

# Package ‘SSBtools’

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**Type** Package

**Title** Statistics Norway's Miscellaneous Tools

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**Depends** Matrix

**Imports** stringr, methods

**Description** Functions used by other packages from Statistics Norway are gathered. General data manipulation functions, and functions for hierarchical computations are included. The hierarchy specification functions are useful within statistical disclosure control.

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**URL** <https://github.com/statisticsnorway/SSBtools>

**BugReports** <https://github.com/statisticsnorway/SSBtools/issues>

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AddLeadingZeros	<i>Add leading zeros to numbers while preserving other text</i>
-----------------	---

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## Description

This function is created to fix problems caused by a serious bug in Excel. Editing csv files in that program causes leading zeros to disappear.

## Usage

```
AddLeadingZeros(
  codes,
  places,
  warningText = NULL,
  viaFactor = TRUE,
  nWarning = 6,
```

```

    removeLeadingTrailingWhitespace = TRUE
)

```

**Arguments**

codes	Character vector
places	Number of places for positive numbers. Minus sign is extra
warningText	When non-NULL, warning will be produced
viaFactor	When TRUE, the algorithm uses factor coding internally.
nWarning	Number of elements to be written before ... in warnings.
removeLeadingTrailingWhitespace	Remove leading and trailing whitespace

**Value**

Character vector

**Author(s)**

Øyvind Langsrød

**Examples**

```

AddLeadingZeros(c("1", "ABC", "12345", " 23", "-8", "45 ", " -9", " Agent ", "007",
                 "7 James Bond "), 10)
AddLeadingZeros(c("1", "ABC", "12345", " 23", "-8", "45 ", " -9", " Agent ", "007",
                 "7 James Bond "), 4)
AddLeadingZeros(c("1", "ABC", "12345", " 23", "-8", "45 ", " -9", " Agent ", "007",
                 "7 James Bond "), 4, removeLeadingTrailingWhitespace = FALSE)
AddLeadingZeros(c("1", "ABC", "12345", " 23", "-8", "45 ", " -9", " Agent ", "007",
                 "7 James Bond "), 4, warningText = "string changes")
AddLeadingZeros(c("1", "ABC", "12345", " 23", "-8", "45 ", " -9", " Agent ", "007",
                 "7 James Bond "), 4, warningText = "", nWarning = 2)

```

**Description**

Automatic convert list of hierarchies coded in different ways to standardized to-from coding

## Usage

```
AutoHierarchies(
  hierarchies,
  data = NULL,
  total = "Total",
  hierarchyVarNames = c(mapsFrom = "mapsFrom", mapsTo = "mapsTo", sign = "sign", level
    = "level"),
  combineHierarchies = TRUE,
  unionComplement = FALSE
)
FindHierarchies(data, total = "Total")
```

## Arguments

<code>hierarchies</code>	List of hierarchies
<code>data</code>	Matrix or data frame with data containing codes of relevant variables
<code>total</code>	Within AutoHierarchies: Vector of total codes (possibly recycled) used when running <code>Hrc2DimList</code> .
<code>hierarchyVarNames</code>	Variable names in the hierarchy tables as in <code>HierarchyFix</code>
<code>combineHierarchies</code>	Whether to combine several hierarchies for same variable into a single hierarchy
<code>unionComplement</code>	Logical vector as in <code>Hierarchies2ModelMatrix</code> . The parameter is only in use when hierarchies are combined.

## Details

Input can be to-from coded hierarchies, hierarchies/dimList as in sdcTable, TauArgus coded hierarchies or formulas. Automatic coding from data is also supported. Output is on a from ready for input to `HierarchyCompute`. `FindHierarchies` wraps `FindDimLists` and `AutoHierarchies` into a single function. A single string as hierarchy input is assumed to be a total code. Then, the hierarchy is created as a simple hierarchy where all codes in data sum up to this total. For consistence with `HierarchyCompute`, the codes "rowFactor" and "colFactor" are unchanged. An empty string is recoded to "rowFactor".

## Value

List of hierarchies

## Author(s)

Øyvind Langsrød

## See Also

`DimList2Hierarchy`, `Hierarchy2Formula`.

## Examples

```
# First, create different types of input
z <- SSBtoolsData("sprt_emp_withEU")
yearFormula <- c("y_14 = 2014", "y_15_16 = y_all - y_14", "y_all = 2014 + 2015 + 2016")
yearHier <- Formula2Hierarchy(yearFormula)
geoDimList <- FindDimLists(z[, c("geo", "eu")], total = "Europe")[[1]]
geoDimList2 <- FindDimLists(z[, c("geo", "eu")])[[1]]
geoHrc <- DimList2Hrc(geoDimList)
ageHier <- SSBtoolsData("sprt_emp_ageHier")

h1 <- AutoHierarchies(list(age = ageHier, geo = geoDimList, year = yearFormula))
h2 <- AutoHierarchies(list(age = "Y15-64", geo = geoHrc, year = yearHier), data = z,
                      total = "Europe")
h3 <- AutoHierarchies(list(age = "Total", geo = geoDimList2, year = "Total"), data = z)
h4 <- FindHierarchies(z[, c(1, 2, 3, 5)])
h5 <- AutoHierarchies(list(age = "Total", geo = "", year = "colFactor"), data = z)
identical(h1, h2)
identical(h3, h4)

FindHierarchies(z[, c("geo", "eu", "age")])
```

## AutoSplit

*Creating variables by splitting the elements of a character vector without needing a split string*

## Description

Creating variables by splitting the elements of a character vector without needing a split string

## Usage

```
AutoSplit(
  s,
  split = NULL,
  border = "_",
  revBorder = FALSE,
  noSplit = FALSE,
  varNames = paste("var", 1:100, sep = ""),
  tryReverse = TRUE
)
```

## Arguments

s	The character vector
split	Split string. When NULL (default), automatic splitting without a split string.
border	A split character or an integer (move split) to be used when the exact split position is not unique.

<code>revBorder</code>	When border is integer the split position is moved from the other side.
<code>noSplit</code>	No splitting when TRUE.
<code>varNames</code>	Variable names of the created variables (too many is ok)
<code>tryReverse</code>	When TRUE, the automatic method tries to find more variables by splitting from reversed strings.

**Value**

A data frame with s as row names.

**Author(s)**

Øyvind Langsrød

**Examples**

```
s <- c("A12-3-A-x", "A12-3-B-x", "B12-3-A-x", "B12-3-B-x",
      "A12-3-A-y", "A12-3-B-y", "B12-3-A-y", "B12-3-B-y")
AutoSplit(s)
AutoSplit(s,border="-")
AutoSplit(s,split="-")
AutoSplit(s,border=1)
AutoSplit(s,border=2)
AutoSplit(s,border=2,revBorder=TRUE)
AutoSplit(s,noSplit=TRUE)
AutoSplit(s,varNames=c("A", "B", "C", "D"))
```

**CbindIdMatch**

*Combine several data frames by using id variables to match rows*

**Description**

Combine several data frames by using id variables to match rows

**Usage**

```
CbindIdMatch(
  ...,
  addName = names(x),
  sep = "_",
  idNames = sapply(x, function(x) names(x)[1]),
  idNames1 = idNames,
  addLast = FALSE
)
```

## Arguments

...	Several data frames as several input parameters or a list of data frames
addName	NULL or vector of strings used to name columns according to origin frame
sep	A character string to separate when addName apply
idNames	Names of a id variable within each data frame
idNames1	Names of variables in first data frame that correspond to the id variable within each data frame
addLast	When TRUE addName will be at end

## Details

The first data frame is the basis and the other frames will be matched by using id-variables. The default id-variables are the first variable in each frame. Corresponding variables with the same name in first frame is assumed. An id-variable is not needed if the number of rows is one or the same as the first frame. Then the element of idNames can be set to a string with zero length.

