

Package ‘RGF’

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Type Package

Title Regularized Greedy Forest

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BugReports <https://github.com/RGF-team/rgf/issues>

URL <https://github.com/RGF-team/rgf/tree/master/R-package>

Description

Regularized Greedy Forest wrapper of the 'Regularized Greedy Forest' <<https://github.com/RGF-team/rgf/tree/master/python-package>> 'python' package, which also includes a Multi-core implementation (FastRGF) <<https://github.com/RGF-team/rgf/tree/master/FastRGF>>.

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SystemRequirements Python (2.7 or ≥ 3.4), rgf_python, scikit-learn ($\geq 0.18.0$), scipy, numpy. Detailed installation instructions for each operating system can be found in the README file.

Depends R($\geq 3.2.0$)

Imports reticulate, R6, Matrix

Suggests testthat, covr, knitr, rmarkdown

Encoding UTF-8

LazyData true

RoxygenNote 6.1.0

VignetteBuilder knitr

NeedsCompilation no

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FastRGF_Classifier *A Fast Regularized Greedy Forest classifier*

Description

A Fast Regularized Greedy Forest classifier

Usage

```
# init <- FastRGF_Classifier$new(n_estimators = 500, max_depth = 6,
#                               max_leaf = 50, tree_gain_ratio = 1.0,
#                               min_samples_leaf = 5, loss = "LS", l1 = 1.0,
#                               l2 = 1000.0, opt_algorithm = "rgf",
#                               learning_rate = 0.001, max_bin = NULL,
#                               min_child_weight = 5.0, data_l2 = 2.0,
#                               sparse_max_features = 80000,
#                               sparse_min_occurences = 5,
#                               calc_prob="sigmoid", n_jobs = 1,
#                               verbose = 0)
```

Arguments

x an R matrix (object) or a Python sparse matrix (object) of shape $c(n_samples, n_features)$. The training input samples. The sparse matrix should be a Python sparse matrix. The helper functions *mat_2scipy_sparse* and *TO_scipy_sparse* allow the user to convert an R dense or sparse matrix to a scipy sparse matrix.

y a vector of shape $c(n_samples)$. The target values (real numbers in regression).

n_estimators an integer. The number of trees in the forest (Original name: forest.ntrees.)

| | |
|------------------------|---|
| max_depth | an integer. Maximum tree depth (Original name: dtree.max_level.) |
| max_leaf | an integer. Maximum number of leaf nodes in best-first search (Original name: dtree.max_nodes.) |
| tree_gain_ratio | a float. New tree is created when leaf-nodes gain < this value * estimated gain of creating new tree (Original name: dtree.new_tree_gain_ratio.) |
| min_samples_leaf | an integer or float. Minimum number of training data points in each leaf node. If an integer, then consider min_samples_leaf as the minimum number. If a float, then min_samples_leaf is a percentage and $\text{ceil}(\text{min_samples_leaf} * \text{n_samples})$ are the minimum number of samples for each node (Original name: dtree.min_sample.) |
| loss | a character string. One of "LS" (Least squares loss), "MODLS" (Modified least squares loss) or "LOGISTIC" (Logistic loss) (Original name: dtree.loss.) |
| l1 | a float. Used to control the degree of L1 regularization (Original name: dtree.lamL1.) |
| l2 | a float. Used to control the degree of L2 regularization (Original name: dtree.lamL2.) |
| opt_algorithm | a character string. Either "rgf" or "epsilon-greedy". Optimization method for training forest (Original name: forest.opt.) |
| learning_rate | a float. Step size of epsilon-greedy boosting. Meant for being used with opt_algorithm = "epsilon-greedy" (Original name: forest.stepsize.) |
| max_bin | an integer or NULL. Maximum number of discretized values (bins). If NULL, 65000 is used for dense data and 200 for sparse data (Original name: discretize.(sparse/dense).max_buckets.) |
| min_child_weight | a float. Minimum sum of data weights for each discretized value (bin) (Original name: discretize.(sparse/dense).min_bucket_weights.) |
| data_l2 | a float. Used to control the degree of L2 regularization for discretization (Original name: discretize.(sparse/dense).lamL2.) |
| sparse_max_features | an integer. Maximum number of selected features. Meant for being used with sparse data (Original name: discretize.sparse.max_features.) |
| sparse_min_occurrences | an integer. Minimum number of occurrences for a feature to be selected. Meant for being used with sparse data (Original name: discretize.sparse.min_occurrences.) |
| calc_prob | a character string. Either "sigmoid" or "softmax". Method of probability calculation |
| n_jobs | an integer. The number of jobs to run in parallel for both fit and predict. If -1, all CPUs are used. If -2, all CPUs but one are used. If < -1, (n_cpus + 1 + n_jobs) are used (Original name: set.nthreads.) |
| verbose | an integer. Controls the verbosity of the tree building process (Original name: set.verbose.) |

Format

An object of class R6ClassGenerator of length 24.

