

Manual of Frovedis Python API

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Chapter 1

Introduction

This manual contains Python API documentation. If you are new to Frovedis, please read the `tutorial_python` first.

Currently we only provide part of the API documentation. We are still updating the contents.

- Machine Learning
 - Agglomerative Clustering
 - Matrix Factorization using ALS
 - Bernoulli Naive Bayes
 - DBSCAN
 - DecisionTreeClassifier
 - DecisionTreeRegressor
 - FactorizationMachineClassifier
 - FactorizationMachineRegressor
 - FPGrowth
 - GaussianMixture
 - GradientBoostingClassifier
 - GradientBoostingRegressor
 - KMeans Clustering
 - KNeighborsClassifier
 - KNeighborsRegressor
 - Lasso Regression
 - Latent Dirichlet Allocation
 - Linear Regression
 - LinearSVC
 - LinearSVR
 - Logistic Regression
 - Multinomial Naive Bayes
 - NearestNeighbors
 - Principal Component Analysis
 - RandomForestClassifier
 - RandomForestRegressor
 - Ridge Regression
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 - SGDRegressor
 - Spectral Clustering
 - Spectral Embedding
 - StandardScaler
 - SVC

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Chapter 2

Agglomerative Clustering

2.1 NAME

Agglomerative Clustering - The most common type of hierarchical clustering used to group objects in clusters based on their similarities.

2.2 SYNOPSIS

```
class frovedis.mllib.cluster.AgglomerativeClustering(n_clusters=2, affinity='euclidean',
                                                    memory=None, connectivity=None,
                                                    compute_full_tree='auto',
                                                    linkage='average',
                                                    distance_threshold=None,
                                                    compute_distances=False,
                                                    verbose=0)
```

2.2.1 Public Member Functions

```
fit(X, y = None)
fit_predict(X, y = None)
reassign(ncluster = None)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

2.3 DESCRIPTION

Clustering is a machine learning technique that involves the grouping of data points. Hierarchical clustering is a general family of clustering algorithms that build nested clusters by merging or splitting them successively.

The Agglomerative Clustering object performs a hierarchical clustering using a bottom-up approach. Each observation starts in its own cluster, and clusters are successively merged together.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn Agglomerative Clustering interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Agglomerative Clustering on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

2.3.1 Detailed Description

2.3.1.1 1. AgglomerativeClustering()

Parameters

n_clusters: An integer parameter specifying the number of clusters. The number of clusters should be greater than 0 and less than *n_samples*. (Default: 2)

affinity: An unused parameter. (Default: 'euclidean')

memory: An unused parameter. (Default: None)

connectivity: An unused parameter. (Default: None)

compute_full_tree: An unused parameter. (Default: 'auto')

linkage: A string parameter used to specify linkage criterion. It determines which distance to use between sets of observation. The algorithm will merge the pairs of clusters that minimize this criterion.

- 'average': uses the average of the distances of each observation of the two sets.
- 'complete': linkage uses the maximum distances between all observations of the two sets.
- 'single': uses the minimum of the distances between all observations of the two sets.

Only 'average', 'complete' and 'single' are supported. (Default: 'average')

distance_threshold: A float or double(float64) type parameter, is the linkage distance threshold above which the clusters will not be merged. It must be zero or positive value. (Default: None)

When it is None (not specified explicitly), it will be set as 0.0.

compute_distances: Unlike Scikit-learn, it is always True for frovedis. Hence, this parameter is left unused. (Default: False)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Attributes

n_clusters: A positive integer value specifying the number of clusters found by the algorithm.

labels: A python ndarray of int64 type values and has shape (*n_clusters*,). It contains cluster labels for each point.

children: A python ndarray of int64 type values and has shape (*n_samples* - 1, 2). It contains the children of each non-leaf node.

distances: A python ndarray of float or double(float64) values and has shape (*n_samples* - 1,). It specifies the distances between nodes in the corresponding place in "children".

n_connected_components: An integer value used to provide the estimated number of connected components in the graph.

Purpose

It initializes an Agglomerative Clustering object with the given parameters.

The parameters: “affinity”, “memory”, “connectivity”, “compute_full_tree” and “compute_distances” are simply kept in to make the interface uniform to the Scikit-learn Agglomerative Clustering module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

2.3.1.2 2. fit(X, y = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn as well.

Purpose

It clusters the given data points (X) into a predefined number of clusters.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/sample_data.txt")

# fitting input matrix on AgglomerativeClustering object
from frovedis.mllib.cluster import AgglomerativeClustering
acm = AgglomerativeClustering(n_clusters = 2).fit(mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/sample_data.txt")

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# Agglomerative Clustering with pre-constructed frovedis-like inputs
from frovedis.mllib.cluster import AgglomerativeClustering
acm = AgglomerativeClustering(n_clusters = 2).fit(rmat)
```

Return Value

It simply returns “self” reference.

2.3.1.3 3. fit_predict(X, y = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn as well.

Purpose

It clusters the given data points (X) into a predefined number of clusters. In addition to fitting, it returns the cluster labels for each sample in the training set.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/sample_data.txt")

# fitting input matrix on AgglomerativeClustering object
from frovedis.mllib.cluster import AgglomerativeClustering
acm = AgglomerativeClustering(n_clusters = 2)
print(acm.fit_predict(mat))
```

Output

```
[1 1 0 0 0]
```

Like in fit(), frovedis-like input can be used to speed-up the training at server side.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/sample_data.txt")

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import AgglomerativeClustering
acm = AgglomerativeClustering(n_clusters = 2)
print(acm.fit_predict(rmat))
```

Output

```
[1 1 0 0 0]
```

Return Value

It returns a numpy array of int64 type values containing the cluster labels. It has a shape (**n_samples**,).

2.3.1.4 4. reassign(ncluster = None)

Parameters

nclusters: An integer parameter specifying the number of clusters to be reassigned for the fitted data without computing the tree again. The number of clusters should be greater than 0 and less than n_samples. (Default: None)

When it is None (not specified explicitly), it simply returns the same cluster labels of already fitted clustering model. In this case, 'ncluster' becomes the 'n_cluster' value used during fit().