## Value

A single data frame

## Author(s)

Øyvind Langsrød

## See Also

[RbindAll](#) (same example data)

## Examples

```

zA <- data.frame(idA = 1:10, idB = rep(10 * (1:5), 2), idC = rep(c(100, 200), 5),
                   idC2 = c(100, rep(200, 9)), idC3 = rep(100, 10),
                   idD = 99, x = round(rnorm(10), 3), xA = round(runif(10), 2))
zB <- data.frame(idB = 10 * (1:5), x = round(rnorm(5), 3), xB = round(runif(5), 2))
zC <- data.frame(idC = c(100, 200), x = round(rnorm(2), 3), xC = round(runif(2), 2))
zD <- data.frame(idD = 99, x = round(rnorm(1), 3), xD = round(runif(1), 2))
CbindIdMatch(zA, zB, zC, zD)
CbindIdMatch(a = zA, b = zB, c = zC, d = zD, idNames = c("", "idB", "idC", ""))
CbindIdMatch(a = zA, b = zB, c = zC, d = zD, idNames1 = c("", "idB", "idC2", ""))
CbindIdMatch(a = zA, b = zB, c = zC, d = zD, idNames1 = c("", "idB", "idC3", ""))
CbindIdMatch(zA, zB, zC, zD, addName = c("", "bbb", "ccc", "ddd"), sep = ".", addLast = TRUE)
try(CbindIdMatch(X = zA, Y = zA[, 4:5], Z = zC, idNames = NULL)) # Error
CbindIdMatch(X = zA, Y = zA[, 4:5], Z = zD, idNames = NULL)      # Ok since equal NROW or NROW==1
CbindIdMatch(list(a = zA, b = zB, c = zC, d = zD))               # List is alternative input

```

**DimList2Hierarchy**      *DimList2Hierarchy*

### Description

From hierarchy/dimList as in sdcTable to to-from coded hierarchy

### Usage

```
DimList2Hierarchy(x)
```

### Arguments

x	An element of a dimList as in sdcTable
---	--

### Value

Data frame with to-from coded hierarchy

### Author(s)

Øyvind Langsrød

### Examples

```
# First generate a dimList element
x <- FindDimLists(SSBtoolsData("sprt_emp_withEU")[, c("geo", "eu")], , total = "Europe")[[1]]

DimList2Hierarchy(x)
```

**DimList2Hrc**      *DimList2Hrc/Hrc2DimList*

### Description

Conversion between hierarchies/dimList as in sdcTable and TauArgus coded hierarchies

### Usage

```
DimList2Hrc(dimList)

Hrc2DimList(hrc, total = "Total")
```

**Arguments**

dimList	List of data frames according to the specifications in sdcTable
hrc	List of character vectors
total	String used to name totals.

**Value**

See Arguments

**Author(s)**

Øyvind Langsrud

**Examples**

```
# First generate dimList
dimList <- FindDimLists(SSBtoolsData("sppt_emp_withEU")[, c("geo", "eu", "age")])
hrc <- DimList2Hrc(dimList)
dimList2 <- Hrc2DimList(hrc)
identical(dimList, dimList2)
```

**Description**

A matrix for mapping input codes (columns) to output codes (rows) are created.

**Usage**

```
DummyHierarchy(
  mapsFrom,
  mapsTo,
  sign,
  level,
  mapsInput = NULL,
  inputInOutput = FALSE,
  keepCodes = mapsFrom[integer(0)],
  unionComplement = FALSE,
  reOrder = FALSE
)
```

## Arguments

<code>mapsFrom</code>	Character vector from hierarchy table
<code>mapsTo</code>	Character vector from hierarchy table
<code>sign</code>	Numeric vector of either 1 or -1 from hierarchy table
<code>level</code>	Numeric vector from hierarchy table
<code>mapsInput</code>	All codes in mapsFrom not in mapsTo (created automatically when NULL) and possibly other codes in input data.
<code>inputInOutput</code>	When FALSE all output rows represent codes in mapsTo
<code>keepCodes</code>	To prevent some codes to be removed when inputInOutput = TRUE
<code>unionComplement</code>	When TRUE, sign means union and complement instead of addition or subtraction (see note)
<code>reOrder</code>	When TRUE (FALSE is default) output codes are ordered differently, more similar to a usual model matrix ordering.

## Details

The elements of the matrix specify how columns contribute to rows.

## Value

A sparse matrix with row and column and names

## Note

With `unionComplement` = FALSE (default), the sign of each mapping specifies the contribution as addition or subtraction. Thus, values above one and negative values in output can occur. With `unionComplement` = TRUE, positive is treated as union and negative as complement. Then 0 and 1 are the only possible elements in the output matrix.

## Author(s)

Øyvind Langsrød

## Examples

```
# A hierarchy table
h <- SSBtoolsData("FIFA2018ABCD")

DummyHierarchy(h$mapsFrom, h$mapsTo, h$sign, h$level)
DummyHierarchy(h$mapsFrom, h$mapsTo, h$sign, h$level, inputInOutput = TRUE)
DummyHierarchy(h$mapsFrom, h$mapsTo, h$sign, h$level, keepCodes = c("Portugal", "Spain"))

# Extend the hierarchy table to illustrate the effect of unionComplement
h2 <- rbind(data.frame(mapsFrom = c("EU", "Schengen"), mapsTo = "EUandSchengen",
                       sign = 1, level = 3), h)

DummyHierarchy(h2$mapsFrom, h2$mapsTo, h2$sign, h2$level)
```

```

DummyHierarchy(h2$mapsFrom, h2$mapsTo, h2$sign, h2$level, unionComplement = TRUE)

#' # Extend mapsInput - leading to zero columns.
DummyHierarchy(h$mapsFrom, h$mapsTo, h$sign, h$level,
  mapsInput = c(h$mapsFrom[!(h$mapsFrom %in% h$mapsTo)], "Norway", "Finland"))

```

---

FactorLevCorr      *Factor level correlation*

---

## Description

A sort of correlation matrix useful to detect (hierarchical) relationships between the levels of factor variables.

## Usage

```
FactorLevCorr(x)
```

## Arguments

x	Input matrix or data frame containing the variables
---	---

## Value

Output is a sort of correlation matrix.

Here we refer to  $n_i$  as the number of present levels of variable  $i$  (the number of unique elements) and we refer to  $n_{ij}$  as the number of present levels obtained by crossing variable  $i$  and variable  $j$  (the number unique rows of  $x[,c(i,j)]$ ).

The diagonal elements of the output matrix contains the number of present levels of each variable ( $=n_i$ ).

The absolute values of off-diagonal elements:

0	when $n_{ij} = n_i * n_j$
1	when $n_{ij} = \max(n_i, n_j)$

Other values    Computed as  $(n_i * n_j - n_{ij}) / (n_i * n_j - \max(n_i, n_j))$

So 0 means that all possible level combinations exist in the data and 1 means that the two variables are hierarchically related.

The sign of off-diagonal elements:

positive	when $n_i < n_j$
negative	when $n_i > n_j$

In cases where  $n_i = n_j$  elements will be positive above the diagonal and negative below.

## Author(s)

Øyvind Langsrød

### Examples

```
x <- rep(c("A", "B", "C"), 3)
y <- rep(c(11, 22, 11), 3)
z <- c(1, 1, 1, 2, 2, 2, 3, 3, 3)
zy <- paste(z, y, sep="")
m <- cbind(x, y, z, zy)
FactorLevCorr(m)
```

**FindCommonCells**

*Finding commonCells*

### Description

Finding lists defining common cells as needed for the input parameter commonCells to the function protectLinkedTables in package sdcTable. The function handles two tables based on the same main variables but possibly different aggregating variables.

### Usage

```
FindCommonCells(dimList1, dimList2)
```

### Arguments

dimList1	As input parameter dimList to the function makeProblem in package sdcTable.
dimList2	Another dimList with the same names and using the same level names.

### Value

Output is a list according to the specifications in sdcTable.

