. of 2 variables:
 $ Likes : Factor w/ 5 levels "1,2,3,4,5","1,2,4,5",..: 3 3 3 3 5 5 3 3 3 4
 $ Likes_list:List of 10
 ..$: num 12456
 ..$: num 1 2 4 5 6
```

Advanced Usage

It is also possible to use concat.split to split multiple columns at once. This can be done in stages, or it can be all wrapped in nested statements, as follows:

In the example above (working from the inside of the function outwards):

- First, lapply(2:ncol(concat.test), ...) splits the columns of the data.frame into a list.
- Second, lapply(lapply(...)) does the splitting work.
 - Note the use of sep="; |," to match multiple separators on which to split; if further separators are required, they can be specified by using the pipe symbol (|) with no leading or trailing spaces.
- Finally, do.call(cbind, ...) is evaluated last, "binding" the data together by columns. In this case, the data being bound together is the first column from the concat.test dataset, and the splitted output of the remaining columns.

Alternatively, a similar approach can be taken using the function dfcols.list (see the "Snippets and Tips" section of this manual for the dfcols.list function).

```
# Show just the first few lines, Boolean mode
head(do.call(cbind, c(concat.test[1],
                      lapply(dfcols.list(concat.test[-1]),
                             concat.split, split.col=1, drop=TRUE, sep=";|,"))))
   Name Likes_1 Likes_2 Likes_3 Likes_4 Likes_5 Likes_6 Siblings_1 Siblings_2
1
   Boyd
               1
                      1
                              NA
                                       1
                                               1
                                                        1
                                                            Reynolds
                                                                         Albert
                              NA
2 Rufus
               1
                                                1
                                                               Cohen
                                                                           Bert
                       1
                                       1
                                                        1
3
   Dana
               1
                       1
                              NA
                                       1
                                                1
                                                        1
                                                              Pierce
                                                                           < NA >
4 Carole
               1
                              NA
                                                1
                                                       1
                       1
                                                              Colon
                                                                       Michelle
5 Ramona
               1
                              NA
                                      NA
                                                              Snyder
                       1
                                                1
                                                        1
                                                                          Joann
               1
                              NA
                                      NA
                                                        1
                                                               James
                                                                        Roxanne
6 Kelley
                       1
                                                1
  Siblings_3 Hates_1 Hates_2 Hates_3 Hates_4
                  NA
     Ortega
                           1
2 Montgomery
                  1
                           1
                                   1
                                           1
        <NA>
                  NA
                          1
                                  NA
                                          NA
3
                  1
                                  NA
4
    Ballard
                          NA
                                           1
5
        <NA>
                  1
                          1
                                  1
                                          NA
6
        <NA>
                 1
                          NA
                                           1
                                  NA
```

REFERENCES 7

```
# Show just the first few lines, value mode
head(do.call(cbind, c(concat.test[1],
                     lapply(dfcols.list(concat.test[-1]),
                            concat.split, split.col=1, drop=TRUE,
                            sep=";|,", mode="value"))))
   Name Likes_1 Likes_2 Likes_3 Likes_4 Likes_5 Likes_6 Siblings_1 Siblings_2
                                     4
                                                        Reynolds
   Boyd 1
                      2
                            NA
                                             5
                                                    6
                                                                     Albert
1
                                                           Cohen
2 Rufus
              1
                      2
                            NA
                                     4
                                             5
                                                    6
                                                                       Bert
                      2
  Dana
              1
                            NA
                                     4
                                             5
                                                    6
                                                          Pierce
                                                                       <NA>
4 Carole
              1
                      2
                            NA
                                     4
                                             5
                                                    6
                                                           Colon
                                                                   Michelle
5 Ramona
              1
                      2
                            NA
                                    NA
                                             5
                                                   6
                                                                      Joann
                                                          Snyder
                      2
                            NA
                                    NA
                                             5
                                                    6
6 Kelley
             1
                                                            James
                                                                    Roxanne
 Siblings_3 Hates_1 Hates_2 Hates_3 Hates_4
                        2
     Ortega
                 NA
                                NA
                          2
                                3
                                         4
2 Montgomery
                 1
                        2
3
       <NA>
                 NA
                                NA
                                        NA
4
    Ballard
                 1
                         NA
                                NA
                                         4
5
                         2
                                3
       <NA>
                  1
                                        NA
       <NA>
6
                  1
                        NA
                                NA
                                         4
# Show just the first few lines, list output mode
head(do.call(cbind, c(concat.test[1],
                     lapply(dfcols.list(concat.test[-1]),
                            concat.split, split.col=1, drop=TRUE,
                            sep=";|,", to.list=TRUE))))
           Likes_list
   Name
                                Siblings_list Hates_list
   Boyd 1, 2, 4, 5, 6 Reynolds, Albert, Ortega
2 Rufus 1, 2, 4, 5, 6 Cohen, Bert, Montgomery 1, 2, 3, 4
   Dana 1, 2, 4, 5, 6
                                       Pierce
4 Carole 1, 2, 4, 5, 6 Colon, Michelle, Ballard
                                                    1, 4
5 Ramona 1, 2, 5, 6
                               Snyder, Joann
                                                 1, 2, 3
6 Kelley
           1, 2, 5, 6
                               James, Roxanne
                                                    1, 4
```

References

See: http://stackoverflow.com/q/10100887/1270695

df.sorter

The df.sorter function allows you to sort a data.frame by columns or rows or both. You can also quickly subset data columns by using the var.order argument.

Arguments

- data: the source data.frame.
- var.order: the new order in which you want the variables to appear.
 - Defaults to names (data), which keeps the variables in the original order.
 - Variables can be referred to either by a vector of their index numbers or by a vector of the
 variable name; partial name matching also works, but requires that the partial match identifies
 similar columns uniquely (see examples).
 - Basic subsetting can also be done using var.order simply by omitting the variables you want to drop.
- col.sort: the columns within which there is data that need to be sorted.
 - Defaults to NULL, which means no sorting takes place.
 - Variables can be referred to either by a vector of their index numbers or by a vector of the variable names; full names must be provided.
- at.start: Should the pattern matching be from the start of the variable name? Defaults to "TRUE".

NOTE: If you are sorting both by variables and within the columns, the col.sort order should be based on the location of the columns in the *new* data.frame, not the original data.frame.

Examples

```
# Load the function!
require(RCurl)
baseURL = c("https://raw.github.com/mrdwab/2657-R-Functions/master/")
source(textConnection(getURL(paste0(baseURL, "scripts/df.sorter.R"))))
# Make up some data
set.seed(1)
dat = data.frame(id = rep(1:5, each=3), times = rep(1:3, 5),
```