Purpose

It accepts the number of clusters (nclusters) in order to make prediction with different "nclusters" on same model at frovedis server.

For example,

```
# On the same AgglomerativeClustering object, predicting labels with new nclusters
print(acm.reassign(nclusters = 3))
```

Output

```
[0 0 1 1 2]
```

Return Value

It returns a numpy array of int64 type values containing the cluster labels. It has a shape **(n_samples,)**.

2.3.1.5 5. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: A python ndarray or an instance of FrovedisVector containing the true labels for X. It has shape **(n_samples, 1)**.

sample_weight: An unused parameter whose default value is None. It is simply ignored in frovedis implementation.

Purpose

It uses homogeneity score on given test data and labels i.e homogeneity score of self.predict(X, y) wrt. y.

For example,

```
acm.score(train_mat, [0, 0, 1, 1 1])
```

Output

```
1.0
```

Return Value

It returns a homogeneity score of float type.

2.3.1.6 6. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by AgglomerativeClustering. It is used to get parameters and their values of AgglomerativeClustering class.

For example,

```
print(acm.get_params())
```

Output

```
{'affinity': 'euclidean', 'compute_distances': True, 'compute_full_tree': 'auto',
'connectivity': None, 'distance_threshold': None, 'linkage': 'average', 'memory': None,
'n_clusters': 3, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

2.3.1.7 7. `set_params(**params)`

Parameters

*****params***: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the `BaseEstimator` class inherited by `AgglomerativeClustering`, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(acm.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
acm.set_params(n_clusters = 4)
print("get parameters after setting:")
print(acm.get_params())
```

Output

```
get parameters before setting:
{'affinity': 'euclidean', 'compute_distances': True, 'compute_full_tree': 'auto',
 'connectivity': None, 'distance_threshold': None, 'linkage': 'average', 'memory': None,
 'n_clusters': 3, 'verbose': 0}
get parameters after setting:
{'affinity': 'euclidean', 'compute_distances': True, 'compute_full_tree': 'auto',
 'connectivity': None, 'distance_threshold': None, 'linkage': 'average', 'memory': None,
 'n_clusters': 4, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

2.3.1.8 8. `load(fname, dtype = None)`

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads an agglomerative clustering model stored previously from the specified file (having little-endian binary data).

For example,

```
acm.load("./out/MyAcmClusteringModel", dtype = np.float64)
```

Return Value

It simply returns “self” reference.

2.3.1.9 9. `save(fname)`

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the agglomerative clustering model
acm.save("./out/MyAcmClusteringModel")
```

This will save the agglomerative clustering model on the path “/out/MyAcmClusteringModel”.

It would raise exception if the directory already exists with same name.

The ‘MyAcmClusteringModel’ directory has

MyAcmClusteringModel

```
|—— metadata
|—— model
```

The metadata file contains the number of clusters, number of samples, model kind, input datatype used for trained model.

Here, the **model** directory contains information about dendrogram.

Return Value

It returns nothing.

2.3.1.10 10. debug_print()**Purpose**

It shows the target model information(dendrogram) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
acm.debug_print()
```

Output

```
--- dendrogram ---
      X      Y      distance      size
5:      2      3      0.173205      2
6:      0      1      0.173205      2
7:      4      5      0.259808      3
8:      6      7      15.5019 5
```

This output will be visible on server side. It displays the dendrogram on the trained model which is currently present on the server. Using the dendrogram, the desired number of clusters may be found.

No such output will be visible on client side.

Return Value

It returns nothing.

2.3.1.11 11. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
acm.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

2.3.1.12 12. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, `reassign()` is used before training the model, then it can prompt the user to train the clustering model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

2.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **DBSCAN in Frovedis**
- **KMeans in Frovedis**
- **Spectral Clustering in Frovedis**

Chapter 3

Matrix Factorization using ALS

3.1 NAME

Matrix Factorization using Alternating Least Square (ALS) - is a matrix factorization algorithm commonly used for recommender systems.

3.2 SYNOPSIS

```
class frovedis.mllib.recommendation.ALS(rank=None, max_iter=100, alpha=0.01,
                                         reg_param=0.01, similarity_factor=0.1,
                                         seed=0, verbose=0)
```

3.2.1 Public Member Functions

```
fit(X)
predict(ids)
recommend_users(pid, k)
recommend_products(uid, k)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

3.3 DESCRIPTION

Collaborative filtering is commonly used for recommender systems. These techniques aim to fill in the missing entries of a user-item association matrix. Frovedis currently supports model-based collaborative filtering, in which users and products are described by a small set of latent factors that can be used to predict missing entries.

Frovedis uses the alternating least squares(ALS) algorithm to learn these latent factors. The algorithm is based on a paper “Collaborative Filtering for Implicit Feedback Datasets” by Hu, et al.

This module provides a client-server implementation, where the client application is a normal python program. Scikit-learn does not have any collaborative filtering recommender algorithms like ALS. In this implementation, python side recommender interfaces are provided, where a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ALS call is linked with the frovedis ALS call to get the job done at frovedis server.

Python side calls for ALS on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When recommendation-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

3.3.1 Detailed Description

3.3.1.1 1. ALS()

rank: A positive integer parameter containing the user given rank for the input matrix. (Default: None)
When rank is None (not specified explicitly), it will be the minimum(256, min(M,N)), where M is number of users and N is number of items in input data. It must be within the range of 0 to max(M, N).

max_iter: A positive integer specifying maximum iteration count. (Default: 100)

alpha: A positive double(float64) parameter containing the learning rate. (Default: 0.01)

reg_param: A positive double(float64) parameter, also called as the regularization parameter. (Default: 0.01)

similarity_factor: A double(float64) parameter, which helps to identify whether the algorithm will optimize the computation for similar user/item or not. If similarity percentage of user or item features is more than or equal to the given similarity_factor, the algorithm will optimize the computation for similar user/item. Otherwise, each user and item feature will be treated uniquely. Similarity factor must be in range of ≥ 0.0 to ≤ 1.0 . (Default: 0.1)

seed: An int64 parameter containing the seed value to initialize the model structures with random values. (Default: 0)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Purpose

It initializes an ALS object with the given parameters.