### Author(s)

Øyvind Langsrud

### Examples

```
x <- rep(c('A', 'B', 'C'), 3)
y <- rep(c(11, 22, 11), 3)
z <- c(1, 1, 1, 2, 2, 2, 3, 3, 3)
zy <- paste(z, y, sep='')
m <- cbind(x, y, z, zy)
fg <- FindTableGroup(m, findLinked=TRUE)
dimLists <- FindDimLists(m, fg$groupVarInd)
# Using table1 and table2 in this example cause error,
# but in other cases this may work well
try(FindCommonCells(dimLists[fg$table$table1], dimLists[fg$table$table2]))
FindCommonCells(dimLists[c(1, 2)], dimLists[c(1, 3)])
```

---

FindDimLists*Finding dimList*

---

**Description**

Finding lists of level-hierarchy as needed for the input parameter dimList to the function makeProblem in package sdcTable

**Usage**

```
FindDimLists(
  x,
  groupVarInd = HierarchicalGroups(x = x),
  addName = FALSE,
  sep = ".",
  xReturn = FALSE,
  total = "Total"
)
```

**Arguments**

x	Matrix or data frame containing the variables (micro data or cell counts data).
groupVarInd	List of vectors of indices defining the hierarchical variable groups.
addName	When TRUE the variable name is added to the level names, except for variables with most levels.
sep	A character string to separate when addName apply.
xReturn	When TRUE x is also in output, possibly changed according to addName.
total	String used to name totals.

**Value**

Output is a list according to the specifications in sdcTable. When xReturn is TRUE output has an extra list level and x is the first element.

**Author(s)**

Øyvind Langsrud

**Examples**

```
x <- rep(c('A', 'B', 'C'), 3)
y <- rep(c(11, 22, 11), 3)
z <- c(1, 1, 1, 2, 2, 2, 3, 3, 3)
zy <- paste(z, y, sep = '')
m <- cbind(x, y, z, zy)
FindDimLists(m)
```

**FindTableGroup***Finding table(s) of hierarchical variable groups***Description**

A single table or two linked tables are found

**Usage**

```
FindTableGroup(
  x = NULL,
  findLinked = FALSE,
  mainName = TRUE,
  fCorr = FactorLevCorr(x),
  CheckHandling = warning
)
```

**Arguments**

<code>x</code>	Matrix or data frame containing the variables
<code>findLinked</code>	When TRUE, two linked tables can be in output
<code>mainName</code>	When TRUE the groupVarInd ouput is named according to first variable in group.
<code>fCorr</code>	When non-null x is not needed as input.
<code>CheckHandling</code>	Function (warning or stop) to be used in problematic situations.

**Value**

Output is a list with items

<code>groupVarInd</code>	List defining the hierarchical variable groups. First variable has most levels.
<code>table</code>	List containing one or two tables. These tables are coded as indices referring to elements of groupVarInd.

**Author(s)**

Øyvind Langsrød

**Examples**

```
x <- rep(c('A', 'B', 'C'), 3)
y <- rep(c(11, 22, 11), 3)
z <- c(1, 1, 1, 2, 2, 2, 3, 3, 3)
zy <- paste(z, y, sep = '')
m <- cbind(x, y, z, zy)
FindTableGroup(m)
FindTableGroup(m, findLinked=TRUE)
```

---

FormulaSums	<i>Sums (aggregates) and/or sparse model matrix with possible cross table</i>
-------------	---

---

## Description

By default this function return sums if the formula contains a response part and a model matrix otherwise

## Usage

```
FormulaSums(
  data,
  formula,
  makeNames = TRUE,
  crossTable = FALSE,
  total = "Total",
  printInc = FALSE,
  dropResponse = FALSE,
  makeModelMatrix = NULL,
  sep = "-",
  sepCross = ":")

Formula2ModelMatrix(data, formula, dropResponse = TRUE, ...)
```

## Arguments

data	data frame
formula	A model formula
makeNames	Column/row names made when TRUE
crossTable	Cross table in output when TRUE
total	String used to name totals
printInc	Printing "..." to console when TRUE
dropResponse	When TRUE response part of formula ignored.
makeModelMatrix	Make model matrix when TRUE. NULL means automatic.
sep	String to separate when creating column names
sepCross	String to separate when creating column names involving crossing
...	Further arguments to be passed to FormulaSums

## Details

The model matrix is constructed by calling fac2sparse() repeatedly. The sums are computed by calling aggregate() repeatedly. Hierarchical variables handled when constructing cross table. Column names constructed from the cross table.

**Value**

A matrix of sums, a sparse model matrix or a list of three elements (model matrix, cross table and sums).

**Author(s)**

Øyvind Langsrød

**Examples**

```
x <- SSBtoolsData("sprt_emp_withEU")

FormulaSums(x, ths_per ~ year*geo + year*eu)
FormulaSums(x, ~ year*age*eu)
FormulaSums(x, ths_per ~ year*age*geo + year*age*eu, crossTable = TRUE, makeModelMatrix = TRUE)
FormulaSums(x, ths_per ~ year:age:geo -1)
```

GaussSuppression

*Secondary suppression by Gaussian elimination*

**Description**

Sequentially the secondary suppression candidates (columns in x) are used to reduce the x-matrix by Gaussian elimination. Candidates who completely eliminate one or more primary suppressed cells (columns in x) are omitted and made secondary suppressed. This ensures that the primary suppressed cells do not depend linearly on the non-suppressed cells. How to order the input candidates is an important choice. The singleton problem and the related problem of zeros are also handled.

**Usage**

```
GaussSuppression(
  x,
  candidates = 1:ncol(x),
  primary = NULL,
  forced = NULL,
  hidden = NULL,
  singleton = rep(FALSE, NROW(x)),
  singletonMethod = "anySum",
  printInc = TRUE,
  ...
)
```

**Arguments**

- |            |   |
|------------|---|
| x          | Matrix that relates cells to be published or suppressed to inner cells. $y_{Publish} = crossprod(x, y_{Inner})$ |
| candidates | Indices of candidates for secondary suppression   |

primary	Indices of primary suppressed cells
forced	Indices forced to be not suppressed
hidden	Indices to be removed from the above candidates input (see details)
singleton	Logical vector specifying inner cells for singleton handling. Normally, this means cells with 1s when 0s are non-suppressed and cells with 0s when 0s are suppressed.
singletonMethod	Method for handling the problem of singletons and zeros: "anySum" (default), "subSum", "subSpace" or "none" (see details).
printInc	Printing "..." to console when TRUE
...	Extra unused parameters

## Details

It is possible to specify too many (all) indices as candidates. Indices specified as primary or hidden will be removed. Hidden indices (not candidates or primary) refer to cells that will not be published, but do not need protection. The singleton method "subSum" makes new imaginary primary suppressed cells, which are the sum of the singletons within each group. The "subSpace" method is conservative and ignores the singleton dimensions when looking for linear dependency. The default method, "anySum", is between the other two. Instead of making imaginary cells of sums within groups, the aim is to handle all possible sums, also across groups. In addition, "subSumSpace" and "subSumAny" are possible methods, primarily for testing. These methods are similar to "subSpace" and "anySum", and additional cells are created as in "subSum". It is believed that the extra cells are redundant.