```
measure1 = rnorm(15), score1 = sample(300, 15),
               code1 = replicate(15, paste(sample(LETTERS[1:5], 3),
                                        sep="", collapse="")),
               measure2 = rnorm(15), score2 = sample(150:300, 15),
               code2 = replicate(15, paste(sample(LETTERS[1:5], 3),
                                        sep="", collapse="")))
# Preview your data
dat
  id times measure1 score1 code1 measure2 score2 code2
        1 -0.6265
                     145 DAB -0.7075
                                         299
                                              CEB
1
   1
                                         224
                                              ECD
        2 0.1836
                     180 DCB 0.3646
2
  1
        3 -0.8356
                               0.7685
                                         222 DAE
3
  1
                   148 EBA
   2
        1 1.5953
                     56 AED -0.1123
                                       175 DBA
5
   2
        2 0.3295
                     245 CEB
                               0.8811
                                         260 DAC
        3 -0.8205
                   198 EBD
                                         216 DCA
6
   2
                               0.3981
7
   3
        1
            0.4874
                     234
                          BCA
                              -0.6120
                                         300
                                              CEA
8
        2
           0.7383
                     32
                               0.3411
                                         179
                                              CAD
   3
                         CDA
9
   3
        3
           0.5758
                     212 EBC -1.1294
                                       182 BEC
10 4
        1 -0.3054 120 BED 1.4330
                                         234 CDE
11 4
       2 1.5118 239 EDB 1.9804
                                         231 CAB
        3 0.3898 188 DEB -0.3672 160 DBE
12 4
13 5
        1 -0.6212
                     226 DBA -1.0441
                                         154 EDB
        2 -2.2147
                               0.5697
14 5
                                         238 BDE
                     159
                          DAC
15 5
        3
           1.1249
                     152
                           AED -0.1351
                                         277
                                              DCE
# Change the variable order, grouping related columns
# Note that you do not need to specify full variable names,
    just enough that the variables can be uniquely identified
head(df.sorter(dat, var.order = c("id", "ti", "cod", "mea", "sco")))
 id times code1 code2 measure1 measure2 score1 score2
      1 DAB CEB -0.6265 -0.7075 145
        2 DCB ECD 0.1836 0.3646
2 1
                                       180
                                             224
3 1
        3 EBA DAE -0.8356 0.7685
                                       148
                                             222
               DBA
4 2
           AED
                      1.5953 -0.1123
                                       56
                                             175
           CEB
               DAC
                     0.3295 0.8811
5 2
        2
                                       245
                                             260
        3
           EBD DCA -0.8205 0.3981
                                       198
                                             216
# Same output, but with a more awkward syntax
head(df.sorter(dat, var.order = c(1, 2, 5, 8, 3, 6, 4, 7)))
 id times code1 code2 measure1 measure2 score1 score2
       1 DAB CEB -0.6265 -0.7075
2 1
        2 DCB ECD
                     0.1836
                             0.3646
                                             224
                                       180
3 1
        3 EBA DAE -0.8356
                             0.7685
                                       148
                                             222
4 2
           AED
               DBA
                     1.5953 -0.1123
                                       56
                                              175
           CEB
                DAC
5 2
        2
                     0.3295
                             0.8811
                                       245
                                              260
6 2
           EBD
                DCA -0.8205
                             0.3981
                                       198
                                             216
\# As above, but sorted by ,times, and then ,id,
head(df.sorter(dat, var.order = c("id", "tim", "cod", "mea", "sco"),
             col.sort = c(2, 1))
  id times code1 code2 measure1 measure2 score1 score2
        1 DAB CEB -0.6265 -0.7075
                                        145
  1
```

EXAMPLES 11

```
DBA 1.5953 -0.1123
                                         175
       1
           AED
                                     56
7
                    0.4874 -0.6120
                                          300
   3
      1
           BCA
                CEA
                                     234
10 4
      1
           BED
                CDE -0.3054 1.4330 120
                                         234
13 5
           DBA
                EDB -0.6212 -1.0441 226
                                         154
        2 DCB
                ECD 0.1836 0.3646 180
                                          224
# Drop ,measure1, and ,measure2,, sort by ,times,, and ,score1,
head(df.sorter(dat, var.order = c("id", "tim", "sco", "cod"),
            col.sort = c(2, 3))
  id times score1 score2 code1 code2
4
   2
      1
           56 175 AED DBA
      1 120
                234 BED CDE
10 4
           145 299 DAB CEB
1
 1
      1
13 5
      1
           226 154 DBA EDB
            234 300 BCA CEA
7 3
       1
8 3
        2
           32 179 CDA CAD
# As above, but using names
head(df.sorter(dat, var.order = c("id", "tim", "sco", "cod"),
            col.sort = c("times", "score1")))
  id times score1 score2 code1 code2
  2 1 56 175 AED DBA
4
                  234 BED CDE
10 4
       1
           120
           145 299 DAB CEB
1
  1
       1
13 5
       1
           226 154 DBA EDB
                  300 BCA CEA
7 3
       1
            234
        2
                  179 CDA CAD
8
   3
            32
# Just sort by columns, first by ,times, then by ,id,
head(df.sorter(dat, col.sort = c("times", "id")))
  id times measure1 score1 code1 measure2 score2 code2
      1 -0.6265 145 DAB -0.7075 299 CEB
      1 1.5953
                   56 AED -0.1123 175 DBA
7
  3
       1 0.4874
                   234 BCA -0.6120
                                      300 CEA
10 4
       1 -0.3054
                            1.4330
                  120 BED
                                      234 CDE
       1 -0.6212 226 DBA -1.0441
                                     154 EDB
13 5
        2 0.1836
                                      224 ECD
2
                   180 DCB 0.3646
head(df.sorter(dat, col.sort = c("code1"))) # Sorting by character values
  id times measure1 score1 code1 measure2 score2 code2
4
      1 1.5953 56 AED -0.1123 175 DBA
       3 1.1249
                                     277 DCE
15 5
                   152 AED -0.1351
7 3
       1 0.4874
                   234 BCA -0.6120
                                      300 CEA
10 4
       1 -0.3054
                   120 BED
                            1.4330
                                      234 CDE
        2 0.7383
   3
                   32
                        CDA
                             0.3411
                                     179 CAD
8
        2 0.3295
                                      260 DAC
5
                   245 CEB
                            0.8811
# Pattern matching anywhere in the variable name
head(df.sorter(dat, var.order= "co", at.start=FALSE))
```

	code1	code2	score1	score2
1	DAB	CEB	145	299
2	DCB	ECD	180	224
3	EBA	DAE	148	222
4	AED	DBA	56	175
5	CEB	DAC	245	260
6	EBD	DCA	198	216

To Do

• Add an option to sort ascending or descending—at the moment, not supported.

multi.freq.table

The multi.freq.table function takes a data frame containing Boolean responses to multiple response questions and tabulates the number of responses by the possible combinations of answers. In addition to tabulating the frequency (Freq), there are two other columns in the output: Percent of Responses (Pct.of.Resp) and Percent of Cases (Pct.of.Cases). Percent of Responses is the frequency divided by the total number of answers provided; this column should sum to 100%. In some cases, for instance when a combination table is generated and there are cases where a respondent did not select any option, the Percent of Responses value would be more than 100%. Percent of Cases is the frequency divided by the total number of valid cases; this column would most likely sum to more than 100% when a basic table is produced since each respondent (case) can select multiple answers, but should sum to 100% with other tables.

Arguments

- data: The multiple responses that need to be tabulated.
- sep: The desired separator for collapsing the combinations of options; defaults to "" (collapsing with no space between each option name).
- boolean: Are you tabulating boolean data (see dat examples)? Defaults to TRUE.
- factors: If you are trying to tabulate non-boolean data, and the data are not factors, you can specify the factors here (see dat2 examples).
 - Defaults to NULL and is not used when boolean = TRUE.
- NAtoO: Should NA values be converted to O.
 - Defaults to TRUE, in which case, the number of valid cases should be the same as the number of cases overall.
 - If set to FALSE, any rows with NA values will be dropped as invalid cases.
 - Only applies when boolean = TRUE.
- basic: Should a basic table of each item, rather than combinations of items, be created? Defaults to FALSE.
- dropzero: Should combinations with a frequency of zero be dropped from the final table?
 - Defaults to TRUE.
 - Does not apply when boolean = TRUE.
- clean: Should the original tabulated data be retained or dropped from the final table?
 - Defaults to TRUE.
 - Does not apply when boolean = TRUE.

Examples

Boolean Data

