

Intro to DatABEL

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Contents

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> library(DatABEL)
> make_random_matrix <- function(range_dim1 = c(2, 10), range_dim2 = c(2,
+   10), range_data = c(-10, 10), type = "double") {
+   dim1 <- round(runif(1, range_dim1[1], range_dim1[2]))
+   dim2 <- round(runif(1, range_dim2[1], range_dim2[2]))
+   data <- runif(dim1 * dim2, range_data[1], range_data[2])
+   data <- as(data, type)
+   data <- matrix(data, nrow = dim1, ncol = dim2)
+   namesCol <- paste("col", c(1:dim2), sep = "_")
+   namesRow <- paste("row", c(1:dim1), sep = "_")
+   dimnames(data) <- list(namesRow, namesCol)
+   return(data)
+ }
> testmatr <- make_random_matrix()
> testmatr

      col_1      col_2      col_3      col_4      col_5
row_1 -8.510850 -7.371706 -0.5829301 6.774361  1.793244
row_2 -5.033636  9.213437  0.1687509 5.115450 -5.191560

> test_fv <- as(testmatr, "datababel_base_R")

[1] "./tmp269886"
WARNING: you appear to work on 32-bit system... large files not supported
WARNING: you appear to work on 32-bit system... large files not supported
finalizing AbstractMatrix: 0x16ac800
coersion from 'matrix' to 'datababel_base_R' of type DOUBLE ; object connected to file ./tmp269886

> test_fv

backingfilename = ./tmp269886
cachesizeMb = 1
number of columns (variables) =  5
number of rows (observations) =  2
```

```

Upper-left 5 columns and 2 rows:
[,1]      [,2]      [,3]      [,4]      [,5]
[1,] -8.510850 -7.371706 -0.5829301 6.774361 1.793244
[2,] -5.033636  9.213437  0.1687509 5.115449 -5.191560

> as(test_fv, "matrix")

[,1]      [,2]      [,3]      [,4]      [,5]
[1,] -8.510850 -7.371706 -0.5829301 6.774361 1.793244
[2,] -5.033636  9.213437  0.1687509 5.115449 -5.191560

> abs(testmatr - as(test_fv, "matrix")) < 1e-06

col_1 col_2 col_3 col_4 col_5
row_1 TRUE TRUE TRUE TRUE TRUE
row_2 TRUE TRUE TRUE TRUE TRUE

> write.table(testmatr, file = "test_matrix_dimnames.dat", row.names = TRUE,
+   col.names = TRUE, quote = FALSE)
> text2filevector(infile = "test_matrix_dimnames.dat", outfile = "test_matrix_dimnames",
+   R_matrix = TRUE)

Options in effect:
--infile    = test_matrix_dimnames.dat
--outfile   = test_matrix_dimnames
--skiprows  = 1
--skipcols  = 1
--cnrow     = ON, using line 1 of 'test_matrix_dimnames.dat'
--rncol     = ON, using column 1 of 'test_matrix_dimnames.dat'
--transpose = OFF
--Rmatrix   = ON

WARNING: you appear to work on 32-bit system... large files not supported
WARNING: you appear to work on 32-bit system... large files not supported
WARNING: you appear to work on 32-bit system... large files not supported
WARNING: you appear to work on 32-bit system... large files not supported
unique.names = TRUE
unique.rownames = TRUE
unique.colnames = TRUE
backingfilename = test_matrix_dimnames
cachesizeMb = 1
number of columns (variables) = 5
number of rows (observations) = 2
usedRowIndex: 1 2
usedColIndex: 1 2 3 4 5
Upper-left 5 columns and 2 rows:
[,1]      [,2]      [,3]      [,4]      [,5]
[1,] -8.510850 -7.371706 -0.5829301 6.774361 1.793244
[2,] -5.033636  9.213437  0.1687509 5.115449 -5.191560

```

```

> x <- databel_filtered_R("test_matrix_dimnames")