## Value

Secondary suppression indices

## Examples

```
# Input data
df <- data.frame(values = c(1, 1, 1, 5, 5, 9, 9, 9, 9, 9, 0, 0, 0, 7, 7),
                  var1 = rep(1:3, each = 5),
                  var2 = c("A", "B", "C", "D", "E"), stringsAsFactors = FALSE)

# Make output data frame and x
fs <- FormulaSums(df, values ~ var1 * var2, crossTable = TRUE, makeModelMatrix = TRUE)
x <- fs$modelMatrix
datF <- data.frame(fs$crossTable, values = as.vector(fs$allSums))

# Add primary suppression
datF$primary <- datF$values
datF$primary[datF$values < 5 & datF$values > 0] <- NA
datF$suppressedA <- datF$primary
datF$suppressedB <- datF$primary
datF$suppressedC <- datF$primary

# zero secondary suppressed
```

```

datF$suppressedA[GaussSuppression(x, primary = is.na(datF$primary))] <- NA

# zero not secondary suppressed by first in ordering
datF$suppressedB[GaussSuppression(x, c(which(datF$values == 0), which(datF$values > 0)),
                                     primary = is.na(datF$primary))] <- NA

# with singleton
datF$suppressedC[GaussSuppression(x, c(which(datF$values == 0), which(datF$values > 0)),
                                     primary = is.na(datF$primary), singleton = df$values == 1)] <- NA

datF

```

## Description

According to the (factor) levels of the variables

## Usage

```

HierarchicalGroups(
  x = NULL,
  mainName = TRUE,
  eachName = FALSE,
  fCorr = FactorLevCorr(x)
)

```

## Arguments

<code>x</code>	Matrix or data frame containing the variables
<code>mainName</code>	When TRUE output list is named according to first variable in group.
<code>eachName</code>	When TRUE variable names in output instead of indices.
<code>fCorr</code>	When non-null x is not needed as input.

## Value

Output is a list containing the groups. First variable has most levels.

## Author(s)

Øyvind Langsrød

## Examples

```
x <- rep(c("A", "B", "C"), 3)
y <- rep(c(11, 22, 11), 3)
z <- c(1, 1, 1, 2, 2, 2, 3, 3, 3)
zy <- paste(z, y, sep="")
m <- cbind(x, y, z, zy)
HierarchicalGroups(m)
```

## HierarchicalWildcardGlobbing

*Find variable combinations by advanced wildcard/globbing specifications.*

## Description

Find combinations present in an input data frame or, when input is a list, find all possible combinations that meet the requirements.

## Usage

```
HierarchicalWildcardGlobbing(
  z,
  wg,
  useUnique = NULL,
  useFactor = FALSE,
  makeWarning = TRUE,
  printInfo = FALSE,
  useMatrixToDataFrame = TRUE
)
```

## Arguments

<code>z</code>	list or data.frame
<code>wg</code>	data.frame with data globbing and wildcards
<code>useUnique</code>	Logical variable about recoding within the algorithm. By default (NULL) an automatic decision is made.
<code>useFactor</code>	When TRUE, internal factor recoding is used.
<code>makeWarning</code>	When TRUE, warning is made in cases of unused variables. Only variables common to <code>z</code> and <code>wg</code> are used.
<code>printInfo</code>	When TRUE, information is printed during the process.
<code>useMatrixToDataFrame</code>	When TRUE, special functions (DataFrameToMatrix/MatrixToDataFrame) for improving speed and memory is utilized.

## Details

The final variable combinations must meet the requirements in each positive sign group and must not match the requirements in the negative sign groups. The function is implemented by calling [WildcardGlobbing](#) several times within an algorithm that uses hierarchical clustering ([hclust](#)).

## Value

`data.frame`

## Author(s)

Øyvind Langsrud

## Examples

```
# useUnique=NULL betyr valg ut fra antall rader i kombinasjonsfil
data(precip)
data(mtcars)
codes <- as.character(c(100, 200, 300, 600, 700, 101, 102, 103, 104, 134, 647, 783,
                      13401, 13402, 64701, 64702))

# Create list input
zList <- list(car = rownames(mtcars), wt = as.character(1000 * mtcars$wt),
              city = names(precip), code = codes)

# Create data.frame input
m <- cbind(car = rownames(mtcars), wt = as.character(1000 * mtcars$wt))
zFrame <- data.frame(m[rep(1:NROW(m), each = 35), ],
                      city = names(precip), code = codes, stringsAsFactors = FALSE)

# Create globbing/wildcards input
wg <- data.frame(rbind(c("Merc*", "", "", "?00"), 
                        c("F*", "", "", "?????"),
                        c("", "??0", "C*", ""),
                        c("", "", "!Co*"),
                        c("", "", "?i*", "????2"),
                        c("", "", "?h*", "????1")),
                  sign = c("+", "+", "+", "+", "-", "-"), stringsAsFactors = FALSE)
names(wg)[1:4] <- names(zList)

# =====
#   Finding unique combinations present in the input data frame
# =====

# Using first row of wg. Combinations of car starting with Merc
# and three-digit code ending with 00
HierarchicalWildcardGlobbing(zFrame[, c(1, 4)], wg[1, c(1, 4, 5)])
```

```

# Using first row of wg. Combinations of all four variables
HierarchicalWildcardGlobbing(zFrame, wg[1, ])

# More combinations when using second row also
HierarchicalWildcardGlobbing(zFrame, wg[1:2, ])

# Less combinations when using third row also
# since last digit of wt must be 0 and only cities starting with C
HierarchicalWildcardGlobbing(zFrame, wg[1:3, ])

# Less combinations when using fourth row also since city cannot start with Co
HierarchicalWildcardGlobbing(zFrame, wg[1:4, ])

# Less combinations when using fourth row also
# since specific combinations of city and code are removed
HierarchicalWildcardGlobbing(zFrame, wg)

# =====
# Using list input to create all possible combinations
# =====

dim(HierarchicalWildcardGlobbing(zList, wg))

# same result with as.list since same unique values of each variable
dim(HierarchicalWildcardGlobbing(as.list(zFrame), wg))

```

**Hierarchies2ModelMatrix***Model matrix representing crossed hierarchies***Description**

Make a model matrix,  $x$ , that corresponds to data and represents all hierarchies crossed. This means that aggregates corresponding to numerical variables can be computed as  $t(x) \%*\% y$ , where  $y$  is a matrix with one column for each numerical variable.

**Usage**

```

Hierarchies2ModelMatrix(
  data,
  hierarchies,
  inputInOutput = TRUE,
  crossTable = FALSE,
  total = "Total",
  hierarchyVarNames = c(mapsFrom = "mapsFrom", mapsTo = "mapsTo", sign = "sign", level
    = "level"),

```

```

    unionComplement = FALSE,
    reOrder = TRUE,
    select = NULL,
    selectionByMultiplicationLimit = 10^7,
    makeColnames = TRUE,
    verbose = FALSE
)

```

## Arguments

<code>data</code>	Matrix or data frame with data containing codes of relevant variables
<code>hierarchies</code>	List of hierarchies, which can be converted by <a href="#">AutoHierarchies</a> . Thus, the variables can also be coded by "rowFactor" or "", which correspond to using the categories in the data.
<code>inputInOutput</code>	Logical vector (possibly recycled) for each element of hierarchies. TRUE means that codes from input are included in output. Values corresponding to "rowFactor" or "" are ignored.
<code>crossTable</code>	Cross table in output when TRUE
<code>total</code>	Vector of total codes (possibly recycled) used when running <a href="#">Hrc2DimList</a>
<code>hierarchyVarNames</code>	Variable names in the hierarchy tables as in <a href="#">HierarchyFix</a>
<code>unionComplement</code>	Logical vector (possibly recycled) for each element of hierarchies. When TRUE, sign means union and complement instead of addition or subtraction. Values corresponding to "rowFactor" and "colFactor" are ignored.
<code>reOrder</code>	When TRUE (default) output codes are ordered in a way similar to a usual model matrix ordering.
<code>select</code>	Data frame specifying variable combinations for output.
<code>selectionByMultiplicationLimit</code>	With non-NULL select and when the number of elements in the model matrix exceeds this limit, the computation is performed by a slower but more memory efficient algorithm.
<code>makeColnames</code>	Colnames included when TRUE (default).
<code>verbose</code>	Whether to print information during calculations. FALSE is default.

## Details

This function makes use of [AutoHierarchies](#) and [HierarchyCompute](#) via [HierarchyComputeDummy](#). Since the dummy matrix is transposed in comparison to [HierarchyCompute](#), the parameter `rowSelect` is renamed to `select` and `makeRownames` is renamed to `makeColnames`.