Return Value

It simply returns “self” reference.

3.3.1.2 2. fit(X)

Parameters

X: A scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix. It has shape (n_samples, n_features).

Purpose

It accepts the training sparse matrix (X) and trains a matrix factorization model on that at frovedis server.

It starts with initializing the model structures of the size MxF and NxF (where M is the number of users, N is the products in the given rating matrix and F is the given rank) with random values and keeps updating them until maximum iteration count is reached.

For example,

```
# creating csr matrix
import numpy as np
from scipy.sparse import csr_matrix
row = np.array([0, 0, 1, 2, 2, 3, 3, 3, 4, 4, 5, 6, 6, 7, 7, 7])
col = np.array([0, 4, 0, 2, 3, 0, 1, 6, 0, 4, 0, 2, 3, 0, 1, 6])
data = np.array([2.0, 9.0, 1.0, 4.0, 8.0, 2.0, 3.0, 8.9, 2.0,
                 9.0, 1.0, 4.0, 8.0, 2.0, 3.0, 8.9])
csr_matrix = csr_matrix((data, (row, col)), shape = (8, 7))

# fitting input matrix on ALS object
from frovedis.ml.lib.recommendation import ALS
als = ALS(rank = 4).fit(csr_matrix)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# Since "mat" is scipy csr sparse matrix, we have created FrovedisCRSMatrix.
from frovedis.matrix.crs import FrovedisCRSMatrix
crs_mat = FrovedisCRSMatrix(mat)

# ALS with pre-constructed frovedis-like inputs
from frovedis.ml.lib.recommendation import ALS
als = ALS(rank = 4).fit(crs_mat)
```

Return Value

It simply returns “self” reference.

3.3.1.3 3. predict(ids)

Parameters

ids: A python tuple or list object containing the pairs of user id and product id to predict.

Purpose

It accepts a list of pair of user ids and product ids (0-based Id) in order to make prediction for their ratings from the trained model at frovedis server.

For example,

```
# prints the predicted ratings for the given list of id pairs
als.predict([(1,1),(0,1),(2,3),(3,1)])
```

Output:

```
[ 0.00224735  0.00152505  0.99515575  0.99588757]
```

Return Value

It returns a numpy array containing the predicted ratings, of float or double (float64) type depending upon the input type.

3.3.1.4 4. recommend_users(pid, k)

Parameters

pid: An integer parameter specifying the product ID (0-based Id) for which to recommend users.

k: An integer parameter specifying the number of users to be recommended.

Purpose

It recommends the best “k” users with highest rating confidence in sorted order for the given product.

If $k > \text{number of rows}$ (number of users in the given matrix when training the model), then it resets the k as “number of rows in the given matrix”. This is done in order to recommend all the users with rating confidence values in descending order.

For example,

```
# recommend 2 users for second product
als.recommend_users(1,2)
```

Output:

```
('uids:', array([7, 3], dtype=int32))
('scores:', array([ 0.99588757,  0.99588757]))
```

Return Value

It returns a python list containing the pairs of recommended users and their corresponding rating confidence values (double (float64)) in descending order.

3.3.1.5 5. recommend_products(uid, k)

Parameters

uid: An integer parameter specifying the user ID (0-based Id) for which to recommend products.

k: An integer parameter specifying the number of products to be recommended.

Purpose

It recommends the best “k” products with highest rating confidence in sorted order for the given user.

If $k > \text{number of columns}$ (number of products in the given matrix when training the model), then it resets the k as “number of columns in the given matrix”. This is done in order to recommend all the products with rating confidence values in descending order.

For example,

```
# recommend 2 products for second user
print als.recommend_products(1,2)
```

Output:

```
(' recommend_product pids:', array([0, 4], dtype=int32))
('scores:', array([ 0.99494576,  0.0030741 ]))[(0, 0.9949457612078868), (4, 0.0030740973144160397)]
```

Return Value

It returns a python list containing the pairs of recommended products and their corresponding rating confidence values (double (float64)) in descending order.

3.3.1.6 6. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information(metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
als.save("./out/MyALSModel")
```

The MyALSModel contains below directory structure:

MyALSModel

```
|— metadata
|— model
    |— X
    |— Y
```

‘metadata’ represents the detail about model_kind and datatype of training vector. Here, the **model** directory contains information about user and product features.

If the directory already exists with the same name then it will raise an exception.

Return Value

It returns nothing.

3.3.1.7 7. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the same model
als.load("./out/MyALSModel")
```

Return Value

It simply returns “self” instance.

3.3.1.8 8. debug__print()

Purpose

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
als.debug_print()
```

Output:

```
===== Matrix::X =====
0.829524 -0.84477 -0.152624 0.569863
0.829528 -0.844775 -0.152625 0.569867
0.829921 -0.845174 -0.152697 0.570136
===== Matrix::Y =====
0.473117 -0.481813 -0.087049 0.32502
0.473117 -0.481813 -0.087049 0.32502
0.473117 -0.481813 -0.087049 0.32502
```

This output will be visible on server side. It will print the matrix and labels of training data.

No such output will be visible on client side.

Return Value

It returns nothing.

3.3.1.9 9. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
als.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

3.3.1.10 10. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

3.4 SEE ALSO

- Introduction to FrovedisCRSMatrix

Chapter 4

Bernoulli Naive Bayes

4.1 NAME

BernoulliNB - One of the variations of Naive Bayes algorithm. It is a classification algorithm to predict only binary output.

4.2 SYNOPSIS

```
frovedis.mllib.naive_bayes.BernoulliNB(alpha=1.0, fit_prior=True,  
                                         class_prior=None, binarize=0.0,  
                                         verbose=0)
```

4.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)  
predict(X)  
predict_proba(X)  
score(X, y, sample_weight = None)  
get_params(deep = True)  
set_params(**params)  
debug_print()  
load(fname, dtype = None)  
save(fname)  
release()  
is_fitted()
```

4.3 DESCRIPTION

Naive Bayes classifier for bernoulli models.