```
# Load the function!
require(RCurl)
baseURL = c("https://raw.github.com/mrdwab/2657-R-Functions/master/")
source(textConnection(getURL(paste0(baseURL, "scripts/multi.freq.table.R"))))
# Make up some data
set.seed(1)
dat = data.frame(A = sample(c(0, 1), 20, replace=TRUE),
                B = sample(c(0, 1, NA), 20,
                           prob=c(.3, .6, .1), replace=TRUE),
                C = sample(c(0, 1, NA), 20,
                           prob=c(.7, .2, .1), replace=TRUE),
                D = sample(c(0, 1, NA), 20,
                           prob=c(.3, .6, .1), replace=TRUE),
                E = sample(c(0, 1, NA), 20,
                           prob=c(.4, .4, .2), replace=TRUE))
# View your data
dat
  A B C D E
1 O NA 1 NA O
2 0 1 0 1 0
3 1 0 1 1 1
4 1 1 0 1 1
5 0 1 0 0 0
6 1 1 1 1 1
7 \quad 1 \quad 1 \quad 0 \quad 1 \quad 0
8 1 1 0 0 1
9 1 0 1 1 1
10 0 1 0 0 1
11 0 1 0 1 1
12 0 1 1 0 1
13 1 1 0 1 0
14 0 1 0 1 NA
15 1 0 0 1 0
16 0 0 0 0 0
17 1 0 0 0 0
18 1 1 0 1 0
19 0 0 0 0 NA
20 1 1 0 NA 0
# How many cases have "NA" values?
table(is.na(rowSums(dat)))
FALSE TRUE
  16 4
# Apply the function with all defaults accepted
multi.freq.table(dat)
```

Combn Freq Weighted.Freq Pct.of.Resp Pct.of.Cases

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```
2
                          4.167
                                       10
1
2
     Α
        1
                   1
                           2.083
                                        5
                   1
3
     В
       1
                           2.083
                                        5
    AB 1
4
                   2
                         4.167
                                        5
    C 1
5
                   1
                         2.083
6
   AD
        1
                   2
                          4.167
                                        5
7
                   4
                         8.333
   BD
         2
                                       10
                   9
                        18.750
   ABD
8
                                        15
                         4.167
9
   BE
                   2
         1
                                        5
10 ABE
                                        5
       1
                   3
                          6.250
11 BCE 1
                         6.250
                                        5
                   3
12 BDE 1
                   3
                         6.250
                                       5
13 ABDE 1
                   4
                          8.333
                                        5
14 ACDE
         2
                   8
                          16.667
                                        10
15 ABCDE
                                        5
                    5
                          10.417
# Tabulate only on variables "A", "B", and "D", with a different
# separator, keep any zero frequency values, and keeping the
# original tabulations. There are no solitary "D" responses.
multi.freq.table(dat[c(1, 2, 4)], sep="-", dropzero=FALSE, clean=FALSE)
 A B D Freq Combn Weighted.Freq Pct.of.Resp Pct.of.Cases
1 0 0 0 3
                       3
                             8.571
2 1 0 0
                              2.857
       1
            Α
                        1
3 0 1 0 3
                              8.571
                                           15
            В
                       3
      2 A-B
4 1 1 0
                       4
                             11.429
                                           10
       0
5001
           D
                        0
                              0.000
      3 A-D
                       6
6 1 0 1
                              17.143
                                           15
7 0 1 1
      3 B-D
                       6
                              17.143
                                           15
8 1 1 1 5 A-B-D
                              42.857
                      15
                                           25
# As above, but without converting "NA" to "O".
# Note the difference in the number of valid cases.
multi.freq.table(dat[c(1, 2, 4)], NAtoO=FALSE,
             sep="-", dropzero=FALSE, clean=FALSE)
 A B D Freq Combn Weighted.Freq Pct.of.Resp Pct.of.Cases
1 0 0 0 2
               2
                            6.061
                                    11.111
       1
2 1 0 0
             Α
                        1
                              3.030
                                         5.556
                       3
3 0 1 0
       3
            В
                             9.091
                                        16.667
4 1 1 0 1 A-B
                       2
                             6.061
                                        5.556
5 0 0 1 0 D
                       0
                              0.000
                                        0.000
                       6 18.182
6 1 0 1 3 A-D
                                       16.667
7 0 1 1
        3 B-D
                       6
                             18.182
                                        16.667
                    15
8 1 1 1
        5 A-B-D
                              45.455
                                        27.778
# View a basic table.
multi.freq.table(dat, basic=TRUE)
 Freq Pct.of.Resp Pct.of.Cases
A 11 22.92 55
 13
         27.08
В
                       65
```

25

40

C 5

D 11

E 8

10.42

22.92

16.67

Non-Boolean Data

```
# Make up some data
dat2 = structure(list(Reason.1 = c("one", "one", "two", "one", "two",
                              "three", "one", "one", NA, "two"),
                  Reason.3 = c("three", NA, NA, NA, NA,
                             NA, NA, "three", NA, NA)),
               .Names = c("Reason.1", "Reason.2", "Reason.3"),
              class = "data.frame",
              row.names = c(NA, -10L))
# View your data
dat2
  Reason.1 Reason.2 Reason.3
1
            two three
     one
2
      one three
                    <NA>
3
      two three
                    <NA>
                    <NA>
4
             <NA>
      one
            <NA>
                    <NA>
5
      two
6
   three
             two <NA>
7
                    <NA>
     one three
     one
8
            two three
9
     <NA> <NA> <NA>
      two <NA>
10
                    <NA>
# The following will not work.
# The data are not factored.
multi.freq.table(dat2, boolean=FALSE)
Error: Input variables must be factors. Please provide factors using the
'factors' argument or convert your data to factor before using function.
# Factor create the factors.
multi.freq.table(dat2, boolean=FALSE,
              factors = c("one", "two", "three"))
      Combos Freq Weighted.Freq Pct.of.Resp Pct.of.Cases
1
              1
                          1
                                5.882
         one 1
                                 5.882
8
                          1
                                               10
         two 2
                          2
                                11.765
12
                                               20
    onethree 2
                          4
                                23.529
15
                                               20
                                23.529
17
     threetwo 2
                           4
                                                20
                          6
                               35.294
                                                20
22 onethreetwo 2
# And, a basic table.
multi.freq.table(dat2, boolean=FALSE,
              factors = c("one", "two", "three"),
              basic=TRUE)
  Item Freq Pct.of.Resp Pct.of.Cases
1 one 5 29.41 50
2 two 6
               35.29
3 three 6
              35.29
                              60
```

EXAMPLES 17

Extended Examples

The following example is based on some data available from the University of Auckland's Student Learning Resources¹.

When the data are read into R, the factor labels are very long, which makes it difficult to see on the screen. Thus, in the first example that follows, the factor levels are first recoded before the multiple frequency tables are created. Additionally, the data for the binary information in the second example was coded in a common 1 = Yes and 2 = No format, but we need 0 = No instead, so we need to do some recoding there too before using the function.

```
# Get the data
library(foreign)
temp = "http://cad.auckland.ac.nz/file.php/content/files/slc/"
computer = read.spss(paste0(temp,
                            "computer_multiple_response.sav"),
                     to.data.frame=TRUE)
rm(temp)
# Preview
dim(computer)
[1] 100 20
names(computer)
 [1] "id"
                           "ms_excel" "ms_ppt" "ms_outlk" "ms_pub"
                "ms word"
 [7] "ms_proj"
                           "netscape" "int expl" "adobe rd" "endnote"
                "ms acc"
                "quality1" "quality2" "quality3" "quality4" "quality5"
[13] "spss"
[19] "quality6" "gender"
# First, let,s just tabulate the instructor qualities.
# Extract the relevant columns, and relevel the factors.
instructor.quality =
 computer[, grep("quali", names(computer))]
# View the existing levels.
lapply(instructor.quality, levels)[[1]]
[1] "Ability to provide practical examples"
[2] "Ability to answer questions positively"
[3] "Ability to clearly explain concepts"
[4] "Ability to instruct at a suitable pace"
[5] "Knowledge of software"
[6] "Humour"
[7] "Other"
instructor.quality = lapply(instructor.quality,
                            function(x) { levels(x) =
 list(Q1 = "Ability to provide practical examples",
       Q2 = "Ability to answer questions positively",
       Q3 = "Ability to clearly explain concepts",
       Q4 = "Ability to instruct at a suitable pace",
       Q5 = "Knowledge of Software",
       Q6 = "Humour", Q7 = "Other"); x })
# Now, apply multi.freq.table to the data.
multi.freq.table(data.frame(instructor.quality),
                 boolean=FALSE, basic=TRUE)
```