## Value

A sparse model matrix or a list of two elements (model matrix and cross table)

**Author(s)**

Øyvind Langsrød

**See Also**

[HierarchiesAndFormula2ModelMatrix](#)

**Examples**

```
# Create some input
z <- SSBtoolsData("sprt_emp_withEU")
ageHier <- SSBtoolsData("sprt_emp_ageHier")
geoDimList <- FindDimLists(z[, c("geo", "eu")], total = "Europe")[[1]]

# First example has list output
Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList), inputInOutput = FALSE,
                        crossTable = TRUE)

m1 <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList), inputInOutput = FALSE)
m2 <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList))
m3 <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList, year = ""),
                               inputInOutput = FALSE)
m4 <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList, year = "allYears"),
                               inputInOutput = c(FALSE, FALSE, TRUE))

# Illustrate the effect of unionComplement, geoHier2 as in the examples of HierarchyCompute
geoHier2 <- rbind(data.frame(mapsFrom = c("EU", "Spain"), mapsTo = "EUandSpain", sign = 1),
                   SSBtoolsData("sprt_emp_geoHier")[, -4])
m5 <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoHier2, year = "allYears"),
                               inputInOutput = FALSE) # Spain is counted twice
m6 <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoHier2, year = "allYears"),
                               inputInOutput = FALSE, unionComplement = TRUE)

# Compute aggregates
ths_per <- as.matrix(z[, "ths_per", drop = FALSE]) # matrix with the values to be aggregated
t(m1) %*% ths_per # crossprod(m1, ths_per) is equivalent and faster
t(m2) %*% ths_per
t(m3) %*% ths_per
t(m4) %*% ths_per
t(m5) %*% ths_per
t(m6) %*% ths_per

# Example using the select parameter
select <- data.frame(age = c("Y15-64", "Y15-29", "Y30-64"), geo = c("EU", "nonEU", "Spain"))
m2a <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList), select = select)

# Same result by slower alternative
m2B <- Hierarchies2ModelMatrix(z, list(age = ageHier, geo = geoDimList), crossTable = TRUE)
```

```
m2b <- m2B$modelMatrix[, Match(select, m2B$crossTable), drop = FALSE]
t(m2b) %*% ths_per
```

## HierarchiesAndFormula2ModelMatrix

*Model matrix representing crossed hierarchies according to a formula*

### Description

How to cross the hierarchies are defined by a formula. The formula is automatically simplified when totals are involved.

### Usage

```
HierarchiesAndFormula2ModelMatrix(
  data,
  hierarchies,
  formula,
  inputInOutput = TRUE,
  makeColNames = TRUE,
  crossTable = FALSE,
  total = "Total",
  simplify = TRUE,
  hierarchyVarNames = c(mapsFrom = "mapsFrom", mapsTo = "mapsTo", sign = "sign", level
    = "level"),
  unionComplement = FALSE,
  reOrder = TRUE,
  sep = "-"
)
```

### Arguments

<code>data</code>	Matrix or data frame with data containing codes of relevant variables
<code>hierarchies</code>	List of hierarchies, which can be converted by <a href="#">AutoHierarchies</a> . Thus, the variables can also be coded by "rowFactor" or "", which correspond to using the categories in the data.
<code>formula</code>	A model formula
<code>inputInOutput</code>	Logical vector (possibly recycled) for each element of hierarchies. TRUE means that codes from input are included in output. Values corresponding to "rowFactor" or "" are ignored.
<code>makeColNames</code>	Colnames included when TRUE (default).
<code>crossTable</code>	Cross table in output when TRUE
<code>total</code>	Vector of total codes (possibly recycled) used when running <a href="#">Hrc2DimList</a>
<code>simplify</code>	When TRUE (default) the model can be simplified when total codes are found in the hierarchies (see examples).

<code>hierarchyVarNames</code>	Variable names in the hierarchy tables as in <a href="#">HierarchyFix</a>
<code>unionComplement</code>	Logical vector (possibly recycled) for each element of hierarchies. When TRUE, sign means union and complement instead of addition or subtraction. Values corresponding to "rowFactor" and "colFactor" are ignored.
<code>reOrder</code>	When TRUE (default) output codes are ordered in a way similar to a usual model matrix ordering.
<code>sep</code>	String to separate when creating column names

**Value**

A sparse model matrix or a list of two elements (model matrix and cross table)

**Author(s)**

Øyvind Langsrud

**See Also**

[Hierarchies2ModelMatrix](#), [Formula2ModelMatrix](#).

**Examples**

```
# Create some input
z <- SSBtoolsData("sppt_emp_withEU")
ageHier <- SSBtoolsData("sppt_emp_ageHier")
geoDimList <- FindDimLists(z[, c("geo", "eu")], total = "Europe")[[1]]

# Shorter function name
H <- HierarchiesAndFormula2ModelMatrix

# Small dataset example. Two dimensions.
s <- z[z$geo == "Spain", ]
geoYear <- list(geo = geoDimList, year = "")
m <- H(s, geoYear, ~geo * year, inputInOutput = c(FALSE, TRUE))
print(m, col.names = TRUE)
attr(m, "total")      # Total code 'Europe' is found
attr(m, "startCol")   # Two model terms needed

# Another model and with crossTable in output
H(s, geoYear, ~geo + year, crossTable = TRUE)

# Three dimensions
ageGeoYear <- list(age = ageHier, geo = geoDimList, year = "allYears")
m <- H(z, ageGeoYear, ~age * geo + geo * year)
head(colnames(m))
attr(m, "total")
attr(m, "startCol")

# With simplify = FALSE
```

```

m <- H(z, ageGeoYear, ~age * geo + geo * year, simplify = FALSE)
head(colnames(m))
attr(m, "total")
attr(m, "startCol")

# Compute aggregates
m <- H(z, ageGeoYear, ~geo * age, inputInOutput = c(TRUE, FALSE, TRUE))
t(m) %*% z$ths_per

# Without hierarchies. Only factors.
ageGeoYearFactor <- list(age = "", geo = "", year = "")
t(H(z, ageGeoYearFactor, ~geo * age + year:geo))

```

**Hierarchy2Formula***Hierarchy2Formula***Description**

Conversion between to-from coded hierarchy and formulas written with =, - and +.

**Usage**

```

Hierarchy2Formula(
  x,
  hierarchyVarNames = c(mapsFrom = "mapsFrom", mapsTo = "mapsTo", sign = "sign", level
    = "level")
)
Formula2Hierarchy(s)

```

**Arguments**

x	Data frame with to-from coded hierarchy
hierarchyVarNames	Variable names in the hierarchy tables as in <a href="#">HierarchyFix</a> .
s	Character vector of formulas written with =, - and +.

**Value**

See Arguments

**Author(s)**

Øyvind Langsrød

## Examples

```
x <- SSBtoolsData("sppt_emp_geoHier")
s <- Hierarchy2Formula(x)
Formula2Hierarchy(s)
```

HierarchyCompute

*Hierarchical Computations*

## Description

This function computes aggregates by crossing several hierarchical specifications and factorial variables.

## Usage

```
HierarchyCompute(
  data,
  hierarchies,
  valueVar,
  colVar = NULL,
  rowSelect = NULL,
  colSelect = NULL,
  select = NULL,
  inputInOutput = FALSE,
  output = "data.frame",
  autoLevel = TRUE,
  unionComplement = FALSE,
  constantsInOutput = NULL,
  hierarchyVarNames = c.mapsFrom = "mapsFrom", mapsTo = "mapsTo", sign = "sign", level
    = "level"),
  selectionByMultiplicationLimit = 10^7,
  colNotInDataWarning = TRUE,
  useMatrixToDataFrame = TRUE,
  handleDuplicated = "sum",
  asInput = FALSE,
  verbose = FALSE,
  reOrder = FALSE,
  reduceData = TRUE,
  makeRownames = NULL
)
```

## Arguments

<code>data</code>	The input data frame
<code>hierarchies</code>	A named (names in <code>data</code> ) list with hierarchies. Variables can also be coded by " <code>rowFactor</code> " and " <code>colFactor</code> ".