Details

the *fit* function builds a classifier from the training set (x, y).

the *predict* function predicts the class for x.

the *predict_proba* function predicts class probabilities for x.

the *cleanup* function removes tempfiles used by this model. See the issue <https://github.com/RGF-team/rgf/issues/75>, which explains in which cases the *cleanup* function applies.

the *get_params* function returns the parameters of the model.

the *score* function returns the mean accuracy on the given test data and labels.

Methods

```
FastRGF_Classifier$new(n_estimators = 500, max_depth = 6, max_leaf = 50, tree_gain_ratio = 1.0, min
```

```
-----
fit(x, y, sample_weight = NULL)
-----
```

```
predict(x)
-----
```

```
predict_proba(x)
-----
```

```
cleanup()
-----
```

```
get_params(deep = TRUE)
-----
```

```
score(x, y, sample_weight = NULL)
-----
```

References

<https://github.com/RGF-team/rgf/tree/master/python-package>, Tong Zhang, *FastRGF: Multi-core Implementation of Regularized Greedy Forest* (<https://github.com/RGF-team/rgf/tree/master/FastRGF>)

Examples

```
if (reticulate::py_available() && reticulate::py_module_available("rgf.sklearn")) {
  library(RGF)
  set.seed(1)
  x = matrix(runif(100000), nrow = 100, ncol = 1000)
  y = sample(1:2, 100, replace = TRUE)
```

```

fast_RGF_class = FastRGF_Classifier$new(max_leaf = 50)

fast_RGF_class$fit(x, y)

preds = fast_RGF_class$predict_proba(x)
}

```

FastRGF_Regressor *A Fast Regularized Greedy Forest regressor*

Description

A Fast Regularized Greedy Forest regressor

Usage

```

# init <- FastRGF_Regressor$new(n_estimators = 500, max_depth = 6,
#                               max_leaf = 50, tree_gain_ratio = 1.0,
#                               min_samples_leaf = 5, l1 = 1.0,
#                               l2 = 1000.0, opt_algorithm = "rgf",
#                               learning_rate = 0.001, max_bin = NULL,
#                               min_child_weight = 5.0, data_l2 = 2.0,
#                               sparse_max_features = 80000,
#                               sparse_min_occurrences = 5,
#                               n_jobs = 1, verbose = 0)

```

Arguments

| | |
|-------------------------------|--|
| <code>x</code> | an R matrix (object) or a Python sparse matrix (object) of shape $c(n_samples, n_features)$. The training input samples. The sparse matrix should be a Python sparse matrix. The helper functions <i>mat_2scipy_sparse</i> and <i>TO_scipy_sparse</i> allow the user to convert an R dense or sparse matrix to a scipy sparse matrix. |
| <code>y</code> | a vector of shape $c(n_samples)$. The target values (real numbers in regression). |
| <code>n_estimators</code> | an integer. The number of trees in the forest (Original name: forest.ntrees.) |
| <code>max_depth</code> | an integer. Maximum tree depth (Original name: dtree.max_level.) |
| <code>max_leaf</code> | an integer. Maximum number of leaf nodes in best-first search (Original name: dtree.max_nodes.) |
| <code>tree_gain_ratio</code> | a float. New tree is created when leaf-nodes gain $<$ this value * estimated gain of creating new tree (Original name: dtree.new_tree_gain_ratio.) |
| <code>min_samples_leaf</code> | an integer or float. Minimum number of training data points in each leaf node. If an integer, then consider <code>min_samples_leaf</code> as the minimum number. If a float, then <code>min_samples_leaf</code> is a percentage and $\text{ceil}(\text{min_samples_leaf} * n_samples)$ are the minimum number of samples for each node (Original name: dtree.min_sample.) |
| <code>l1</code> | a float. Used to control the degree of L1 regularization (Original name: dtree.lamL1.) |