¹See: http://www.cad.auckland.ac.nz/index.php?p=spss

```
Item Freq Pct.of.Resp Pct.of.Cases
   Q1
      47
               18.077
1
2
   Q2
      59
               22.692
                              59
3
  Q3 55
              21.154
                             55
4
  Q4 43
              16.538
                              43
5 Q5 0
               0.000
                              0
        47
              18.077
                               47
6
   Q6
7
   07
       9
               3.462
                               9
list(head(multi.freq.table(data.frame(instructor.quality),
                        boolean=FALSE, sep="-")),
    tail(multi.freq.table(data.frame(instructor.quality),
                        boolean=FALSE, sep="-")))
[[1]]
  Combos Freq Weighted.Freq Pct.of.Resp Pct.of.Cases
1
      Q1
           1
                1
                             0.3846
21
      Q2
                       3
           3
                              1.1538
          2
                       2
                                               2
31
      QЗ
                              0.7692
                       2
37
      Q4
          2
                              0.7692
                                               2
         3
                       3
                                               3
39
      Q6
                              1.1538
41 Q1-Q2 8
                       16
                              6.1538
                                               8
[[2]]
             Combos Freq Weighted.Freq Pct.of.Resp Pct.of.Cases
133
         Q1-Q3-Q6-Q7
                          4
                                      1.538
                     1
141
         Q2-Q3-Q4-Q6
                      4
                                  16
                                           6.154
                                                          4
                                      1.538
1.923
1.923
151
         Q3-Q4-Q6-Q7
                     1
                                  4
                                                          1
                                  5
161
      Q1-Q2-Q3-Q4-Q6
                    1
                                                          1
                                  5
164
      Q1-Q2-Q3-Q6-Q7
                    1
                                                         1
201 Q1-Q2-Q3-Q4-Q6-Q7
                                   6
                                          2.308
# Now. let,s look at the software.
instructors.sw = computer[2:13]
# These columns are coded as 1 = Yes and 2 = No,
# so, convert to integers, and subtract two, and
# take the absolute value to convert to binary.
instructors.sw = lapply(instructors.sw,
                      function(x) abs(as.integer(x)-2))
# Apply multi.freq.table
multi.freq.table(data.frame(instructors.sw), basic=TRUE)
        Freq Pct.of.Resp Pct.of.Cases
ms word
         77
               13.975
                                48
ms_excel 48
                8.711
          55
                9.982
                                55
ms_ppt
                9.437
ms_outlk 52
                                52
                3.448
         19
                                19
ms_pub
          21
                3.811
                                21
ms_proj
ms_acc
          57
               10.345
                                57
netscape 10
                1.815
                                10
int_expl
          84
                15.245
                                84
                 8.711
                                48
         48
adobe_rd
endnote
          55
                  9.982
                                 55
          25
                  4.537
                                 25
spss
# The output here is not pretty. To get prettier (or more meaningful)
```

output, provide shorter names for the variables or use just a

REFERENCES 19

```
meaningful subset of the variables.
list(head(multi.freq.table(data.frame(instructors.sw), sep="-")),
     tail(multi.freq.table(data.frame(instructors.sw), sep="-")))
[[1]]
                                            Combn Freq Weighted.Freq Pct.of.Resp
1
                  ms_word-ms_excel-ms_ppt-ms_acc
                                                                    4
                                                                           0.7260
                                                                           1.0889
2 ms_word-ms_excel-ms_ppt-ms_outlk-ms_pub-ms_acc
                                                                    6
                                                     1
                                                                    2
3
                                         int_expl
                                                     2
                                                                           0.3630
                                                                    2
4
                                ms_word-int_expl
                                                     1
                                                                           0.3630
5
                         ms_word-ms_ppt-int_expl
                                                     1
                                                                    3
                                                                           0.5445
                                                                    3
                                                                           0.5445
6
                       ms_word-ms_outlk-int_expl
  Pct.of.Cases
2
             1
3
             2
4
             1
5
             1
             1
[[2]]
91 ms_word-ms_excel-ms_outlk-ms_pub-ms_proj-int_expl-adobe_rd-endnote-spss
             ms_word-ms_excel-ms_ppt-ms_acc-int_expl-adobe_rd-endnote-spss
                                                                                 1
93
                    ms_word-ms_outlk-ms_acc-int_expl-adobe_rd-endnote-spss
94
             {\tt ms\_word\_ms\_ppt\_ms\_outlk\_ms\_acc\_int\_expl\_adobe\_rd\_endnote\_spss}
                                                                                1
95
                      ms_word-ms_pub-ms_acc-int_expl-adobe_rd-endnote-spss
                                                                                 1
96
                    ms_outlk-ms_proj-ms_acc-int_expl-adobe_rd-endnote-spss
   Weighted.Freq Pct.of.Resp Pct.of.Cases
               9
91
                      1.633
92
               8
                       1.452
                                         1
93
               7
                       1.270
                                         1
94
               8
                       1.452
                                         1
95
               7
                                         1
                       1.270
96
               7
                                         1
                       1.270
```

References

apply shortcut for creating the Combn column in the output by Justin See: $http://stackoverflow.com/q/11348391/1270695 \ and \ http://stackoverflow.com/q/11622660/1270695$

row.extractor

The row.extractor function takes a data.frame and extracts rows with the min, median, or max values of a given variable, or extracts rows with specific quantiles of a given variable.

Arguments

- data: the source data.frame.
- extract.by: the column which will be used as the reference for extraction; can be specified either by the column number or the variable name.
- what: options are min (for all rows matching the minimum value), median (for the median row or rows), max (for all rows matching the maximum value), or all (for min, median, and max); alternatively, a numeric vector can be specified with the desired quantiles, for instance c(0, .25, .5, .75, 1)

Examples

```
# Load the function!
require(RCurl)
baseURL = c("https://raw.github.com/mrdwab/2657-R-Functions/master/")
source(textConnection(getURL(paste0(baseURL, "scripts/row.extractor.R"))))
# Make up some data
set.seed(1)
dat = data.frame(V1 = 1:50, V2 = rnorm(50),
                 V3 = round(abs(rnorm(50)), digits=2),
                 V4 = sample(1:30, 50, replace=TRUE))
# Get a sumary of the data
summary(dat)
                      V2
                                        VЗ
                                                        ۷4
       V1
                                       :0.000
                      :-2.215
                                                  Min. : 2.00
Min.
      : 1.0
                Min.
                                 \mathtt{Min}.
 1st Qu.:13.2
                1st Qu.:-0.372
                                 1st Qu.:0.347
                                                  1st Qu.: 8.25
 Median:25.5
                Median : 0.129
                                 Median :0.590
                                                  Median :13.00
Mean
       :25.5
                      : 0.100
                                 Mean
                                         :0.774
                                                        :14.80
                Mean
                                                  Mean
 3rd Qu.:37.8
                3rd Qu.: 0.728
                                 3rd Qu.:1.175
                                                  3rd Qu.:20.75
       :50.0
                Max.
                       : 1.595
                                 Max.
                                        :2.400
                                                  Max.
                                                         :29.00
\# Get the rows corresponding to the ,min, ,median, , and ,max, of ,V4,
row.extractor(dat, 4)
```

```
V1
         V2 V3 V4
28 28 -1.4708 0.00 2
47 47 0.3646 1.28 13
29 29 -0.4782 0.07 13
11 11 1.5118 2.40 29
14 14 -2.2147 0.03 29
18 18 0.9438 1.47 29
19 19 0.8212 0.15 29
50 50 0.8811 0.47 29
\# Get the ,min, rows only, referenced by the variable name
row.extractor(dat, "V4", "min")
   V1
         V2 V3 V4
28 28 -1.471 0 2
# Get the ,median, rows only. Notice that there are two rows
# since we have an even number of cases and true median
# is the mean of the two central sorted values
row.extractor(dat, "V4", "median")
          V2 V3 V4
   V1
47 47 0.3646 1.28 13
29 29 -0.4782 0.07 13
# Get the rows corresponding to the deciles of , V3,
row.extractor(dat, "V3", seq(0.1, 1, 0.1))
           V2 V3 V4
   V1
10 10 -0.30539 0.14 22
26 26 -0.05613 0.29 16
39 39 1.10003 0.37 13
41 41 -0.16452 0.54 10
30 30 0.41794 0.59 26
44 44 0.55666 0.70 5
37 37 -0.39429 1.06 21
49 49 -0.11235 1.22 14
34 34 -0.05381 1.52 19
11 11 1.51178 2.40 29
```