<code>valueVar</code>	Name of the variable(s) to be aggregated.
<code>colVar</code>	When non-NULL, the function <a href="#">HierarchyCompute2</a> is called. See its documentation for more information.
<code>rowSelect</code>	Data frame specifying variable combinations for output. The colFactor variable is not included. In addition <code>rowSelect=="removeEmpty"</code> removes combinations corresponding to empty rows (only zeros) of <code>dataDummyHierarchy</code> .
<code>colSelect</code>	Vector specifying categories of the colFactor variable for output.
<code>select</code>	Data frame specifying variable combinations for output. The colFactor variable is included.
<code>inputInOutput</code>	Logical vector (possibly recycled) for each element of hierarchies. TRUE means that codes from input are included in output. Values corresponding to "rowFactor" and "colFactor" are ignored.
<code>output</code>	One of "data.frame" (default), "dummyHierarchies", "outputMatrix", "dataDummyHierarchy", "valueMatrix", "fromCrossCode", "toCrossCode", "crossCode" (as toCrossCode), "outputMatrixWithCrossCode", "matrixComponents", "dataDummyHierarchyWithCodeFrame", "dataDummyHierarchyQuick". The latter two do not require <code>valueVar</code> (reduceData set to FALSE).
<code>autoLevel</code>	Logical vector (possibly recycled) for each element of hierarchies. When TRUE, level is computed by automatic method as in <a href="#">HierarchyFix</a> . Values corresponding to "rowFactor" and "colFactor" are ignored.
<code>unionComplement</code>	Logical vector (possibly recycled) for each element of hierarchies. When TRUE, sign means union and complement instead of addition or subtraction as in <a href="#">DummyHierarchy</a> . Values corresponding to "rowFactor" and "colFactor" are ignored.
<code>constantsInOutput</code>	A single row data frame to be combine by the other output.
<code>hierarchyVarNames</code>	Variable names in the hierarchy tables as in <a href="#">HierarchyFix</a> .
<code>selectionByMultiplicationLimit</code>	With non-NULL <code>rowSelect</code> and when the number of elements in <code>dataDummyHierarchy</code> exceeds this limit, the computation is performed by a slower but more memory efficient algorithm.
<code>colNotInDataWarning</code>	When TRUE, warning produced when elements of <code>colSelect</code> are not in data.
<code>useMatrixToDataFrame</code>	When TRUE (default) special functionality for saving time and memory is used.
<code>handleDuplicated</code>	Handling of duplicated code rows in data. One of: "sum" (default), "sumByAggregate", "sumWithWarning", "stop" (error), "single" or "singleWithWarning". With no colFactor sum and sumByAggregate/sumWithWarning are different (original values or aggregates in "valueMatrix"). When single, only one of the values is used (by matrix subsetting).
<code>asInput</code>	When TRUE (FALSE is default) output matrices match input data. Thus <code>valueMatrix = Matrix(data[,valueVar],ncol=1)</code> . Only possible when no colFactor.

verbose	Whether to print information during calculations. FALSE is default.
reOrder	When TRUE (FALSE is default) output codes are ordered differently, more similar to a usual model matrix ordering.
reduceData	When TRUE (default) unnecessary (for the aggregated result) rows of valueMatrix are allowed to be removed.
makeRownames	When TRUE dataDummyHierarchy contains rownames. By default, this is decided based on the parameter output.

## Details

A key element of this function is the matrix multiplication: `outputMatrix = dataDummyHierarchy %*% valueMatrix`. The matrix, `valueMatrix` is a re-organized version of the `valueVar` vector from input. In particular, if a variable is selected as `colFactor`, there is one column for each level of that variable. The matrix, `dataDummyHierarchy` is constructed by crossing dummy coding of hierarchies ([DummyHierarchy](#)) and factorial variables in a way that matches `valueMatrix`. The code combinations corresponding to rows and columns of `dataDummyHierarchy` can be obtained as `toCrossCode` and `fromCrossCode`. In the default data frame output, the `outputMatrix` is stacked to one column and combined with the code combinations of all variables.

## Value

As specified by the parameter `output`

## Author(s)

Øyvind Langsrød

## See Also

[Hierarchies2ModelMatrix](#), [AutoHierarchies](#).

## Examples

```
# Data and hierarchies used in the examples
x <- SSBtoolsData("sprt_emp") # Employment in sport in thousand persons from Eurostat database
geoHier <- SSBtoolsData("sprt_emp_geoHier")
ageHier <- SSBtoolsData("sprt_emp_ageHier")

# Two hierarchies and year as rowFactor
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "rowFactor"), "ths_per")

# Same result with year as colFactor (but columns ordered differently)
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per")

# Internally the computations are different as seen when output='matrixComponents'
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "rowFactor"), "ths_per",
                 output = "matrixComponents")
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per",
                 output = "matrixComponents")
```

```

# Include input age groups by setting inputInOutput = TRUE for this variable
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per",
                 inputInOutput = c(TRUE, FALSE))

# Only input age groups by switching to rowFactor
HierarchyCompute(x, list(age = "rowFactor", geo = geoHier, year = "colFactor"), "ths_per")

# Select some years (colFactor) including a year not in input data (zeros produced)
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per",
                 colSelect = c("2014", "2016", "2018"))

# Select combinations of geo and age including a code not in data or hierarchy (zeros produced)
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per",
                 rowSelect = data.frame(geo = "EU", age = c("Y0-100", "Y15-64", "Y15-29")))

# Select combinations of geo, age and year
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per",
                 select = data.frame(geo = c("EU", "Spain"), age = c("Y15-64", "Y15-29"), year = 2015))

# Extend the hierarchy table to illustrate the effect of unionComplement
# Omit level since this is handled by autoLevel
geoHier2 <- rbind(data.frame(mapsFrom = c("EU", "Spain"), mapsTo = "EUandSpain", sign = 1),
                   geoHier[, -4])

# Spain is counted twice
HierarchyCompute(x, list(age = ageHier, geo = geoHier2, year = "colFactor"), "ths_per")

# Can be seen in the dataDummyHierarchy matrix
HierarchyCompute(x, list(age = ageHier, geo = geoHier2, year = "colFactor"), "ths_per",
                 output = "matrixComponents")

# With unionComplement=TRUE Spain is not counted twice
HierarchyCompute(x, list(age = ageHier, geo = geoHier2, year = "colFactor"), "ths_per",
                 unionComplement = TRUE)

# With constantsInOutput
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "colFactor"), "ths_per",
                 constantsInOutput = data.frame(c1 = "AB", c2 = "CD"))

# More than one valueVar
x$y <- 10*x$ths_per
HierarchyCompute(x, list(age = ageHier, geo = geoHier), c("y", "ths_per"))

```

## Description

Extended variant of [HierarchyCompute](#) with several column variables (not just "colFactor"). Parameter colVar splits the hierarchy variables in two groups and this variable overrides the difference

between "rowFactor" and "colFactor".

## Usage

```
HierarchyCompute2(
  data,
  hierarchies,
  valueVar,
  colVar,
  rowSelect = NULL,
  colSelect = NULL,
  select = NULL,
  output = "data.frame",
  ...
)
```

## Arguments

data	The input data frame
hierarchies	A named list with hierarchies
valueVar	Name of the variable(s) to be aggregated
colVar	Name of the column variable(s)
rowSelect	Data frame specifying variable combinations for output
colSelect	Data frame specifying variable combinations for output
select	Data frame specifying variable combinations for output
output	One of "data.frame" (default), "outputMatrix", "matrixComponents".
...	Further parameters sent to <a href="#">HierarchyCompute</a>

## Details

Within this function, [HierarchyCompute](#) is called two times. By specifying output as "matrixComponents", output from the two runs are retuned as a list with elements hcRow and hcCol. The matrix multiplication in [HierarchyCompute](#) is extended to  $\text{outputMatrix} = \text{hcRow} \$ \text{data} \$ \text{DummyHierarchy} \%*% \text{hcRow} \$ \text{valueMatrix} \%*% t(\text{hcCol} \$ \text{data} \$ \text{DummyHierarchy})$ . This is modified in cases with more than a single valueVar.