The Bernoulli Naive Bayes classifier is suitable for classification with binary/boolean features.

This model is popular for document classification tasks, where binary term occurrence features (i.e a word occurs in a document or not) are used rather than finding the frequency of a word in document.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn BernoulliNB interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for BernoulliNB on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

4.3.1 Detailed Description

4.3.1.1 1. BernoulliNB()

Parameters

alpha: A positive double (float64) smoothing parameter (0 for no smoothing). It must be greater than or equal to 1. (Default: 1.0)

fit_prior: A boolean parameter specifying whether to learn class prior probabilities or not. If False, a uniform prior will be used. (Default: True)

class_prior: A numpy ndarray of double (float64) type values and must be of shape (***n_classes***,). It gives prior probabilities of the classes. (Default: None)

When it is None (not specified explicitly), the priors are adjusted according to the data.

binarize: A double (float64) parameter specifying the threshold for binarizing sample features. (Default: 0.0)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Attributes

class_log_prior_: A python ndarray of double (float64) type values and has shape (***n_classes***,). It contains log probability of each class (smoothed).

feature_log_prob_: A python ndarray of double (float64) type values and has shape (***n_classes***, ***n_features***). It contains empirical log probability of features given a class, $P(x_i|y)$.

class_count_: A python ndarray of double (float64) type values and has shape (***n_classes***,). It contains the number of samples encountered for each class during fitting. This value is weighted by the sample weight when provided.

classes_: A python ndarray of double (float64) type values and has shape (***n_classes***,). It contains the of unique labels given to the classifier during training.

feature_count_: A python ndarray of double (float64) type values and has shape (***n_classes***, ***n_features***). It contains the number of samples encountered for each (class, feature) during fitting. This value is weighted by the sample weight when provided.

coef_: A python ndarray of double (float64) type values.

- If '***classes_***' is 2, then its shape (**1**, ***n_features***).

- If '***classes_***' is more than 2, then its shape is (***n_classes***, ***n_features***).

It mirrors '***feature_log_prob_***' for interpreting BernoulliNB as a linear model.

intercept_: A python ndarray of double (float64) type values.

- If '***classes_***' is 2, the its shape (**1**,).

- If '***classes_***' is more than 2, then its shape is (***n_classes***,).

It mirrors '***class_log_prior_***' for interpreting BernoulliNB as a linear model.

Purpose

It initializes a BernoulliNB object with the given parameters.

Return Value

It simply returns “self” reference.

4.3.1.2 2. fit(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: Any python array-like object or an instance of FrovedisDvector. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training matrix (X) with labels (y) and trains a BernoulliNB model.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# fitting input matrix and label on BernoulliNB object
from frovedis.mllib.linear_model import BernoulliNB
bnb = BernoulliNB(alpha = 1.0).fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisRowmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# BernoulliNB with pre-constructed frovedis-like inputs
from frovedis.mllib.linear_model import BernoulliNB
bnb = BernoulliNB(alpha = 1.0).fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

4.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
# predicting on BernoulliNB model
bnb.predict(mat)
```

Output

```
[1 1 1 ... 1 1 1]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# predicting on BernoulliNB model using pre-constructed input
bnb.predict(rmat)
```

Output

```
[1 1 1 ... 1 1 1]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**,) containing the predicted outputs.

4.3.1.4 4. predict_proba(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server. Unlike Scikit-learn, it performs the classification on an array and returns the probability estimates for the test feature matrix (X).

For example,

```
# finds the probability sample for each class in the model
bnb.predict_proba(mat)
```

Output

```
[[0.35939685 0.64060315]
 [0.35939685 0.64060315]
 [0.35939685 0.64060315]
 ...]
```

```
[0.35939685 0.64060315]
[0.35939685 0.64060315]
[0.35939685 0.64060315]]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# finds the probability sample for each class in the model
bnb.predict_proba(rmat)
```

Output

```
[[0.35939685 0.64060315]
 [0.35939685 0.64060315]
 [0.35939685 0.64060315]
 ...
 [0.35939685 0.64060315]
 [0.35939685 0.64060315]
 [0.35939685 0.64060315]]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape **(n_samples, n_classes)** containing the prediction probability values.

4.3.1.5 5. `score(X, y)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data.

y: Any python array-like object containing true labels for X. It has shape **(n_samples,)**.

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of `self.predict(X)` wrt. `y`.

For example,

```
# calculate mean accuracy score on given test data and labels
bnb.score(mat, lbl)
```

Output

```
0.6274
```

Return Value

It returns an accuracy score of float type.

4.3.1.6 6. `debug_print()`

Purpose

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
bnb.debug_print()
```

Output

```
model_type: bernoulli
binarize: -1
feature_count: 212 212 212 212 212 ... 357 357 357 357 357
theta: node = 0, local_num_row = 2, local_num_col = 30, val = -0.00468385 -0.00468385
-0.00468385 -0.00468385 -0.00468385 ... -0.0027894 -0.0027894 -0.0027894 -0.0027894 -0.0027894
pi: -0.987294 -0.466145
label: 0 1
class count: 212 357
theta_minus_negtheta: node = 0, local_num_row = 2, local_num_col = 30, val = 5.36129
5.36129 5.36129 5.36129 5.36129 ... 5.88053 5.88053 5.88053 5.88053 5.88053
negtheta_sum: -160.979 -176.5
```

This output will be visible on server side. It displays the target model information like `model_type`, `binarize`, `feature_count`, `theta`, `pi`, etc. values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

4.3.1.7 7. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If `True`, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: `True`)

Purpose

This method belongs to the `BaseEstimator` class inherited by `BernoulliNB`. It is used to get parameters and their values of `BernoulliNB` class.