To Do

• Add some error checking to make sure a valid what is provided.

References

which.quantile function by cheleites See: http://stackoverflow.com/q/10256503/1270695

sample.size

The sample.size function either calculates the optimum survey sample size when provided with a population size, or the confidence interval of using a certain sample size with a given population. It can be used to generate tables (data.frames) of different combinations of inputs of the following arguments, which can be useful for showing the effect of each of these in sample size calculation.

The Arguments

- population: The population size for which a sample size needs to be calculated.
- samp.size: The sample size.
 - This argument is only used when calculating the confidence interval, and defaults to NULL.
- c.lev: The desired confidence level. Defaults to a reasonable 95%.
- c.int: The confidence interval.
 - This argument is only used when calculating the sample size.
 - If not specified when calculating the sample size, defaults to 5% and a message is provided indicating this; this is also the default action if c.int = NULL.
- what: Should the function calculate the desired sample size or the confidence interval?
 - Accepted values are "sample" and "confidence" (quoted), and defaults to "sample".
- distribution: Response distribution. Defaults to 50%, which will give you the largest sample size.

Examples

```
# Load the function!
require(RCurl)
baseURL = c("https://raw.github.com/mrdwab/2657-R-Functions/master/")
source(textConnection(getURL(pasteO(baseURL, "scripts/sample.size.R"))))
# What should our sample size be for a population of 300?
# All defaults accepted.
sample.size(population = 300)

population conf.level conf.int distribution sample.size
1 300 95 5 50 169
```

```
# What sample should we take for a population of 300
# at a confidence level of 97%?
sample.size(population = 300, c.lev = 97)
  population conf.level conf.int distribution sample.size
                     97
1
         300
                               5
                                           50
                                                      183
# What about if we change our confidence interval?
sample.size(population = 300, c.int = 2.5, what = "sample")
 population conf.level conf.int distribution sample.size
1
                     95
                             2.5
# What about if we want to determine the confidence interval
   of a sample of 140 from a population of 300? A confidence
    level of 95% is assumed.
sample.size(population = 300, samp.size = 140, what = "confidence")
  population conf.level conf.int distribution sample.size
1
                     95
                            6.06
                                           50
                                                      140
```

Advanced Usage

10.0

As the function is vectorized, it is possible to easily make tables with multiple scenarios.

```
# What should the sample be for populations of 300 to 500 by 50?
sample.size(population=c(300, 350, 400, 450, 500))
  population conf.level conf.int distribution sample.size
1
         300
                      95
                                5
                                              50
                                                          169
         350
                      95
                                 5
                                                          183
2
                                              50
                      95
                                 5
3
         400
                                              50
                                                         196
4
         450
                      95
                                 5
                                              50
                                                         207
5
         500
                      95
                                5
                                              50
                                                         217
# How does varying confidence levels or confidence intervals
# affect the sample size?
sample.size(population=300,
            c.lev=rep(c(95, 96, 97, 98, 99), times = 3),
            c.int=rep(c(2.5, 5, 10), each=5))
   population conf.level conf.int distribution sample.size
          300
                       95
                                2.5
                                               50
1
                                                           251
2
          300
                       96
                                2.5
                                               50
                                                           255
3
          300
                       97
                                               50
                                                           259
                                2.5
4
          300
                       98
                                               50
                                2.5
                                                           264
5
          300
                       99
                                2.5
                                               50
                                                           270
6
          300
                       95
                                5.0
                                               50
                                                           169
7
          300
                       96
                                5.0
                                               50
                                                           176
          300
                       97
                                5.0
                                               50
8
                                                           183
9
          300
                       98
                                5.0
                                               50
                                                           193
10
          300
                       99
                                5.0
                                               50
                                                           207
11
          300
                       95
                              10.0
                                               50
                                                           73
12
          300
                       96
                              10.0
                                               50
                                                           78
13
          300
                       97
                              10.0
                                               50
                                                           85
          300
                              10.0
14
                       98
                                               50
                                                           93
```

REFERENCES 25

	population	conf.level	conf.int	distribution	sample.size
1	300	95	5.67	50	150
2	300	95	5.30	50	160
3	300	95	4.96	50	170

Note that the use of rep() is required in constructing the arguments for the advanced usage examples where more than one argument takes on multiple values.

References

See the 2657 Productions News site for how this function progressively developed¹. The sample.size function is based on the following formulas²:

$$ss = \frac{-Z^2 \times p \times (1-p)}{c^2}$$

$$pss = \frac{ss}{1 + \frac{ss - 1}{pop}}$$

 $[\]frac{1}{2} \text{http://news.mrdwab.com/2010/09/10/a-sample-size-calculator-function-for-r/}$

 $^{^2 \}rm See:$ Creative Research Systems. (n.d.). Sample size formulas for our sample size calculator. Retrieved from: http://www.surveysystem.com/sample-size-formula.htm. Archived on 07 August 2012 at http://www.webcitation.org/69kNjMuKe.

Part II The Functions

Where to Get the Functions

The most current source code for the functions described in this document follow.

To load the functions, you can directly source them from the 2657 R Functions page at github: https://github.com/mrdwab/2657-R-Functions

You should be able to load the functions using the following (replace ----- with the function name¹):

 $^{^1\}mathrm{The}$ "snippets" in Part III of this document can all be loaded from the script $\mathtt{snippets.R.}$

concat.split

```
concat.split = function(data, split.col, to.list=FALSE, mode=NULL,
                       sep=",", drop.col=FALSE) {
 # Takes a column with multiple values, splits the values into
    separate columns, and returns a new data.frame.
  # ,data, is the source data.frame; ,split.col, is the variable that
     needs to be split; ,to.list, is whether the split output should
    be added as a single variable list (defaults to "FALSE");
    mode, can be either ,binary, or ,value, (where ,binary, is
  # default and it recodes values to 1 or NA); , sep, is the
  # character separating each value (defaults to ,,,);
  # and ,drop.col, is logical (whether to remove the original
    variable from the output or not.
  # === EXAMPLES ===
          dat = data.frame(V1 = c("1, 2, 4", "3, 4, 5",
  #
                                  "1, 2, 5", "4", "1, 2, 3, 5"),
  #
                           V2 = c("1;2;3;4", "1", "2;5",
  #
                                  "3;2", "2;3;4"))
         dat2 = data.frame(V1 = c("Fred, John, Sue", "Jerry, Jill",
                                   "Sally, Ryan", "Susan, Amos, Ben"))
         concat.split(dat, 1)
         concat.split(dat, 2, sep=";")
         concat.split(dat, "V2", sep=";", mode="value")
         concat.split(dat, "V1", mode="binary")
         concat.split(dat2, 1)
          concat.split(dat2, "V1", drop.col=TRUE)
  # See: http://stackoverflow.com/q/10100887/1270695
  # Check to see if split.col is specified by name or position
  if (is.numeric(split.col)) split.col = split.col
 else split.col = which(colnames(data) %in% split.col)
  # Split the data
 a = as.character(data[ , split.col])
 b = strsplit(a, sep)
 #### LIST ####
 if (isTRUE(to.list)) {
```