## Value

As specified by the parameter output

## Note

There is no need to call [HierarchyCompute2](#) directly. The main function [HierarchyCompute](#) can be used instead.

## Author(s)

Øyvind Langsrød

**See Also**

[Hierarchies2ModelMatrix](#), [AutoHierarchies](#).

**Examples**

```
x <- SSBtoolsData("sprt_emp")
geoHier <- SSBtoolsData("sprt_emp_geoHier")
ageHier <- SSBtoolsData("sprt_emp_ageHier")

HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "rowFactor"), "ths_per",
                 colVar = c("age", "year"))
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "rowFactor"), "ths_per",
                 colVar = c("age", "geo"))
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "rowFactor"), "ths_per",
                 colVar = c("age", "year"), output = "matrixComponents")
HierarchyCompute(x, list(age = ageHier, geo = geoHier, year = "rowFactor"), "ths_per",
                 colVar = c("age", "geo"), output = "matrixComponents")
```

**HierarchyFix**

*Change the hierarchy table to follow the standard*

**Description**

Make sure that variable names and sign coding follow an internal standard. Level may be computed automatically

**Usage**

```
HierarchyFix(
  hierarchy,
  hierarchyVarNames = c(mapsFrom = "mapsFrom", mapsTo = "mapsTo", sign = "sign", level
    = "level"),
  autoLevel = TRUE
)
```

**Arguments**

<code>hierarchy</code>	data frame with hierarchy table
<code>hierarchyVarNames</code>	variable names
<code>autoLevel</code>	When TRUE, level is computed by automatic method

**Value**

data frame with hierarchy table

**Author(s)**

Øyvind Langsrud

**Examples**

```
# Make input data by changing variable names and sign coding.
h <- SSBtoolsData("FIFA2018ABCD")[, 1:3]
names(h)[1:2] <- c("from", "to")
minus <- h$sign < 0
h$sign <- "+"
h$sign[minus] <- "-"

# Run HierarchyFix - Two levels created
HierarchyFix(h, c(mapsFrom = "from", mapsTo = "to", sign = "sign"))

# Extend the hierarchy table
h2 <- rbind(data.frame(from = c("Oceania", "Asia", "Africa", "America", "Europe"),
                        to = "World", sign = "+"),
             data.frame(from = c("World", "Europe"),
                        to = "nonEurope", sign = c("+", "-")), h)

# Run HierarchyFix - Three levels created
HierarchyFix(h2, c(mapsFrom = "from", mapsTo = "to", sign = "sign"))
```

**MakeHierFormula**

*Make model formula from data taking into account hierarchical variables*

**Description**

Make model formula from data taking into account hierarchical variables

**Usage**

```
MakeHierFormula(
  data = NULL,
  hGroups = HierarchicalGroups2(data),
  n = length(hGroups),
  sim = TRUE
)
```

**Arguments**

<b>data</b>	data frame
<b>hGroups</b>	Output from HierarchicalGroups2()
<b>n</b>	Interaction level or 0 (all levels)
<b>sim</b>	Include "~" when TRUE

**Value**

Formula as character string

**Author(s)**

Øyvind Langsrud

**Examples**

```
x <- SSBtoolsData("sprt_emp_withEU")[, -4]
MakeHierFormula(x)
MakeHierFormula(x, n = 2)
MakeHierFormula(x, n = 0)
```

**Match**

*Matching rows in data frames*

**Description**

The algorithm is based on converting variable combinations to whole numbers. The final matching is performed using [match](#).

**Usage**

```
Match(x, y)
```

**Arguments**

x	data frame
y	data frame

**Details**

When the result of multiplying together the number of unique values in each column of x exceeds 9E15 (largest value stored exactly by the numeric data type), the algorithm is recursive.

**Value**

An integer vector giving the position in y of the first match if there is a match, otherwise NA.

**Author(s)**

Øyvind Langsrud

## Examples

```

a <- data.frame(x = c("a", "b", "c"), y = c("A", "B"), z = 1:6)
b <- data.frame(x = c("b", "c"), y = c("B", "K", "A", "B"), z = c(2, 3, 5, 6))

Match(a, b)
Match(b, a)

# Slower alternative
match(data.frame(t(a), stringsAsFactors = FALSE), data.frame(t(b), stringsAsFactors = FALSE))
match(data.frame(t(b), stringsAsFactors = FALSE), data.frame(t(a), stringsAsFactors = FALSE))

# More comprehensive example (n, m and k may be changed)
n <- 10^4
m <- 10^3
k <- 10^2
data(precip)
data(mtcars)
y <- data.frame(car = sample(rownames(mtcars), n, replace = TRUE),
                 city = sample(names(precip), n, replace = TRUE),
                 n = rep_len(1:k, n), a = rep_len(c("A", "B", "C", "D"), n),
                 b = rep_len(as.character(rnorm(1000)), n),
                 d = sample.int(k + 10, n, replace = TRUE),
                 e = paste(sample.int(k * 2, n, replace = TRUE),
                           rep_len(c("Green", "Red", "Blue"), n), sep = "_"),
                 r = rnorm(k)^99)
x <- y[sample.int(n, m), ]
row.names(x) <- NULL
ix <- Match(x, y)

```

## Description

Functions to generate increasing sequences

## Usage

```
matlabColon(from, to)

SeqInc(from, to)
```

## Arguments

<code>from</code>	numeric. The start value
<code>to</code>	numeric. The end value.

**Details**

`matlabColon(a,b)` returns `a:b` (R's version) unless  $a > b$ , in which case it returns `integer(0)`. `SeqInc(a,b)` is similar, but results in error when the calculated length of the sequence (1+to-from) is negative.

**Value**

A numeric vector, possibly empty.

**Author(s)**

Bjørn-Helge Mevik (`matlabColon`) and Øyvind Langsrød (`SeqInc`)

**See Also**

[seq](#)

**Examples**

```
identical(3:5, matlabColon(3, 5)) ## => TRUE
3:1 ## => 3 2 1
matlabColon(3, 1) ## => integer(0)
try(SeqInc(3, 1)) ## => Error
SeqInc(3, 2)      ## => integer(0)
```

**Matrix2list**

*Convert matrix to sparse list*

**Description**

Convert matrix to sparse list

**Usage**

`Matrix2list(x)`

`Matrix2listInt(x)`

**Arguments**

`x` Input matrix

**Details**

Within the function, the input matrix is first converted to a `dgTMatrix` matrix (Matrix package).

**Value**

A two-element list: List of row numbers (`r`) and a list of numeric or integer values (`x`)

**Note**

`Matrix2listInt` converts the values to integers by `as.integer` and no checking is performed. Thus, zeros are possible.