For example,

```
print(bnb.get_params())
```

Output

```
{'alpha': 1.0, 'binarize': -1.0, 'class_prior': None, 'fit_prior': True, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

4.3.1.8 8. `set_params(**params)`

Parameters

*****params***: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the `BaseEstimator` class inherited by `BernoulliNB`, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(bnb.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
bnb.set_params(n_clusters = 4)
print("get parameters after setting:")
print(bnb.get_params())
```

Output

```
get parameters before setting:
{'alpha': 1.0, 'binarize': -1.0, 'class_prior': None, 'fit_prior': True, 'verbose': 0}
get parameters after setting:
{'alpha': 1.0, 'binarize': 0.5, 'class_prior': None, 'fit_prior': True, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

4.3.1.9 9. load(fname, dtype=None)

Parameters

fname: A string object containing the name of the file having model information such as theta, cls_count, feature_count, label, pi, type to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
bnb.load("./out/BNBModel")
```

Return Value

It simply returns “self” reference.

4.3.1.10 10. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the bernoulli naive bayes model
bnb.save("./out/BNBModel")
```

This will save the naive bayes model on the path ‘/out/BNBModel’. It would raise exception if the directory already exists with same name.

The ‘BNBModel’ directory has

BNBModel

```
|—label_map
|—metadata
|—model
```

```

|——cls_count
|——feature_count
|——label
|——pi
|——theta
|——type

```

‘label_map’ contains information about labels mapped with their encoded value.

The metadata file contains the model kind, input datatype used for trained model.

Here, the **model** directory contains information about class count, feature count, labels, pi, **theta** and thier datatype.

Return Value

It returns nothing

4.3.1.11 11. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
bnb.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

4.3.1.12 12. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the naive bayes model first.

Return Value

It returns ‘True’, if the model is already fitted, otherwise, it returns ‘False’.

4.4 SEE ALSO

- Introduction to FrovedisRowmajorMatrix
- Introduction to FrovedisCRSMatrix
- Introduction to FrovedisDvector
- Multinomial Naive Bayes in Frovedis

Chapter 5

DBSCAN

5.1 NAME

DBSCAN clustering - Density-Based Spatial Clustering of Applications with Noise (DBSCAN) is a clustering algorithm commonly used in EDA (exploratory data analysis). It is a base algorithm for density-based clustering, which can discover clusters of different shapes and sizes from a large amount of data, containing noise and outliers.

5.2 SYNOPSIS

```
class frovedis.mllib.cluster.DBSCAN(eps=0.5, min_samples=5, metric='euclidean',
                                     metric_params=None, algorithm='auto',
                                     leaf_size=30, p=None, n_jobs=None,
                                     batch_fraction=None, verbose=0)
```

5.2.1 Public Member Functions

```
fit(X, y = None, sample_weight = None)
fit_predict(X, sample_weight = None)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
release()
is_fitted()
```

5.3 DESCRIPTION

DBSCAN is a Density-Based Clustering unsupervised learning algorithm that identifies distinctive groups/clusters in the data, based on the idea that a cluster in data space is a contiguous region of high point-density, separated from other such clusters by contiguous regions of low point-density.

It is able to find arbitrary shaped clusters and clusters with noise (i.e. outliers). The main idea behind DBSCAN is that a point belongs to a cluster if it is close to many points from that cluster.

DBSCAN algorithm requires two key parameters: **eps** and **min_samples**.

Based on these two parameters, the points are classified as core point, border point, or outlier:

- **Core point:** A point is a core point if there are at least minPts number of points (including the point itself) in its surrounding area with radius eps.
- **Border point:** A point is a border point if it is reachable from a core point and there are less than minPts number of points within its surrounding area.
- **Outlier:** A point is an outlier if it is not a core point and not reachable from any core points.

The main advantage of using DBSCAN is that it is able to find arbitrarily size and arbitrarily shaped clusters and identify noise data while clustering.

This module provides a client-server implementation, where the client application is a normal python program. Frovedis is almost same as Scikit-learn clustering module providing DBSCAN support, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for DBSCAN on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, the python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

5.3.1 Detailed Description

5.3.1.1 1. DBSCAN()

Parameters

eps: A positive double (float64) parameter containing the epsilon value or distance that specifies the neighborhoods. (Default: 0.5)

Two points are considered to be neighbors if the distance between them is less than or equal to eps.

min_samples: A positive integer parameter which specifies the number of samples in a neighborhood for a point to be considered as a core point.

This includes the point itself. (Default: 5)

metric: A string object parameter used when calculating distance between instances in a feature array. (Default: 'euclidean')

It only supports 'euclidean' distance.

metric_params: It is unused parameter which is an additional keyword arguments for the metric function. (Default: None)

algorithm: A string object parameter, used to compute pointwise distances and find nearest neighbors. (Default: 'auto')

When it is 'auto', it will be set as 'brute'(brute-force search approach). Unlike Scikit-learn, currently it supports only 'brute'.

leaf_size: An unused parameter, which is used to affect the speed of the construction and query. (Default: 30)

p: An unused parameter specifying the power of the Minkowski metric to be used to calculate distance between points. (Default: None)

n_jobs: An unused parameter specifying the number of parallel jobs to run. (Default: None)

batch_fraction: A positive double (float64) parameter used to calculate the batches of specific size. These batches are used to construct the distance matrix. (Default: None)

It must be within the range of 0.0 to 1.0. When it is None (not specified explicitly), it will be set as np.finfo(np.float64).max value.

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO

mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Attributes

labels_: It is a python ndarray, containing int64 typed values and has shape (**n_samples**,). These are the coordinates of cluster centers.

core_sample_indices_: It is a python ndarray, containing int32 or int64 typed values and has shape (**n_samples**,). These are the core samples indices.

components_: It is a numpy array or FrovedisRowmajorMatrix, containing float or double (float64) typed values and has shape (**n_samples**, **n_features**).

These are the copy of each core sample found by training.

Purpose

It initializes a DBSCAN object with the given parameters.

The parameters: “metric_params”, “leaf_size”, “p” and “n_jobs” are simply kept to make the interface uniform to Scikit-learn DBSCAN clustering module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

5.3.1.2 2. fit(X, y = None, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: Python array-like containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It clusters the given data points (X).

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on DBSCAN object
from frovedis.mllib.cluster import DBSCAN
dbm = DBSCAN(eps = 5, min_samples = 2).fit(train_mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
```

```

rmat = FrovedisRowmajorMatrix(train_mat)

# DBSCAN with pre-constructed Frovedis-like inputs
from frovedis.mllib.cluster import DBSCAN
dbm = DBSCAN(eps = 5, min_samples = 2).fit(rmat)

```

Return Value

It simply returns “self” reference.