}

```
varname = paste(names(data[split.col]), "_list", sep="")
  if (suppressWarnings(is.na(try(max(as.numeric(unlist(b))))))) {
    data[varname] = list(lapply(lapply(b, as.character),
                                function(x) gsub("^{\s+|\s+$"},
                                                 "", x)))
  } else if (!is.na(try(max(as.numeric(unlist(b)))))) {
    data[varname] = list(lapply(b, as.numeric))
  if (isTRUE(drop.col)) data[-split.col]
  else data
### SEPARATE COLUMNS ###
} else if (!isTRUE(to.list)) {
  if (suppressWarnings(is.na(try(max(as.numeric(unlist(b))))))) {
    what = "string"
   ncol = max(unlist(lapply(b, function(i) length(i))))
  } else if (!is.na(try(max(as.numeric(unlist(b)))))) {
    what = "numeric"
    ncol = max(as.numeric(unlist(b)))
  if (identical(what, "string")) {
    temp = as.data.frame(t(sapply(b, |[,, 1:ncol)))
   names(temp) = paste(names(data[split.col]), "_", 1:ncol, sep="")
   temp = apply(temp, 2, function(x) gsub("^{\s+|\s+$", "", x})
   temp1 = cbind(data, temp)
  } else if (identical(what, "numeric")) {
    temp = lapply(b, as.numeric)
    m = matrix(nrow = nrow(data), ncol = ncol)
    for (i in 1:nrow(data)) {
      m[i, temp[[i]]] = temp[[i]]
    m = setNames(data.frame(m),
                 paste(names(data[split.col]), "_", 1:ncol, sep=""))
    if (is.null(mode) || identical(mode, "binary")) {
      temp1 = cbind(data, replace(m, m != "NA", 1))
    } else if (identical(mode, "value")) {
      temp1 = cbind(data, m)
    }
  if (isTRUE(drop.col)) temp1[-split.col]
  else temp1
}
```

df.sorter

```
df.sorter = function(data, var.order=names(data), col.sort=NULL, at.start=TRUE ) {
  # Sorts a data.frame by columns or rows or both.
  # Can also subset the data columns by using ,var.order,.
  # Can refer to variables either by names or number.
  # If referring to variable by number, and sorting both the order
     of variables and the sorting within variables, refer to the
      variable numbers of the final data.frame.
  # === EXAMPLES ===
      library (foreign)
      temp = "http://www.ats.ucla.edu/stat/stata/modules/kidshtwt.dta"
      kidshtwt = read.dta(temp); rm(temp)
       df.sorter(kidshtwt, var.order = c("fam", "bir", "wt", "ht"))
       df.sorter(kidshtwt, var.order = c("fam", "bir", "wt", "ht"),
                 col.sort = c("birth", "famid")) # USE FULL NAMES HERE
       df.sorter(kidshtwt, var.order = c(1:4), \# DROP THE WT COLUMNS)
                 col.sort = 3)
                                                 # SORT BY HT1
  if (is.numeric(var.order))
    var.order = colnames(data)[var.order]
  else var.order = var.order
  if (isTRUE(at.start)) {
    x = unlist(lapply(var.order, function(x)
      sort(grep(paste("^", x, sep="", collapse=""),
                names(data), value = TRUE))))
  } else if (!isTRUE(at.start)) {
    x = unlist(lapply(var.order, function(x)
      sort(grep(x, names(data), value = TRUE))))
  y = data[, x]
  if (is.null(col.sort)) {
  } else if (is.numeric(col.sort)) {
    y[do.call(order, y[colnames(y)[col.sort]]), ]
  } else if (!is.numeric(col.sort)) {
    y[do.call(order, y[col.sort]), ]
}
```

multi.freq.table

```
multi.freq.table = function(data, sep="", boolean=TRUE,
                            factors=NULL,
                            NAtoO=TRUE, basic=FALSE,
                            dropzero=TRUE, clean=TRUE) {
  # Takes multiple-response data and tabulates it according
     to the possible combinations of each variable.
  # === EXAMPLES ===
  #
       set.seed(1)
       dat = data.frame(A = sample(c(0, 1), 20, replace=TRUE),
                         B = sample(c(0, 1), 20, replace=TRUE),
                         C = sample(c(0, 1), 20, replace=TRUE),
                         D = sample(c(0, 1), 20, replace=TRUE),
                         E = sample(c(0, 1), 20, replace=TRUE))
     multi.freq.table(dat)
     multi.freq.table(dat[1:3], sep="-", dropzero=TRUE)
  # See: http://stackoverflow.com/q/11348391/1270695
         http://stackoverflow.com/q/11622660/1270695
  if (!is.data.frame(data)) {
    stop("Input must be a data frame.")
  if (isTRUE(boolean)) {
    CASES = nrow(data)
    RESPS = sum(data, na.rm=TRUE)
    if(isTRUE(NAtoO)) {
      data[is.na(data)] = 0
      VALID = CASES
     VRESP = RESPS
    } else if(!isTRUE(NAto0)) {
      data = data[complete.cases(data), ]
      VALID = CASES - (CASES - nrow(data))
      VRESP = sum(data)
    if(isTRUE(basic)) {
      counts = data.frame(Freq = colSums(data),
                          Pct.of.Resp = (colSums(data)/sum(data))*100,
```

```
Pct.of.Cases = (colSums(data)/nrow(data))*100)
  } else if (!isTRUE(basic)) {
    counts = data.frame(table(data))
    Z = counts[, c(intersect(names(data), names(counts)))]
    Z = rowSums(sapply(Z, as.numeric)-1)
    if(Z[1] == 0) \{ Z[1] = 1 \}
    N = ncol(counts)
    counts$Combn = apply(counts[-N] == 1, 1,
                         function(x) paste(names(counts[-N])[x],
                                           collapse=sep))
    counts$Weighted.Freq = Z*counts$Freq
    counts$Pct.of.Resp = (counts$Weighted.Freq/sum(data))*100
    counts$Pct.of.Cases = (counts$Freq/nrow(data))*100
    if (isTRUE(dropzero)) {
      counts = counts[counts$Freq != 0, ]
    } else if (!isTRUE(dropzero)) {
      counts = counts
    }
    if (isTRUE(clean)) {
      counts = data.frame(Combn = counts$Combn, Freq = counts$Freq,
                          Weighted.Freq = counts$Weighted.Freq,
                          Pct.of.Resp = counts$Pct.of.Resp,
                          Pct.of.Cases = counts$Pct.of.Cases)
   }
  message("Total cases:
                            ", CASES, "\n",
          "Valid cases:
                           ", VALID, "\n",
          "Total responses: ", RESPS, "\n",
          "Valid responses: ", VRESP, "\n")
  counts
} else if (!isTRUE(boolean)) {
  CASES = nrow(data)
  RESPS = length(data[!is.na(data)])
  if (!isTRUE(any(sapply(data, is.factor)))) {
    if (is.null(factors)) {
      stop("Input variables must be factors.
      Please provide factors using the ,factors, argument or
           convert your data to factor before using function.")
    } else {
      data[sapply(data, is.character)] =
        lapply(data[sapply(data, is.character)],
               function(x) factor(x, levels=factors))
    }
  }
  if (isTRUE(basic)) {
   ROWS = levels(unlist(data))
    OUT = table(unlist(data))
   PCT = (OUT/sum(OUT)) * 100
    OUT = data.frame(ROWS, OUT, PCT, row.names=NULL)
    OUT = data.frame(Item = OUT[, 1], Freq = OUT[, 3],
                     Pct.of.Resp = OUT[, 5],
                     Pct.of.Cases = (OUT[, 3]/CASES)*100)
    message("Total cases: ", CASES, "\n",
            "Total responses: ", RESPS, "\n")
    OUT
  } else if (!isTRUE(basic)) {
    Combos = apply(data, 1, function(x) paste0(sort(x), collapse = sep))
    Weight = as.numeric(rowSums(!is.na(data)))
```

