

2657 Functions

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Contents

| | |
|----------------------------|-----------|
| concat.split | 3 |
| What it Does | 3 |
| Arguments | 3 |
| Examples | 3 |
| To Do | 5 |
| References | 5 |
| df.sorter | 6 |
| What it Does | 6 |
| Arguments | 6 |
| Examples | 6 |
| To Do | 8 |
| multi.freq.table | 9 |
| What it Does | 9 |
| Arguments | 9 |
| Examples | 9 |
| References | 10 |
| row.extractor | 11 |
| What it Does | 11 |
| Arguments | 11 |
| Examples | 11 |
| References | 12 |
| The Functions | 13 |
| concat.split | 14 |
| df.sorter | 16 |
| multi.freq.table | 17 |
| row.extractor | 18 |

concat.split

What it Does

The `concat.split` function takes a column with multiple values, splits the values 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.
- `mode`: can be either `binary` or `value` (where `binary` is default and it recodes values to 1 or NA).
- `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](https://raw.githubusercontent.com/mrdwab/2657-R-Functions/master/data/concatenated-cells.csv). The URL is an HTTPS, so we need to use `getURL` from `RCurl`.

```
require(RCurl)

## Loading required package: RCurl

## Loading required package: bitops

baseURL = c("https://raw.githubusercontent.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)

##      Name      Likes      Siblings      Hates
## 1  Boyd 1,2,4,5,6 Reynolds , Albert , Ortega 2;4;
## 2  Rufus 1,2,4,5,6 Cohen , Bert , Montgomery 1;2;3;4;
## 3   Dana 1,2,4,5,6                Pierce 2;
## 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.githubusercontent.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
## 1   Boyd 1,2,4,5,6 Reynolds , Albert , Ortega      2;4;      1      1
## 2   Rufus 1,2,4,5,6 Cohen , Bert , Montgomery 1;2;3;4;      1      1
## 3    Dana 1,2,4,5,6      Pierce      2;      1      1
## 4   Carole 1,2,4,5,6 Colon , Michelle , Ballard      1;4;      1      1
## 5   Ramona 1,2,5,6      Snyder , Joann ,      1;2;3;      1      1
## 6   Kelley 1,2,5,6      James , Roxanne ,      1;4;      1      1
## Likes_3 Likes_4 Likes_5 Likes_6
## 1      NA      1      1      1
## 2      NA      1      1      1
## 3      NA      1      1      1
## 4      NA      1      1      1
## 5      NA      NA      1      1
## 6      NA      NA      1      1
```

```
# ... or by name, and drop the offensive first column
head(concat.split(concat.test, "Likes", drop.col = TRUE))
```

```
##      Name      Siblings      Hates Likes_1 Likes_2 Likes_3
## 1   Boyd Reynolds , Albert , Ortega      2;4;      1      1      NA
## 2   Rufus Cohen , Bert , Montgomery 1;2;3;4;      1      1      NA
## 3    Dana      Pierce      2;      1      1      NA
## 4   Carole Colon , Michelle , Ballard      1;4;      1      1      NA
## 5   Ramona      Snyder , Joann ,      1;2;3;      1      1      NA
## 6   Kelley      James , Roxanne ,      1;4;      1      1      NA
## Likes_4 Likes_5 Likes_6
## 1      1      1      1
## 2      1      1      1
## 3      1      1      1
## 4      1      1      1
## 5      NA      1      1
## 6      NA      1      1
```

```
# The 'Hates' column uses a different separator:
head(concat.split(concat.test, "Hates", sep = ";", drop.col = TRUE))
```

```
##      Name      Likes      Siblings Hates_1 Hates_2 Hates_3
## 1   Boyd 1,2,4,5,6 Reynolds , Albert , Ortega      NA      1      NA
## 2   Rufus 1,2,4,5,6 Cohen , Bert , Montgomery      1      1      1
## 3    Dana 1,2,4,5,6      Pierce      NA      1      NA
## 4   Carole 1,2,4,5,6 Colon , Michelle , Ballard      1      NA      NA
## 5   Ramona 1,2,5,6      Snyder , Joann ,      1      1      1
## 6   Kelley 1,2,5,6      James , Roxanne ,      1      NA      NA
## Hates_4
## 1      1
## 2      1
## 3      NA
## 4      1
## 5      NA
## 6      1
```

```
# Retain the original values
head(concat.split(concat.test, 2, mode = "value", drop.col = TRUE))
```