5.3.1.3 3. fit_predict(X, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape **(n_samples, n_features)**.

sample_weight: Python array-like containing the intended weights for each input samples and it should be the shape of **(n_samples,)**.

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It computes the clusters from the given data points (X) or distance matrix and predict labels.

For example,

```

# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on DBSCAN object
from frovedis.mllib.cluster import DBSCAN
dbm = DBSCAN(eps = 5, min_samples = 2)
print(dbm.fit_predict(train_mat))

```

Output

```
[0 0 1 1 1]
```

Like in fit() frovedis-like input can be used to speed-up training at server side.

For example,

```

# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import DBSCAN
dbm = DBSCAN(eps = 5, min_samples = 2)
print(dbm.fit_predict(rmat))

```

Output

```
[0 0 1 1 1]
```

Return Value

It returns a numpy array of int64 type containing the cluster labels, with shape **(n_samples,)**.

5.3.1.4 4. score(X, y, sample_weight = None)

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (n_samples, n_features).

y: A python ndarray or an instance of FrovedisVector containing the true labels for X. It has shape (n_samples,).

sample_weight: An unused parameter whose default value is None. It is simply ignored in frovedis implementation, like in Scikit-learn as well.

Purpose

It calculates homogeneity score on given test data and labels i.e homogeneity score of self.predict(X, y) wrt. y.

For example,

```
dbm.score(train_mat, [0 0 1 1 1])
```

Output

```
1.0
```

Return Value

It returns a score of float type.

5.3.1.5 5. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by DBSCAN. It is used to get parameters and their values of DBSCAN class.

For example,

```
print(dbm.get_params())
```

Output

```
{'algorithm': 'auto', 'batch_fraction': None, 'eps': 5, 'leaf_size': 30,
'metric': 'euclidean', 'metric_params': None, 'min_samples': 2, 'n_jobs': None,
'p': None, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

5.3.1.6 6. set_params(params)****Parameters**

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by DBSCAN, used to set parameter values.

For example,

```

print("Get parameters before setting:")
print(kmeans.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
kmeans.set_params(n_clusters=4, n_init=5)
print("Get parameters after setting:")
print(kmeans.get_params())

```

Output

```

Get parameters before setting:
{'algorithm': 'auto', 'copy_x': True, 'init': 'random', 'max_iter': 300,
'n_clusters': 2, 'n_init': 1, 'n_jobs': 1, 'precompute_distances': 'auto',
'random_state': None, 'tol': 0.0001, 'use_shrink': False, 'verbose': 0}
Get parameters after setting:
{'algorithm': 'auto', 'copy_x': True, 'init': 'random', 'max_iter': 300,
'n_clusters': 4, 'n_init': 5, 'n_jobs': 1, 'precompute_distances': 'auto',
'random_state': None, 'tol': 0.0001, 'use_shrink': False, 'verbose': 0}

```

Return Value

It simply returns “self” reference.

5.3.1.7 7. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
dbm.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

5.3.1.8 8. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, `fit_predict()` is used before training the model, then it can prompt the user to train the clustering model first.

Return Value

It returns ‘True’, if the model is already fitted otherwise, it returns ‘False’.

5.4 SEE ALSO

- [Introduction to FrovedisRowmajorMatrix](#)
- [Introduction to FrovedisCRSMatrix](#)
- [Agglomerative Clustering in Frovedis](#)
- [Spectral Clustering in Frovedis](#)

- **KMeans in Frovedis**

Chapter 6

DecisionTreeClassifier

6.1 NAME

DecisionTreeClassifier - A classification algorithm that predicts the binary and multi-class output using conditional control statements. A decision tree is a decision support tool that uses a tree-like model of decisions and their possible consequences, including chance, event, outcomes, resource costs, and utility.

6.2 SYNOPSIS

```
class frovedis.mllib.tree.DecisionTreeClassifier(criterion='gini', splitter='best',
                                                max_depth=None, min_samples_split=2,
                                                min_samples_leaf=1,
                                                min_weight_fraction_leaf=0.0,
                                                max_features=None,
                                                random_state=None, max_leaf_nodes=None,
                                                min_impurity_decrease=0.0,
                                                class_weight=None, presort='deprecated',
                                                ccp_alpha=0.0, max_bins=32,
                                                verbose=0, categorical_info={})
```

6.2.1 Public Member Functions