| | |
|-------------------------------------|---|
| <code>l2</code> | a float. Used to control the degree of L2 regularization (Original name: <code>dtree.lamL2</code> .) |
| <code>opt_algorithm</code> | a character string. Either <code>"rgf"</code> or <code>"epsilon-greedy"</code> . Optimization method for training forest (Original name: <code>forest.opt</code> .) |
| <code>learning_rate</code> | a float. Step size of epsilon-greedy boosting. Meant for being used with <code>opt_algorithm = "epsilon-greedy"</code> (Original name: <code>forest.stepsize</code> .) |
| <code>max_bin</code> | an integer or NULL. Maximum number of discretized values (bins). If NULL, 65000 is used for dense data and 200 for sparse data (Original name: <code>discretize.(sparse/dense).max_buckets</code> .) |
| <code>min_child_weight</code> | a float. Minimum sum of data weights for each discretized value (bin) (Original name: <code>discretize.(sparse/dense).min_bucket_weights</code> .) |
| <code>data_l2</code> | a float. Used to control the degree of L2 regularization for discretization (Original name: <code>discretize.(sparse/dense).lamL2</code> .) |
| <code>sparse_max_features</code> | an integer. Maximum number of selected features. Meant for being used with sparse data (Original name: <code>discretize.sparse.max_features</code> .) |
| <code>sparse_min_occurrences</code> | an integer. Minimum number of occurrences for a feature to be selected. Meant for being used with sparse data (Original name: <code>discretize.sparse.min_occurrences</code> .) |
| <code>n_jobs</code> | an integer. The number of jobs to run in parallel for both fit and predict. If -1, all CPUs are used. If -2, all CPUs but one are used. If < -1, (<code>n_cpus + 1 + n_jobs</code>) are used (Original name: <code>set.nthreads</code> .) |
| <code>verbose</code> | an integer. Controls the verbosity of the tree building process (Original name: <code>set.verbose</code> .) |

Format

An object of class `R6ClassGenerator` of length 24.

Details

the *fit* function builds a regressor from the training set (x, y).

the *predict* function predicts the regression target for x .

the *cleanup* function removes tempfiles used by this model. See the issue <https://github.com/RGF-team/rgf/issues/75>, which explains in which cases the *cleanup* function applies.

the *get_params* function returns the parameters of the model.

the *score* function returns the coefficient of determination (R^2) for the predictions.

Methods

```
FastRGF_Regressor$new(n_estimators = 500, max_depth = 6, max_leaf = 50, tree_gain_ratio = 1.0, min_
```

```
-----
fit(x, y, sample_weight = NULL)
-----
```

```

predict(x)
-----
cleanup()
-----
get_params(deep = TRUE)
-----
score(x, y, sample_weight = NULL)
-----

```

References

<https://github.com/RGF-team/rgf/tree/master/python-package>, Tong Zhang, *FastRGF: Multi-core Implementation of Regularized Greedy Forest* (<https://github.com/RGF-team/rgf/tree/master/FastRGF>)

Examples

```

if (reticulate::py_available() && reticulate::py_module_available("rgf.sklearn")) {
  library(RGF)

  set.seed(1)
  x = matrix(runif(100000), nrow = 100, ncol = 1000)

  y = runif(100)

  fast_RGF_regr = FastRGF_Regressor$new(max_leaf = 50)

  fast_RGF_regr$fit(x, y)

  preds = fast_RGF_regr$predict(x)
}

```

mat_2scipy_sparse *conversion of an R matrix to a scipy sparse matrix*

Description

conversion of an R matrix to a scipy sparse matrix

Usage

```
mat_2scipy_sparse(x, format = "sparse_row_matrix")
```

Arguments

x a data matrix
format a character string. Either *"sparse_row_matrix"* or *"sparse_column_matrix"*