| ## | Name | Siblings | Hates | Likes_1 | Likes_2 | Likes_3 |
|------|-----------------------------------|----------|-------|---------|---------|---------|
| ## 1 | Boyd Reynolds , Albert , Ortega | 2;4; | | 1 | 2 | NA |
| ## 2 | Rufus Cohen , Bert , Montgomery | 1;2;3;4; | | 1 | 2 | NA |
| ## 3 | Dana | Pierce | 2; | 1 | 2 | NA |
| ## 4 | Carole Colon , Michelle , Ballard | 1;4; | | 1 | 2 | NA |
| ## 5 | Ramona Snyder , Joann , | 1;2;3; | | 1 | 2 | NA |
| ## 6 | Kelley James , Roxanne , | 1;4; | | 1 | 2 | NA |

| ## | Likes_4 | Likes_5 | Likes_6 |
|------|---------|---------|---------|
| ## 1 | 4 | 5 | 6 |
| ## 2 | 4 | 5 | 6 |
| ## 3 | 4 | 5 | 6 |
| ## 4 | 4 | 5 | 6 |
| ## 5 | NA | 5 | 6 |
| ## 6 | NA | 5 | 6 |

```
# Let's try splitting some strings... Same syntax
head(concat.split(concat.test, 3, drop.col = TRUE))
```

| ## | Name | Likes | Hates | Siblings_1 | Siblings_2 | Siblings_3 |
|------|--------|-----------|----------|------------|------------|------------|
| ## 1 | Boyd | 1,2,4,5,6 | 2;4; | Reynolds | Albert | Ortega |
| ## 2 | Rufus | 1,2,4,5,6 | 1;2;3;4; | Cohen | Bert | Montgomery |
| ## 3 | Dana | 1,2,4,5,6 | 2; | Pierce | <NA> | <NA> |
| ## 4 | Carole | 1,2,4,5,6 | 1;4; | Colon | Michelle | Ballard |
| ## 5 | Ramona | 1,2,5,6 | 1;2;3; | Snyder | Joann | <NA> |
| ## 6 | Kelley | 1,2,5,6 | 1;4; | James | Roxanne | <NA> |

To Do

- Add the option to put the output as lists instead of adding multiple columns to the `data.frame`.
- Modify the function so that you can split multiple columns in one go?