WARNING: you appear to work on 32-bit system... large files not supported

> x

unique.names = TRUE
unique.rownames = TRUE
unique.colnames = TRUE
backingfilename = test_matrix_dimnames
cachesizeMb = 1
number of columns (variables) = 5
number of rows (observations) = 2
usedRowIndex: 1 2
usedColIndex: 1 2 3 4 5
Upper-left 5 columns and 2 rows:
 [,1]      [,2]      [,3]      [,4]      [,5]
[1,] -8.510850 -7.371706 -0.5829301 6.774361 1.793244
[2,] -5.033636  9.213437  0.1687509 5.115449 -5.191560

> tmp <- as(x, "matrix")
> tmp

      col_1      col_2      col_3      col_4      col_5
row_1 -8.510850 -7.371706 -0.5829301 6.774361 1.793244
row_2 -5.033636  9.213437  0.1687509 5.115449 -5.191560

> abs(testmatr - tmp) < 1e-06

      col_1 col_2 col_3 col_4 col_5
row_1  TRUE  TRUE  TRUE  TRUE  TRUE
row_2  TRUE  TRUE  TRUE  TRUE  TRUE

> text2filevector(infile = "test_matrix_dimnames.dat", outfile = "test_matrix_dimnames_T",
+      R_matrix = TRUE, transpose = TRUE)

Options in effect:
  --infile    = test_matrix_dimnames.dat
  --outfile   = test_matrix_dimnames_T
  --skiprows  = 1
  --skipcols  = 1
  --cnrow     = ON, using line 1 of 'test_matrix_dimnames.dat'
  --rncol     = ON, using column 1 of 'test_matrix_dimnames.dat'
  --transpose = ON
  --Rmatrix   = ON

WARNING: you appear to work on 32-bit system... large files not supported
WARNING: you appear to work on 32-bit system... large files not supported
unique.names = TRUE

```

```

unique.rownames = TRUE
unique.colnames = TRUE
backingfilename = test_matrix_dimnames_T
cachesizeMb = 1
number of columns (variables) = 2
number of rows (observations) = 5
usedRowIndex: 1 2 3 4 5
usedColIndex: 1 2
Upper-left 2 columns and 5 rows:
      [,1]      [,2]
[1,] -8.5108500 -5.0336356
[2,] -7.3717060  9.2134371
[3,] -0.5829301  0.1687509
[4,]  6.7743611  5.1154494
[5,]  1.7932440 -5.1915603

> x <- databel_filtered_R("test_matrix_dimnames_T")

WARNING: you appear to work on 32-bit system... large files not supported

> t(testmatr)

      row_1      row_2
col_1 -8.5108500 -5.0336355
col_2 -7.3717062  9.2134366
col_3 -0.5829301  0.1687509
col_4  6.7743612  5.1154496
col_5  1.7932441 -5.1915602

> x

finalizing AbstractMatrix: 0x147f200
finalizing AbstractMatrix: 0x154b000
finalizing AbstractMatrix: 0x14abc00
unique.names = TRUE
unique.rownames = TRUE
unique.colnames = TRUE
backingfilename = test_matrix_dimnames_T
cachesizeMb = 1
number of columns (variables) = 2
number of rows (observations) = 5
usedRowIndex: 1 2 3 4 5
usedColIndex: 1 2
Upper-left 2 columns and 5 rows:
      [,1]      [,2]
[1,] -8.5108500 -5.0336356
[2,] -7.3717060  9.2134371

```

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[3,] -0.5829301  0.1687509
[4,]  6.7743611  5.1154494
[5,]  1.7932440 -5.1915603

> tmp <- as(x, "matrix")
> tmp

      row_1      row_2
col_1 -8.5108500 -5.0336356
col_2 -7.3717060  9.2134371
col_3 -0.5829301  0.1687509
col_4  6.7743611  5.1154494
col_5  1.7932440 -5.1915603

> abs(t(testmatr) - tmp) < 1e-06

      row_1 row_2
col_1  TRUE  TRUE
col_2  TRUE  TRUE
col_3  TRUE  TRUE
col_4  TRUE  TRUE
col_5  TRUE  TRUE

> unlink("*.fv?")
> unlink("test_matrix_*")

```