Arguments

| | |
|------------------|--|
| x | an R matrix (object) or a Python sparse matrix (object) of shape $c(n_samples, n_features)$. The training input samples. The sparse matrix should be a Python sparse matrix. The helper functions <i>mat_2scipy_sparse</i> and <i>TO_scipy_sparse</i> allow the user to convert an R dense or sparse matrix to a scipy sparse matrix. |
| y | a vector of shape $c(n_samples)$. The target values (class labels in classification). |
| sample_weight | a vector of shape $c(n_samples)$ or NULL. Individual weights for each sample. |
| max_leaf | an integer. Training will be terminated when the number of leaf nodes in the forest reaches this value. |
| test_interval | an integer. Test interval in terms of the number of leaf nodes. |
| algorithm | a character string specifying the <i>Regularization algorithm</i> . One of "RGF" (RGF with L2 regularization on leaf-only models), "RGF_Opt" (RGF with min-penalty regularization) or "RGF_Sib" (RGF with min-penalty regularization with the sum-to-zero sibling constraints). |
| loss | a character string specifying the <i>Loss function</i> . One of "LS" (Square loss), "Expo" (Exponential loss) or "Log" (Logistic loss). |
| reg_depth | a float. Must be no smaller than 1.0. Meant for being used with the algorithm <i>RGF Opt</i> or <i>RGF Sib</i> . A larger value penalizes deeper nodes more severely. |
| l2 | a float. Used to control the degree of L2 regularization. |
| sl2 | a float or NULL. Override L2 regularization parameter l2 for the process of growing the forest. That is, if specified, the weight correction process uses l2 and the forest growing process uses sl2. If NULL, no override takes place and l2 is used throughout training. |
| normalize | a boolean. If True, training targets are normalized so that the average becomes zero. |
| min_samples_leaf | an integer or a float. Minimum number of training data points in each leaf node. If an integer, then consider <i>min_samples_leaf</i> as the minimum number. If a float, then <i>min_samples_leaf</i> is a percentage and $\text{ceil}(\text{min_samples_leaf} * n_samples)$ are the minimum number of samples for each node. |
| n_iter | an integer or NULL. The number of iterations of coordinate descent to optimize weights. If NULL, 10 is used for loss = "LS" and 5 for loss = "Expo" or "Log". |
| n_tree_search | an integer. The number of trees to be searched for the nodes to split. The most recently grown trees are searched first. |
| opt_interval | an integer. Weight optimization interval in terms of the number of leaf nodes. For example, by default, weight optimization is performed every time approximately 100 leaf nodes are newly added to the forest. |
| learning_rate | a float. Step size of Newton updates used in coordinate descent to optimize weights. |
| calc_prob | a character string. One of "sigmoid" or "softmax". Method of probability calculation. |

| | |
|----------------------------|--|
| <code>n_jobs</code> | an integer. The number of jobs (threads) to use for the computation. The substantial number of the jobs depends on <code>classes_</code> (The number of classes when <i>fit</i> is performed). If <code>classes_ = 2</code> , the substantial max number of the jobs is one. If <code>classes_ > 2</code> , the substantial max number of the jobs is the same as <code>classes_</code> . If <code>n_jobs = 1</code> , no parallel computing code is used at all regardless of <code>classes_</code> . If <code>n_jobs = -1</code> and <code>classes_ >=</code> number of CPU, all CPUs are used. For <code>n_jobs = -2</code> , all CPUs but one are used. For <code>n_jobs</code> below <code>-1</code> , <code>(n_cpus + 1 + n_jobs)</code> are used. |
| <code>memory_policy</code> | a character string. One of <i>"conservative"</i> (it uses less memory at the expense of longer runtime. Try only when with default value it uses too much memory) or <i>"generous"</i> (it runs faster using more memory by keeping the sorted orders of the features on memory for reuse). Memory using policy. |
| <code>verbose</code> | an integer. Controls the verbosity of the tree building process. |
| <code>init_model</code> | either NULL or a character string, optional (default=NULL). Filename of a previously saved model from which training should do warm-start. If model has been saved into multiple files, do not include numerical suffixes in the filename. <i>NOTE</i> : Make sure you haven't forgotten to increase the value of the <code>max_leaf</code> parameter regarding to the specified warm-start model because warm-start model trees are counted in the overall number of trees. |
| <code>filename</code> | a character string specifying a valid path to a file where the fitted model should be saved |

Format

An object of class `R6ClassGenerator` of length 24.