References

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

df.sorter

What it Does

The `df.sorter` function allows you to sort a `data.frame` by columns or rows or both. You can also quickly subset data solums 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.githubusercontent.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 | 1 | 1 | -0.6265 | 145 | DAB | -0.7075 | 299 | CEB |
| ## 2 | 1 | 2 | 0.1836 | 180 | DCB | 0.3646 | 224 | ECD |
| ## 3 | 1 | 3 | -0.8356 | 148 | EBA | 0.7685 | 222 | DAE |
| ## 4 | 2 | 1 | 1.5953 | 56 | AED | -0.1123 | 175 | DBA |
| ## 5 | 2 | 2 | 0.3295 | 245 | CEB | 0.8811 | 260 | DAC |
| ## 6 | 2 | 3 | -0.8205 | 198 | EBD | 0.3981 | 216 | DCA |

```
## 7 3 1 0.4874 234 BCA -0.6120 300 CEA
## 8 3 2 0.7383 32 CDA 0.3411 179 CAD
## 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
## 12 4 3 0.3898 188 DEB -0.3672 160 DBE
## 13 5 1 -0.6212 226 DBA -1.0441 154 EDB
## 14 5 2 -2.2147 159 DAC 0.5697 238 BDE
## 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 1 1 DAB CEB -0.6265 -0.7075 145 299
## 2 1 2 DCB ECD 0.1836 0.3646 180 224
## 3 1 3 EBA DAE -0.8356 0.7685 148 222
## 4 2 1 AED DBA 1.5953 -0.1123 56 175
## 5 2 2 CEB DAC 0.3295 0.8811 245 260
## 6 2 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 1 1 DAB CEB -0.6265 -0.7075 145 299
## 2 1 2 DCB ECD 0.1836 0.3646 180 224
## 3 1 3 EBA DAE -0.8356 0.7685 148 222
## 4 2 1 AED DBA 1.5953 -0.1123 56 175
## 5 2 2 CEB DAC 0.3295 0.8811 245 260
## 6 2 3 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 1 1 DAB CEB -0.6265 -0.7075 145 299
## 4 2 1 AED DBA 1.5953 -0.1123 56 175
## 7 3 1 BCA CEA 0.4874 -0.6120 234 300
## 10 4 1 BED CDE -0.3054 1.4330 120 234
## 13 5 1 DBA EDB -0.6212 -1.0441 226 154
## 2 1 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
## 10 4 1 120 234 BED CDE
## 1 1 1 145 299 DAB CEB
## 13 5 1 226 154 DBA EDB
## 7 3 1 234 300 BCA CEA
## 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
## 4      2      1      56      175  AED  DBA
## 10     4      1     120      234  BED  CDE
## 1      1      1     145      299  DAB  CEB
## 13     5      1     226      154  DBA  EDB
## 7      3      1     234      300  BCA  CEA
## 8      3      2       32      179  CDA  CAD
```

```
# 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      1      1 -0.6265     145  DAB  -0.7075     299  CEB
## 4      2      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     120  BED   1.4330     234  CDE
## 13     5      1 -0.6212     226  DBA  -1.0441     154  EDB
## 2      1      2  0.1836     180  DCB   0.3646     224  ECD
```

```
head(df.sorter(dat, col.sort = c("code1"))) # Sorting by character values
```

```
##      id times measure1 score1 code1 measure2 score2 code2
## 4      2      1  1.5953       56  AED  -0.1123     175  DBA
## 15     5      3  1.1249     152  AED  -0.1351     277  DCE
## 7      3      1  0.4874     234  BCA  -0.6120     300  CEA
## 10     4      1 -0.3054     120  BED   1.4330     234  CDE
## 8      3      2  0.7383       32  CDA   0.3411     179  CAD
## 5      2      2  0.3295     245  CEB   0.8811     260  DAC
```

```
# 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

What it Does

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.

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).
- `dropzero`: Should combinations with a frequency of zero be dropped from the final table? Defaults to FALSE.
- `clean`: Should the original tabulated data be retained or dropped from the final table? Defaults to TRUE.

Examples

```
# Load the function! require(RCurl) baseURL =
# c('https://raw.githubusercontent.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), 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))
# View your data
dat

##      A B C D E
## 1  0 1 1 1 0
## 2  0 0 1 0 1
## 3  1 1 1 0 0
## 4  1 0 1 0 0
## 5  0 0 1 1 1
## 6  1 0 1 0 0
## 7  1 0 0 0 1
## 8  1 0 0 1 0
## 9  1 1 1 0 0
## 10 0 0 1 1 0
## 11 0 0 0 0 0
## 12 0 1 1 1 0
## 13 1 0 0 0 1
## 14 0 0 0 0 1
## 15 1 1 0 0 1
## 16 0 1 0 1 1
## 17 1 1 0 1 0
## 18 1 0 1 0 0
## 19 0 1 1 1 1
## 20 1 0 0 1 1

# Apply the function with all defaults accepted
multi.freq.table(dat)
```

```
##      Combn Freq
## 1      1
## 2      A    0
## 3      B    0
## 4     AB    0
## 5      C    0
## 6     AC    3
## 7     BC    0
## 8    ABC    2
## 9      D    0
## 10     AD    1
## 11     BD    0
## 12    ABD    1
## 13     CD    1
## 14    ACD    0
## 15    BCD    2
## 16   ABCD    0
## 17     E    1
## 18    AE    2
## 19     BE    0
## 20    ABE    1
## 21     CE    1
## 22    ACE    0
## 23    BCE    0
## 24   ABCE    0
## 25     DE    0
## 26    ADE    1
## 27    BDE    1
## 28   ABDE    0
## 29     CDE    1
## 30    ACDE    0
## 31    BCDE    1
## 32   ABCDE    0
```

```
# Tabulate only on variables 'A', 'B', and 'D', with a different
# separator, dropping any zero frequency values, and keeping the original
# tabulations. Note that there are no solitary 'B' responses.
multi.freq.table(dat[c(1, 2, 4)], sep = "-", dropzero = TRUE, clean = FALSE)
```