row.extractor

```
row.extractor = function(data, extract.by, what="all") {
  # Extracts rows with min, median, and max values, or by quantiles.
  # Values for "what" can be "min", "median", "max", "all", or a
    vector specifying the desired quantiles.
  # Values for "extract.by" can be the variable name or number.
  # === EXAMPLES ===
     set.seed(1)
  # dat = data.frame(V1 = 1:10, V2 = rnorm(10), V3 = rnorm(10),
                        V4 = sample(1:20, 10, replace=T))
    dat2 = dat[-10,]
    row.extractor(dat, 4, "all")
      row.extractor(dat1, 4, "min")
     row.extractor(dat, "V4", "median")
     row.extractor(dat, 4, c(0, .5, 1))
     row.extractor(dat, "V4", c(0, .25, .5, .75, 1))
  # "which.quantile" function by cheleites:
  # http://stackoverflow.com/users/755257/cbeleites
  # See: http://stackoverflow.com/q/10256503/1270695
  if (is.numeric(extract.by)) {
    extract.by = extract.by
  } else if (is.numeric(extract.by) != 0) {
    extract.by = which(colnames(data) %in% "extract.by")
  }
  if (is.character(what)) {
    which.median = function(data, extract.by) {
      a = data[, extract.by]
      if (length(a) %% 2 != 0) {
       which(a == median(a))
      } else if (length(a) %% 2 == 0) {
        b = sort(a)[c(length(a)/2, length(a)/2+1)]
        c(max(which(a == b[1])), min(which(a == b[2])))
    }
    X1 = data[which(data[extract.by] == min(data[extract.by])), ] # min
    X2 = data[which(data[extract.by] == max(data[extract.by])), ] # max
                                                              # median
    X3 = data[which.median(data, extract.by), ]
```

```
if (identical(what, "min")) {
     X1
    } else if (identical(what, "max")) {
    } else if (identical(what, "median")) {
    } else if (identical(what, "all")) {
      rbind(X1, X3, X2)
  } else if (is.numeric(what)) {
    which.quantile <- function (data, extract.by, what, na.rm = FALSE) {</pre>
      x = data[ , extract.by]
      if (! na.rm & any (is.na (x)))
        return (rep (NA_integer_, length (what)))
      o <- order (x)
      n <- sum (! is.na (x))
      o <- o [seq_len (n)]
      nppm \leftarrow n * what - 0.5
      j <- floor(nppm)</pre>
      h \leftarrow ifelse((nppm == j) & ((j\%2L) == 0L), 0, 1)
      j \leftarrow j + h
      j [j == 0] <- 1
     o[j]
    data[which.quantile(data, extract.by, what), ]
                                                               # quantile
}
```

sample.size

```
sample.size = function(population, samp.size=NULL, c.lev=95,
                       c.int=NULL, what = "sample",
                       distribution=50) {
  # Returns a data.frame of sample sizes or confidence
  # intervals for different conditions provided by
     the following arguments.
  # population: Population size
  # samp.size: Sample size
 # c.lev: Confidence level
  # c.int: Confidence interval (+/-)
  # what: Whether sample size or confidence interval
          is being calculated.
  # distribution: Response distribution
  # === EXAMPLES ===
    sample.size(300)
    sample.size(300, 150, what="confidence")
     sample.size(c(300, 400, 500), c.lev=97)
 z = qnorm(.5+c.lev/200)
 if (identical(what, "sample")) {
   if (is.null(c.int)) {
      c.int = 5
      message("NOTE! Confidence interval set to 5.
     To override, set c.int to desired value.\n")
    } else if (!is.null(c.int) == 1) {
      c.int = c.int
   if (!is.null(samp.size)) {
     message("NOTE! ,samp.size, value provided but ignored.
      See output for actual sample size(s).\n")
   ss = (z^2 * (distribution/100) *
      (1-(distribution/100)))/((c.int/100)^2)
    samp.size = ss/(1 + ((ss-1)/population))
```

```
} else if (identical(what, "confidence")) {
    if (is.null(samp.size)) {
     stop("Missing ,samp.size, with no default value.")
    if (!is.null(c.int)) {
     message("NOTE! ,c.int, value provided but ignored.
     See output for actual confidence interval value(s).\n")
    ss = ((population*samp.size-samp.size)/(population-samp.size))
    c.int = round(sqrt((z^2 * (distribution/100) *
      (1-(distribution/100)))/ss)*100, digits = 2)
  } else if (what %in% c("sample", "confidence") == 0) {
    stop(",what, must be either ,sample, or ,confidence,")
  RES = data.frame(population = population,
                   conf.level = c.lev,
                   conf.int = c.int,
                   distribution = distribution,
                   sample.size = round(samp.size, digits = 0))
  RES
}
```

Part III Snippets and Tips

Snippets

Load All Scripts and Data Files From Multiple Directories

```
load.scripts.and.data <- function(path,</pre>
                                 pattern=list(scripts = "*.R$",
                                               data = "*.rda$|*.Rdata$"),
                                 ignore.case=TRUE) {
  # Reads all the data files and scripts from specified directories.
       In general, should only need to specify the directories.
        Specify directories without trailing slashes.
  # === EXAMPLE ===
  #
       load.scripts.and.data(c("~/Dropbox/Public",
                               "~/Dropbox/Public/R Functions"))
 file.sources = list.files(path, pattern=pattern$scripts,
                            full.names=TRUE, ignore.case=ignore.case)
 data.sources = list.files(path, pattern=pattern$data,
                            full.names=TRUE, ignore.case=ignore.case)
 sapply(data.sources, load, .GlobalEnv)
  sapply(file.sources, source, .GlobalEnv)
```

Convert a List of Data Frames Into Individual Data Frames

```
unlist.dfs <- function(data) {
    # Specify the quoted name of the source list.
    q = get(data)
    prefix = pasteO(data, "_", 1:length(q))
    for (i in 1:length(q)) assign(prefix[i], q[[i]], envir=.GlobalEnv)
}</pre>
```

Example

Note that the list name must be quoted.

```
# Sample data
temp = list(A = data.frame(A = 1:2, B = 3:4),
```

```
B = data.frame(C = 5:6, D = 7:8))
temp
$A
 A B
1 1 3
2 2 4
$B
 C D
1 5 7
2 6 8
\# Remove any files with similar names to output
rm(list=ls(pattern="temp_"))
# The following should not work
temp_1
Error: object 'temp_1' not found
# Split it up!
unlist.dfs("temp")
# List files with the desired pattern
ls(pattern="temp_")
[1] "temp_1" "temp_2"
# View the new files
temp_1
 A B
1 1 3
2 2 4
temp_2
 C D
1 5 7
2 6 8
```

Convert a Data Frame Into a List With Each Column Becoming a List Item

```
dfcols.list <- function(data, vectorize=FALSE) {
    # Specify the unquoted name of the data.frame to convert
    if (isTRUE(vectorize)) {
        dat.list = sapply(1:ncol(data), function(x) data[x])
    } else if (!isTRUE(vectorize)) {
        dat.list = lapply(names(data), function(x) data[x])
    }
    dat.list
}</pre>
```