Details

the *fit* function builds a classifier from the training set (x, y) .

the *predict* function predicts the class for x .

the *predict_proba* function predicts class probabilities for x .

the *cleanup* function removes tempfiles used by this model. See the issue <https://github.com/RGF-team/rgf/issues/75>, which explains in which cases the *cleanup* function applies.

the *get_params* function returns the parameters of the model.

the *score* function returns the mean accuracy on the given test data and labels.

the *feature_importances* function returns the feature importances for the data.

the *dump_model* function currently prints information about the fitted model in the console

the *save_model* function saves a model to a file from which training can do warm-start in the future.

Methods

```
RGF_Classifier$new(max_leaf = 1000, test_interval = 100, algorithm = "RGF", loss = "Log", reg_depth
```

```
-----
```

```
fit(x, y, sample_weight = NULL)
```

```

-----
predict(x)
-----
predict_proba(x)
-----
cleanup()
-----
get_params(deep = TRUE)
-----
score(x, y, sample_weight = NULL)
-----
feature_importances()
-----
dump_model()
-----
save_model(filename)
-----

```

References

*<https://github.com/RGF-team/rgf/tree/master/python-package>, Rie Johnson and Tong Zhang, *Learning Nonlinear Functions Using Regularized Greedy Forest**

Examples

```

if (reticulate::py_available() && reticulate::py_module_available("rgf.sklearn")) {
  library(RGF)

  set.seed(1)
  x = matrix(runif(1000), nrow = 100, ncol = 10)

  y = sample(1:2, 100, replace = TRUE)

  RGF_class = RGF_Classifier$new(max_leaf = 50)

  RGF_class$fit(x, y)

  preds = RGF_class$predict_proba(x)
}

```


Arguments

| | |
|------------------|--|
| x | an R matrix (object) or a Python sparse matrix (object) of shape $c(n_samples, n_features)$. The training input samples. The sparse matrix should be a Python sparse matrix. The helper functions <i>mat_2scipy_sparse</i> and <i>TO_scipy_sparse</i> allow the user to convert an R dense or sparse matrix to a scipy sparse matrix. |
| y | a vector of shape $c(n_samples)$. The target values (real numbers in regression). |
| sample_weight | a vector of shape $c(n_samples)$ or NULL. Individual weights for each sample. |
| max_leaf | an integer. Training will be terminated when the number of leaf nodes in the forest reaches this value. |
| test_interval | an integer. Test interval in terms of the number of leaf nodes. |
| algorithm | a character string specifying the <i>Regularization algorithm</i> . One of "RGF" (RGF with L2 regularization on leaf-only models), "RGF_Opt" (RGF with min-penalty regularization) or "RGF_Sib" (RGF with min-penalty regularization with the sum-to-zero sibling constraints). |
| loss | a character string specifying the <i>Loss function</i> . One of "LS" (Square loss), "Expo" (Exponential loss) or "Log" (Logistic loss). |
| reg_depth | a float. Must be no smaller than 1.0. Meant for being used with the algorithm <i>RGF Opt</i> or <i>RGF Sib</i> . A larger value penalizes deeper nodes more severely. |
| l2 | a float. Used to control the degree of L2 regularization. |
| sl2 | a float or NULL. Override L2 regularization parameter l2 for the process of growing the forest. That is, if specified, the weight correction process uses l2 and the forest growing process uses sl2. If NULL, no override takes place and l2 is used throughout training. |
| normalize | a boolean. If True, training targets are normalized so that the average becomes zero. |
| min_samples_leaf | an integer or a float. Minimum number of training data points in each leaf node. If an integer, then consider <i>min_samples_leaf</i> as the minimum number. If a float, then <i>min_samples_leaf</i> is a percentage and $\text{ceil}(\text{min_samples_leaf} * n_samples)$ are the minimum number of samples for each node. |
| n_iter | an integer or NULL. The number of iterations of coordinate descent to optimize weights. If NULL, 10 is used for loss = "LS" and 5 for loss = "Expo" or "Log". |
| n_tree_search | an integer. The number of trees to be searched for the nodes to split. The most recently grown trees are searched first. |
| opt_interval | an integer. Weight optimization interval in terms of the number of leaf nodes. For example, by default, weight optimization is performed every time approximately 100 leaf nodes are newly added to the forest. |
| learning_rate | a float. Step size of Newton updates used in coordinate descent to optimize weights. |
| memory_policy | a character string. One of "conservative" (it uses less memory at the expense of longer runtime. Try only when with default value it uses too much memory) or "generous" (it runs faster using more memory by keeping the sorted orders of the features on memory for reuse). Memory using policy. |

| | |
|------------|---|
| verbose | an integer. Controls the verbosity of the tree building process. |
| init_model | either NULL or a character string, optional (default=NULL). Filename of a previously saved model from which training should do warm-start. If model has been saved into multiple files, do not include numerical suffixes in the filename. <i>NOTE</i> : Make sure you haven't forgotten to increase the value of the max_leaf parameter regarding to the specified warm-start model because warm-start model trees are counted in the overall number of trees. |
| filename | a character string specifying a valid path to a file where the fitted model should be saved |

Format

An object of class R6ClassGenerator of length 24.