```
##   A B D Freq Combn
## 1 0 0 0     3
## 2 1 0 0     5      A
## 4 1 1 0     3    A-B
## 5 0 0 1     2      D
## 6 1 0 1     2    A-D
## 7 0 1 1     4    B-D
## 8 1 1 1     1  A-B-D
```

References

apply shortcut for creating the Combn column in the output by [Justin](#)
 See: <http://stackoverflow.com/q/11348391/1270695>

row.extractor

What it Does

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.githubusercontent.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 summary of the data  
summary(dat)
```

```
##           V1           V2           V3           V4  
## Min.      : 1.0      Min.    :-2.215      Min.     :0.000      Min.      : 2.00  
## 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      Mean   : 0.100      Mean   :0.774      Mean   :14.80  
## 3rd Qu.:37.8      3rd Qu.: 0.728      3rd Qu.:1.175      3rd Qu.:20.75  
## Max.    :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")

##      V1      V2  V3 V4
## 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))

##      V1      V2  V3 V4
## 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
```

References

which.quantile function by [cbeleites](#)
 See: <http://stackoverflow.com/q/10256503/1270695>

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 name):

```
require(RCurl)
baseURL = c("https://raw.githubusercontent.com/mrdwab/2657-R-Functions/master/")
source(textConnection(getURL(paste0(baseURL, "scripts/-----R"))))
```

concat.split

```
concat.split = function(data, split.col, 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; '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

  if (is.numeric(split.col))
    split.col = split.col else split.col = which(colnames(data) %in% split.col)

  a = as.character(data[, split.col])
  b = strsplit(a, sep)

  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)))
  }

  m = matrix(nrow = nrow(data), ncol = ncol)
  v = vector("list", nrow(data))

  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")) {
    for (i in 1:nrow(data)) {
      v[[i]] = as.numeric(strsplit(a, sep)[[i]])
    }

    temp = v

    for (i in 1:nrow(data)) {
      m[i, temp[[i]]] = temp[[i]]
    }
  }
}
```

```

m = data.frame(m)
names(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

  a = names(data)
  b = length(var.order)
  subs = vector("list", b)

  if (isTRUE(at.start)) {
    for (i in 1:b) {
      subs[[i]] = sort(grep(paste("^", var.order[i], sep = "", collapse = ""),
        a, value = TRUE))
    }
  } else if (!isTRUE(at.start)) {
    for (i in 1:b) {
      subs[[i]] = sort(grep(var.order[i], a, value = TRUE))
    }
  }

  x = unlist(subs)
  y = data[, x]

  if (is.null(col.sort)) {
    y
  } else if (is.numeric(col.sort)) {
    col.sort = colnames(y)[col.sort]
    y[do.call(order, y[col.sort]), ]
  } else if (!is.numeric(col.sort)) {
    col.sort = col.sort
    y[do.call(order, y[col.sort]), ]
  }
}
```


multi.freq.table

```
multi.freq.table = function(data, sep = "", dropzero = FALSE, clean = TRUE) {  
  # Takes boolean 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  
  
  counts = data.frame(table(data))  
  N = ncol(counts)  
  counts$Combn = apply(counts[-N] == 1, 1, function(x) paste(names(counts[-N])[x],  
    collapse = sep))  
  if (isTRUE(dropzero)) {  
    counts = counts[counts$Freq != 0, ]  
  } else if (!isTRUE(dropzero)) {  
    counts = counts  
  }  
  if (isTRUE(clean)) {  
    counts = data.frame(Combn = counts$Combn, Freq = counts$Freq)  
  }  
  counts  
}
```

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 cbeleites:
  # 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
    X3 = data[which.median(data, extract.by), ] # median

    if (identical(what, "min")) {
      X1
    } else if (identical(what, "max")) {
      X2
    } else if (identical(what, "median")) {
      X3
    } else if (identical(what, "all")) {
      rbind(X1, X3, X2)
    }
  } else if (is.numeric(what)) {
    which.quantile <- function(data, extract.by, what, na.rm = FALSE) {

      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 <- n * what - 0.5
    j <- floor(nppm)
    h <- ifelse((nppm == j) & ((j%2L) == 0L), 0, 1)
    j <- j + h

    j[j == 0] <- 1
    o[j]
  }
  data[which.quantile(data, extract.by, what), ] # quantile
}

```