**Author(s)**

Øyvind Langsrud

**Examples**

```
m = matrix(c(0.5, 1.1, 3.14, 0, 0, 0, 0, 4, 5), 3, 3)
Matrix2list(m)
Matrix2listInt(m)
```

---

Number	<i>Adding leading zeros</i>
--------	-----------------------------

---

**Description**

Adding leading zeros

**Usage**

```
Number(n, width = 3)
```

**Arguments**

n	numeric vector of whole numbers
width	width

**Value**

Character vector

**Author(s)**

Øyvind Langsrud

**Examples**

```
Number(1:3)
```

**RbindAll***Combining several data frames when the columns don't match***Description**

Combining several data frames when the columns don't match

**Usage**

```
RbindAll(...)
```

**Arguments**

...	Several data frames as several input parameters or a list of data frames
-----	--

**Value**

A single data frame

**Note**

The function is an extended version of rbind.all.columns at <https://amywhiteheadresearch.wordpress.com/2013/05/13/combining-dataframes-when-the-columns-dont-match/>

**Author(s)**

Øyvind Langsrød

**See Also**

[CbindIdMatch](#) (same example data)

**Examples**

```
zA <- data.frame(idA = 1:10, idB = rep(10 * (1:5), 2), idC = rep(c(100, 200), 5),
                   idC2 = c(100, rep(200, 9)), idC3 = rep(100, 10),
                   idD = 99, x = round(rnorm(10), 3), xA = round(runif(10), 2))
zB <- data.frame(idB = 10 * (1:5), x = round(rnorm(5), 3), xB = round(runif(5), 2))
zC <- data.frame(idC = c(100, 200), x = round(rnorm(2), 3), xC = round(runif(2), 2))
zD <- data.frame(idD = 99, x = round(rnorm(1), 3), xD = round(runif(1), 2))
RbindAll(zA, zB, zC, zD)
RbindAll(list(zA, zB, zC, zD))
```

---

RowGroups	<i>Create numbering according to unique rows</i>
-----------	--

---

### Description

Create numbering according to unique rows

### Usage

```
RowGroups(x, returnGroups = FALSE, returnGroupsId = FALSE)
```

### Arguments

- |                |   |
|----------------|---|
| x              | Data frame or matrix                        |
| returnGroups   | When TRUE unique rows are returned          |
| returnGroupsId | When TRUE Index of unique rows are returned |

### Value

A vector with the numbering or, according to the arguments, a list with more output.

### Author(s)

Øyvind Langsrud

### Examples

```
a <- data.frame(x = c("a", "b"), y = c("A", "B", "A"), z = rep(1:4, 3))
RowGroups(a)
RowGroups(a, TRUE)
RowGroups(a[, 1:2], TRUE, TRUE)
RowGroups(a[, 1, drop = FALSE], TRUE)
```

---

SSBtoolsData	<i>Function that returns a dataset</i>
--------------	--

---

### Description

Function that returns a dataset

### Usage

```
SSBtoolsData(dataset)
```

### Arguments

- |         |  |
|---------|--|
| dataset | Name of data set within the SSBtools package |
|---------|--|

## Details

**FIFA2018ABCD:** A hierarchy table based on countries within groups A-D in the football championship, 2018 FIFA World Cup.

**sprt\_emp:** Employment in sport in thousand persons. Data from Eurostat database.

**sprt\_emp\_geoHier:** Country hierarchy for the employment in sport data.

**sprt\_emp\_ageHier:** Age hierarchy for the employment in sport data.

**sprt\_emp\_withEU:** The data set sprt\_emp extended with a EU variable.

## Value

data frame

## Author(s)

Øyvind Langsrud

## Examples

```
SSBtoolsData("FIFA2018ABCD")
SSBtoolsData("sprt_emp")
SSBtoolsData("sprt_emp_geoHier")
SSBtoolsData("sprt_emp_ageHier")
SSBtoolsData("sprt_emp_withEU")
```

Stack

*Stack columns from a data frame and include variables.*

## Description

Stack columns from a data frame and include variables.

## Usage

```
Stack(
  data,
  stackVar = 1:NCOL(data),
  blockVar = integer(0),
  rowData = data.frame(stackVar)[, integer(0), drop = FALSE],
  valueName = "values",
  indName = "ind"
)
```

**Arguments**

<code>data</code>	A data frame
<code>stackVar</code>	Indices of variables to be stacked
<code>blockVar</code>	Indices of variables to be replicated
<code>rowData</code>	A separate data frame where NROW(rowData)=length(stackVar) such that each row may contain multiple information of each stackVar variable. The output data frame will contain an extended variant of rowData.
<code>valueName</code>	Name of the stacked/concatenated output variable
<code>indName</code>	Name of the output variable with information of which vector in x the observation originated. When indName is NULL this variable is not included in output.

**Value**

A data frame where the variable ordering corresponds to: blockVar, rowData, valueName, indName

**Author(s)**

Øyvind Langsrød

**See Also**

[Unstack](#)

**Examples**

```

z <- data.frame(n=c(10,20,30), ssb=c('S','S','B'),
Ayes=1:3,Ano=4:6,Byes=7:9,Bno=10:12)
zRow <- data.frame(letter=c('A','A','B','B'),answer=c('yes','no','yes','no') )

x <- Stack(z,3:6,1:2,zRow)

Unstack(x,6,3:4,numeric(0),1:2)
Unstack(x,6,5,numeric(0),1:2)
Unstack(x,6,3:4,5,1:2)

```

[Unstack](#)

*Unstack a column from a data frame and include additional variables.*

**Description**

Unstack a column from a data frame and include additional variables.

**Usage**

```
Unstack(
  data,
  mainVar = 1,
  stackVar = (1:NCOL(data))[-mainVar],
  extraVar = integer(0),
  blockVar = integer(0),
  sep = "_",
  returnRowData = TRUE,
  sorted = FALSE
)
```

**Arguments**

<code>data</code>	A data frame
<code>mainVar</code>	Index of the variable to be unstacked
<code>stackVar</code>	Index of variables defining the unstack grouping
<code>extraVar</code>	Indices of within-replicated variables to be added to the rowData output
<code>blockVar</code>	Indices of between-replicated variables to be added to the data output
<code>sep</code>	A character string to separate when creating variable names
<code>returnRowData</code>	When FALSE output is no list, but only data
<code>sorted</code>	When TRUE the created variables is in sorted order. Otherwise input order is used.

**Value**

When `returnRowData=TRUE` output is list of two elements.

<code>data</code>	Unstacked data
<code>rowData</code>	A separate data frame with one row for each unstack grouping composed of the <code>stackVar</code> variables

**Author(s)**

Øyvind Langsrød

**See Also**

[Stack](#) (examples)

---

<b>WildcardGlobbing</b>	<i>Row selection by wildcard/globbing</i>
-------------------------	---

---

## Description

The selected rows match combined requirements for all variables.

## Usage

```
WildcardGlobbing(x, wg, sign = TRUE, invert = "!")
```

## Arguments

x	data.frame with character data
wg	data.frame with wildcard/globbing
sign	When FALSE, the result is inverted.
invert	Character to invert each single selection.

## Details

This function is used by [HierarchicalWildcardGlobbing](#) and [WildcardGlobbingVector](#) and make use of [grepl](#) and [glob2rx](#).

## Value

Logical vector defining subset of rows.

## Author(s)

Øyvind Langsrød

## Examples

```
# Create data input
data(precip)
data(mtcars)
x <- data.frame(car = rownames(mtcars)[rep(1:NROW(mtcars), each = 35)], city = names(precip),
                 stringsAsFactors = FALSE)

# Create globbing/wildcards input
wg <- data.frame(rbind(c("Merc*", "C*"), c("F*", "??????"), c("!????????*", "!??????*")),
                  stringsAsFactors = FALSE)
names(wg) <- names(x)

# Select the following combinations:
# - Cars starting with Merc and cities starting with C
# - Cars starting with F and six-letter cities
# - Cars with less than nine letters and cities with less than seven letters
x[WildcardGlobbing(x, wg), ]
```

**WildcardGlobbingVector***Selection of elements by wildcard/globbing***Description**

Selection of elements by wildcard/globbing

**Usage**`WildcardGlobbingVector(x, wg, negSign = "-", invert = "!")`**Arguments**

<code>x</code>	Character vector
<code>wg</code>	Character vector with wildcard/globbing
<code>negSign</code>	Character representing selection to be removed
<code>invert</code>	Character to invert each single selection.

**Value**vector with selected elements of `x`**Author(s)**

Øyvind Langsrød

**Examples**

```

data(precip)
x <- names(precip)

# Select the cities starting with B, C and Sa.
WildcardGlobbingVector(x, c("B*", "C*", "Sa*"))

# Remove from the selection cities with o and t in position 2 and 4, respectively.
WildcardGlobbingVector(x, c("B*", "C*", "Sa*", "-?o*", "-???t*"))

# Add to the selection cities not having six or more letters.
WildcardGlobbingVector(x, c("B*", "C*", "Sa*", "-?o*", "-???t*", "!??????*"))

```

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