

2657 Functions

Ananda Mahto

July 6, 2012

Contents

concat.split	3
What it Does	3
Arguments	3
The Function	3
Examples	4
To Do	6
References	6
df.sorter	7
What it Does	7
Arguments	7
The Function	7
Examples	8
To Do	10
multi.freq.table	11
What it Does	11
Arguments	11
The Function	11
Examples	11
References	13
row.extractor	14
What it Does	14
Arguments	14
The Function	14
Examples	15
References	16

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).

The Function

```
concat.split = function(data, split.col, mode = NULL, sep = ",",
  drop.col = FALSE) {

  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 = "")
    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
}

```

Examples

First load some data from a CSV stored at [github](https://raw.githubusercontent.com/mrdwab/2657-R-Functions/master/). 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!

```

# 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

- Modify the function so that you can split multiple columns in one go?
- Strip whitespace from string output.

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`.

The Function

```
df.sorter = function(data, var.order = names(data), col.sort = NULL,
  at.start = TRUE) {
  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]), ]
}
}

```

Examples

```

# 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`.

The Function

```
multi.freq.table = function(data, sep = "", dropzero = FALSE, clean = TRUE) {  
  
  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  
}
```

Examples

```
# 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
## 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)`

The Function

```
row.extractor = function(data, extract.by, what = "all") {  
  
  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
}

```

Examples

```

# 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>