```
fit(X, y)
predict(X)
predict_proba(X)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
score(X, y, sample_weight = None)
debug_print()
release()
is_fitted()
```

6.3 DESCRIPTION

Decision Tree Classifier is a supervised learning method used for classification problems. The aim is to create a model that predicts the value of a target variable by learning simple decision rules inferred from the data features. Like any other tree representation, it has a root node, internal nodes, and leaf nodes. The internal node represents condition on attributes, the branches represent the results of the condition and the leaf node represents the class label. **Frovedis supports both binary and multinomial decision tree classification algorithms.**

During training, the input \mathbf{X} is the training data and \mathbf{y} is the corresponding label values (Frovedis supports any values for labels, but internally it encodes the input binary labels to 0 and 1, and input multinomial labels to 0, 1, 2, ..., N-1 (where N is the no. of classes) before training at Frovedis server) which we want to predict.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn DecisionTreeClassifier interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for DecisionTreeClassifier on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

6.3.1 Detailed Description

6.3.1.1 1. DecisionTreeClassifier()

Parameters

criterion: A string object parameter that specifies the function to measure the quality of a split. Supported criteria are 'gini' and 'entropy'. (Default: 'gini')

- **'gini' impurity:** calculates the amount of probability of a specific feature that is classified incorrectly when selected randomly.

- **'entropy' (information gain):** it is applied to quantify which feature provides maximal information about the classification based on the notion of entropy.

splitter: An unused parameter. (Default: 'best')

max_depth: A positive integer parameter that specifies the maximum depth of the tree. (Default: None) If it is None (not specified explicitly), then 'max_depth' is set to 5.

min_samples_split: An unused parameter. (Default: 2)

min_samples_leaf: A positive integer or float value that specifies the minimum number of samples required to be at a leaf node. A split point at any depth will only be considered, if it leaves, at least 'min_samples_leaf' training samples in each of the left and right branches. (Default: 1)

- If it is an integer, then 'min_samples_leaf' should be greater than or equal to 1.

- If it is float, then 'min_samples_leaf' should be in range (0,0.5].

min_weight_fraction_leaf: An unused parameter. (Default: 0.0)

max_features: An unused parameter. (Default: None)

random_state: An unused parameter. (Default: None)

max_leaf_nodes: An unused parameter. (Default: None)

min_impurity_decrease: A positive double (float64) parameter. A node will be split if this split induces

a decrease of the impurity greater than or equal to this value. (Default: 0.0)

class_weight: An unused parameter. (Default: None)

presort: An unused parameter. (Default: 'deprecated')

ccp_alpha: An unused parameter. (Default: 0.0)

max_bins: A positive integer parameter that specifies the maximum number of bins created by ordered splits. (Default: 32)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

categorical_info: A dictionary that specifies categorical features information. Here, it gives column indices of categorical features and the number of categories for those features.

For example, a dictionary { {0, 2}, {4, 5} } means that the feature[0] takes values 0 or 1 (binary) and the feature[4] has five categories (values 0, 1, 2, 3 or 4). Note that feature indices and category assignments are 0-based. (Default: {})

Attributes

classes: It is a python ndarray (any type) of unique labels given to the classifier during training. It has shape (n_classes,).

n_features: An integer value specifying the number of features when fitting the estimator.

Purpose

It initializes a DecisionTreeClassifier object with the given parameters.

The parameters: “splitter”, “min_samples_split”, “min_weight_fraction_leaf”, “max_features”, “random_state”, “max_leaf_nodes”, “class_weight”, “presort” and “ccp_alpha” are simply kept in to make the interface uniform to the Scikit-learn DecisionTreeClassifier module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

6.3.1.2 2. fit(X, y)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisColmajorMatrix for dense data. It has shape (n_samples, n_features).

y: Any python array-like object or an instance of FrovedisDvector containing the class labels. It has shape (n_samples,).

Purpose

It builds a decision tree classifier from the training data X and labels y.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)
```

```
# fitting input matrix and label on DecisionTreeClassifier object
from frovedis.ml.lib.tree import DecisionTreeClassifier
dtc = DecisionTreeClassifier(max_depth = 5)
dtc.fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# DecisionTreeClassifier with pre-constructed frovedis-like inputs
from frovedis.mllib.tree import DecisionTreeClassifier
dtc = DecisionTreeClassifier(max_depth = 5)
dtc.fit(cmat,dlbl)
```

Return Value

It simply returns “self” reference.

6.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

Predict class for X.

For a classification model, the predicted class value for each sample in X is returned.

For example,

```
# predicting on decision tree classifier model
dtc.predict(mat)
```

Output

```
[0 0 0 ... 0 0 1]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on decision tree classifier model using pre-constructed input
dtc.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[0 0 0 ... 0 0 1]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**,) containing the predicted classes.

6.3.1.4 4. predict_proba(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

Predict class log-probabilities of the input samples X.

Currently, this method is not supported for multinomial classification problems.

For example,

```
# finds the probability sample for each class in the model
dtc.predict_proba(mat)
```

Output

```
[[1.      0.      ]
 [1.      0.      ]
 [1.      0.      ]
 ...
 [1.      0.      ]
 [1.      0.      ]
 [0.009375 0.990625]]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# finds the probability sample for each class in the model
dtc.predict_proba(cmat.to_frovedis_rowmatrix())
```

Output

```
[[1.      0.      ]
 [1.      0.      ]
 [1.      0.      ]
 ...
 [1.      0.      ]
 [1.      0.      ]
 [0.009375 0.990625]]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**, **n_classes**) containing the prediction probability values. Since only binary classification problems (**n_classes = 2**) are supported for this method, so currently its shape will be (**n_samples**, **2**).

6.3.1.5 5. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by DecisionTreeClassifier. It is used to get parameters and their values of DecisionTreeClassifier class.

For example,

```
print(dtc.get_params())
```

Output

```
{'categorical_info': {}, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'GINI',
'max_bins': 32, 'max_depth': 5, 'max_features': None, 'max_leaf_nodes': None,
'min_impurity_decrease': 0.0, 'min_samples_leaf': 1, 'min_samples_split': 2,
'min_weight_fraction_leaf': 0.0, 'presort': 'deprecated', 'random_state': None,
'splitter': 'best', 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

6.3.1.6 6. set_params(**params)

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by DecisionTreeClassifier, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(dtc.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
dtc.set_params(criterion = 'entropy', max_depth = 5)
print("get parameters after setting:")
print(dtc.get_params())
```

Output

```
get parameters before setting:
{'categorical_info': {}, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'GINI',
'max_bins': 32, 'max_depth': 5, 'max_features': None, 'max_leaf_nodes': None,
'min_impurity_decrease': 0.0, 'min_samples_leaf': 1, 'min_samples_split': 2,
'min_weight_fraction_leaf': 0.0, 'presort': 'deprecated', 'random_state': None,
'splitter': 'best', 'verbose': 0}
get parameters after setting:
{'categorical_info': {}, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'entropy',
'max_bins': 32, 'max_depth': 5, 'max_features': None, 'max_leaf_nodes': None,
'min_impurity_decrease': 0.0, 'min_samples_leaf': 1, 'min_samples_split': 2,
'min_weight_fraction_leaf': 0.0, 'presort': 'deprecated', 'random_state': None,
'splitter': 'best', 'verbose': 0}
```

Return Value

It simply returns “self” reference.