Examples

```
# Sample data
dat = data.frame(A = c(1:2), B = c(3:4), C = c(5:6))
  A B C
1 1 3 5
2 2 4 6
# Split into a list, retaining data.frame structure
dfcols.list(dat)
[[1]]
 Α
1 1
2 2
[[2]]
 В
1 3
2 4
[[3]]
 C
1 5
2 6
# Split into a list, converting to vector
dfcols.list(dat, vectorize=TRUE)
$A
[1] 1 2
$B
[1] 3 4
$C
[1] 5 6
```

Rename an Object in the Workplace

```
mv <- function (a, b) {
    # Source: https://stat.ethz.ch/pipermail/r-help/2008-March/156035.html
    anm = deparse(substitute(a))
    bnm = deparse(substitute(b))
    if (!exists(anm, where=1, inherits=FALSE))
        stop(paste(anm, "does not exist.\n"))
    if (exists(bnm, where=1, inherits=FALSE)) {
        ans = readline(paste("Overwrite ", bnm, "? (y/n) ", sep = ""))
        if (ans != "y")
            return(invisible())
    }
    assign(bnm, a, pos = 1)
    rm(list = anm, pos = 1)
    invisible()
}</pre>
```

Basic Usage

If there is already an object with the same name in the workplace, the function will ask you if you want to replace the object or not. Otherwise, the basic usage is:

```
# Rename "object_1" to "object_2"
mv(object_1, object_2)
```

Scrape Data From a Poorly Formatted HTML Page

Reformats a web page using HTML Tidy and uses the XML package to parse the resulting file. Can optionally save the reformatted page.

```
tidyHTML <- function(URL, saveTidy = TRUE) {
  require(XML)
  URL1 = gsub("/", "%2F", URL)
  URL1 = gsub(":", "%3A", URL1)
  URL1 = paste("http://services.w3.org/tidy/tidy?docAddr=", URL1, "&indent=on", sep = "")
  Parsed = htmlParse(URL1)
  if (isTRUE(saveTidy)) saveXML(Parsed, file = basename(URL))
  Parsed
}</pre>
```

Example

```
# Set "saveTidy" to "TRUE" to save the resulting tidied file
URL <- "http://www.bcn.gob.ni/estadisticas/trimestrales_y_mensuales/siec/datos/4.IMAE.htm"
temp <- tidyHTML(URL, saveTidy = FALSE)</pre>
```

"Rounding in Commerce"

R rounds to even—something that some people might not be accustomed to or comfortable with. For the more commonly known rounding rule, use this round2 function.

```
round2 <- function(x, n = 0) {
  posneg = sign(x)
  z = abs(x)*10^n
  z = z + 0.5
  z = trunc(z)
  z = z/10^n
  z*posneg
}</pre>
```

Example

```
x = c(1.85, 1.54, 1.65, 1.85, 1.84)
round(x, 1)
[1] 1.8 1.5 1.6 1.8 1.8
round2(x, 1)
```

```
[1] 1.9 1.5 1.7 1.9 1.8

round(seq(0.5, 9.5, by=1))

[1] 0 2 2 4 4 6 6 8 8 10

round2(seq(0.5, 9.5, by=1))

[1] 1 2 3 4 5 6 7 8 9 10
```

References

Original function: http://www.webcitation.org/68djeLBtJ – see the comments section. See also: http://stackoverflow.com/questions/12688717/round-up-from-5-in-r/.

Tips

Many of the following tips are useful for reducing repetitious tasks. They might seem silly or unnecessary with the small examples provided, but they can be huge time-savers when dealing with larger objects or larger sets of data.

Batch Convert Factor Variables to Character Variables

In the example data below, author and title are automatically converted to factor (unless you add the argument stringsAsFactor = FALSE when you are creating the data). What if you forgot and actually needed the variables to be in mode as.character instead?

Use sapply to identify which variables are currently factors and convert them to as.character.

```
dat = data.frame(title = c("title1", "title2", "title3"),
                author = c("author1", "author2", "author3"),
                customerID = c(1, 2, 1)
str(dat)
'data.frame':
               3 obs. of 3 variables:
           : Factor w/ 3 levels "title1", "title2", ...: 1 2 3
$ author : Factor w/ 3 levels "author1", "author2",..: 1 2 3
$ customerID: num 1 2 1
# Left of the equal sign identifies and extracts the factor variables;
# right converts them from factor to character
dat[sapply(dat, is.factor)] = lapply(dat[sapply(dat, is.factor)],
                                    as.character)
str(dat)
               3 obs. of 3 variables:
'data.frame':
$ title : chr "title1" "title2" "title3"
            : chr "author1" "author2" "author3"
$ customerID: num 1 2 1
```

Using Reduce to Merge Multiple Data Frames at Once

The merge function in R only merges two objects at a time. This is usually fine, but what if you had several data.frames that needed to be merged?

Consider the following data, where we want to take monthly tables and merge them into an annual table:

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```
set.seed(1)
JAN = data.frame(ID = sample(5, 3), JAN = sample(LETTERS, 3))
FEB = data.frame(ID = sample(5, 3), FEB = sample(LETTERS, 3))
MAR = data.frame(ID = sample(5, 3), MAR = sample(LETTERS, 3))
APR = data.frame(ID = sample(5, 3), APR = sample(LETTERS, 3))
```

If we wanted to merge these into a single data.frame using merge, we might end up creating several temporary objects and merging those, like this:

```
temp_1 = merge(JAN, FEB, all=TRUE)
temp_2 = merge(temp_1, MAR, all=TRUE)
temp_3 = merge(temp_2, APR, all=TRUE)
```

Or, we might nest a whole bunch of merge commands together, something like this:

However, that first option requires a lot of unnecessary typing and produces unnecessary objects that we then need to remember to remove, and the second option is not very reader-friendly—try doing a merge like that with, say, 12 data.frames if we had an entire year of data!

Use Reduce instead, simply specifying all the objects to be merged in a list:

```
Reduce(function(x, y) merge(x, y, all=TRUE),
      list(JAN, FEB, MAR, APR))
     JAN FEB MAR APR
1 2
       X
            Ε
                 R
                      F
            F
2 3 <NA>
                 Х
                      D
3 4
       V <NA>
                 Μ
            B <NA> <NA>
```

How Much Memory Are the Objects in Your Workspace Using?

Sometimes you need to just check and see how much memory the objects in your workspace occupy.

```
sort(sapply(ls(), function(x) {object.size(get(x))}))
```

Convert a Table to a Data Frame

Creating tables are easy and fast, but sometimes, it is more convenient to have the output as a data.frame. Get the data.frame by nesting the command in as.data.frame.matrix.

```
# A basic table
x <- with(airquality, table(cut(Temp, quantile(Temp)), Month))
str(x)

'table' int [1:4, 1:5] 24 5 1 0 3 15 7 5 0 2 ...
- attr(*, "dimnames")=List of 2
    ..$ : chr [1:4] "(56,72]" "(72,79]" "(79,85]" "(85,97]"
    ..$ Month: chr [1:5] "5" "6" "7" "8" ...</pre>
```

X

```
Month
          5 6 7 8 9
  (56,72] 24 3 0 1 10
  (72,79] 5 15 2 9 10
  (79,85] 1 7 19 7 5
  (85,97] 0 5 10 14 5
# The same table as a data.frame
y <- as.data.frame.matrix(x)</pre>
str(y)
'data.frame': 4 obs. of 5 variables:
$ 5: int 24 5 1 0
$ 6: int 3 15 7 5
$ 7: int 0 2 19 10
$ 8: int 1 9 7 14
$ 9: int 10 10 5 5
У
        5 6 7 8 9
(56,72] 24 3 0 1 10
(72,79] 5 15 2 9 10
(79,85] 1 7 19 7 5
(85,97] 0 5 10 14 5
```