Details

the *fit* function builds a regressor from the training set (x, y).

the *predict* function predicts the regression target for x.

the *cleanup* function removes tempfiles used by this model. See the issue <https://github.com/RGF-team/rgf/issues/75>, which explains in which cases the *cleanup* function applies.

the *get_params* function returns the parameters of the model.

the *score* function returns the coefficient of determination (R^2) for the predictions.

the *feature_importances* function returns the feature importances for the data.

the *dump_model* function currently prints information about the fitted model in the console

the *save_model* function saves a model to a file from which training can do warm-start in the future.

Methods

```
RGF_Regressor$new(max_leaf = 500, test_interval = 100, algorithm = "RGF", loss = "LS", reg_depth =
```

```
-----
fit(x, y, sample_weight = NULL)
-----
predict(x)
-----
cleanup()
-----
get_params(deep = TRUE)
-----
score(x, y, sample_weight = NULL)
-----
feature_importances()
```

```

-----
dump_model()
-----
save_model(filename)
-----

```

References

<https://github.com/RGF-team/rgf/tree/master/python-package>, Rie Johnson and Tong Zhang, *Learning Nonlinear Functions Using Regularized Greedy Forest*

Examples

```

if (reticulate::py_available() && reticulate::py_module_available("rgf.sklearn")) {
  library(RGF)

  set.seed(1)
  x = matrix(runif(1000), nrow = 100, ncol = 10)

  y = runif(100)

  RGF_regr = RGF_Regressor$new(max_leaf = 50)

  RGF_regr$fit(x, y)

  preds = RGF_regr$predict(x)
}

```

| | |
|-----------------|--|
| TO_scipy_sparse | <i>conversion of an R sparse matrix to a scipy sparse matrix</i> |
|-----------------|--|

Description

conversion of an R sparse matrix to a scipy sparse matrix

Usage

```
TO_scipy_sparse(R_sparse_matrix)
```

Arguments

R_sparse_matrix
 an R sparse matrix. Acceptable input objects are either a *dgCMatrix* or a *dgRMatrix*.

Details

This function allows the user to convert either an R *dgCMatrix* or a *dgRMatrix* to a scipy sparse matrix (*scipy.sparse.csc_matrix* or *scipy.sparse.csr_matrix*). This is useful because the *RGF* package accepts besides an R dense matrix also python sparse matrices as input.

The *dgCMatrix* class is a class of sparse numeric matrices in the compressed, sparse, *column-oriented format*. The *dgRMatrix* class is a class of sparse numeric matrices in the compressed, sparse, *row-oriented format*.

References

<https://stat.ethz.ch/R-manual/R-devel/library/Matrix/html/dgCMatrix-class.html>, <https://stat.ethz.ch/R-manual/R-devel/library/Matrix/html/dgRMatrix-class.html>, <https://docs.scipy.org/doc/scipy/reference/generated/scipy.sparse>

Examples

```
if (reticulate::py_available() && reticulate::py_module_available("scipy")) {
  if (Sys.info()["sysname"] != 'Darwin') {
    library(RGF)

    # 'dgCMatrix' sparse matrix
    #-----

    data = c(1, 0, 2, 0, 0, 3, 4, 5, 6)

    dgcM = Matrix::Matrix(
      data = data
      , nrow = 3
      , ncol = 3
      , byrow = TRUE
      , sparse = TRUE
    )

    print(dim(dgcM))

    res = TO_scipy_sparse(dgcM)

    print(res$shape)

    # 'dgRMatrix' sparse matrix
    #-----

    dgrM = as(dgcM, "RsparseMatrix")

    print(dim(dgrM))

    res_dgr = TO_scipy_sparse(dgrM)
  }
}
```



```
        print(res_dgr$shape)
    }
}
```

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