6.3.1.7 7. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
dtc.load("./out/MyDecisionTreeClassifierModel")
```

Return Value

It simply returns “self” reference.

6.3.1.8 8. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the decision tree classifier model
dtc.save("./out/MyDecisionTreeClassifierModel")
```

This will save the decision tree classifier model on the path “/out/MyDecisionTreeClassifierModel”. It would raise exception if the directory already exists with same name.

The ‘MyDecisionTreeClassifierModel’ directory has

MyDecisionTreeClassifierModel

```
|—label_map
|—metadata
|—model
```

‘label_map’ file contains information about labels mapped with their encoded value.

The ‘metadata’ file contains the number of classes, model kind and input datatype used for trained model.

The ‘model’ file contains the decision tree model saved in binary format.

Return Value

It returns nothing.

6.3.1.9 9. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the true labels for X. It has shape (**n_samples**,).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of self.predict(X) wrt. y.

For example,

```
# calculate mean accuracy score on given test data and labels
dtc.score(mat,lbl)
```

Output

0.9895

Return Value

It returns an accuracy score of float type.

6.3.1.10 10. debug_print()

Purpose

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
dtc.debug_print()
```

Output

```
----- Classification Tree:: -----
# of nodes: 35, height: 5
<1> Split: feature[22] < 113.157, IG: 0.319406
| \_ <2> Split: feature[27] < 0.144844, IG: 0.0626618
|   | \_ <4> Split: feature[22] < 107.223, IG: 0.00955662
|   |   | \_ <8> Split: feature[10] < 0.904737, IG: 0.00582386
|   |   |   | \_ <16> Split: feature[27] < 0.133781, IG: 0.00558482
|   |   |   |   | \_ (32) Predict: 1 (99.0625%)
|   |   |   |   | \_ (33) Predict: 1 (66.6667%)
|   |   |   |   | \_ (17) Predict: 0 (100%)
|   |   |   | \_ <9> Split: feature[14] < 0.00570775, IG: 0.184704
|   |   |   |   | \_ <18> Split: feature[0] < 14.0963, IG: 0.165289
|   |   |   |   |   | \_ (36) Predict: 0 (100%)
|   |   |   |   |   | \_ (37) Predict: 1 (100%)
|   |   |   |   | \_ <19> Split: feature[21] < 21.9025, IG: 0.408163
|   |   |   |   |   | \_ (38) Predict: 1 (100%)
|   |   |   |   |   | \_ (39) Predict: 0 (100%)
|   |   |   | \_ <5> Split: feature[21] < 24.13, IG: 0.189372
|   |   |   |   | \_ <10> Split: feature[4] < 0.109085, IG: 0.32
|   |   |   |   |   | \_ (20) Predict: 1 (100%)
|   |   |   |   |   | \_ (21) Predict: 0 (100%)
|   |   |   |   | \_ <11> Split: feature[9] < 0.0609525, IG: 0.144027
|   |   |   |   |   | \_ (22) Predict: 1 (100%)
|   |   |   |   |   | \_ <23> Split: feature[7] < 0.0404469, IG: 0.0907029
|   |   |   |   |   |   | \_ (46) Predict: 1 (100%)
|   |   |   |   |   |   | \_ (47) Predict: 0 (100%)
|   |   |   | \_ <3> Split: feature[7] < 0.0509028, IG: 0.0364914
|   |   |   |   | \_ <6> Split: feature[17] < 0.00995862, IG: 0.259286
|   |   |   |   |   | \_ <12> Split: feature[1] < 15.855, IG: 0.336735
|   |   |   |   |   |   | \_ (24) Predict: 1 (100%)
|   |   |   |   |   |   | \_ (25) Predict: 0 (100%)
|   |   |   |   |   | \_ (13) Predict: 1 (100%)
|   |   |   | \_ <7> Split: feature[1] < 13.9925, IG: 0.0116213
|   |   |   |   | \_ <14> Split: feature[1] < 12.18, IG: 0.5
|   |   |   |   |   | \_ (28) Predict: 0 (100%)
```

```
| \_ (29) Predict: 1 (100%)
| \_ (15) Predict: 0 (100%)
```

This output will be visible on server side. It displays the decision tree having maximum depth of 5 and total 35 nodes in the tree.

No such output will be visible on client side.

Return Value

It returns nothing.

6.3.1.11 11. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
dtc.release()
```

This will reset the after-fit populated attributes (like `classes__`) to None, along with releasing server side memory.

Return Value

It returns nothing.

6.3.1.12 12. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

6.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Decision Tree Regressor in Frovedis**

Chapter 7

DecisionTreeRegressor

7.1 NAME

DecisionTreeRegressor - A regression algorithm used to predict the output using conditional control statements. A decision tree is a decision support tool that uses a tree-like model of decisions and their possible consequences, including chance, event, outcomes, resource costs, and utility.

7.2 SYNOPSIS

```
class frovedis.mllib.tree.DecisionTreeRegressor(criterion='mse', splitter='best',
                                                max_depth=None, min_samples_split=2,
                                                min_samples_leaf=1,
                                                min_weight_fraction_leaf=0.0,
                                                max_features=None, random_state=None,
                                                max_leaf_nodes=None,
                                                min_impurity_decrease=0.0,
                                                presort='deprecated', ccp_alpha=0.0,
                                                max_bins=32, verbose=0,
                                                categorical_info={})
```

7.2.1 Public Member Functions

```
fit(X, y)
predict(X)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
score(X, y, sample_weight = None)
debug_print()
release()
is_fitted()
```

7.3 DESCRIPTION

Decision Tree Regressor is a supervised learning method used for regression problems. The aim is to create a model that predicts the value of a target variable by learning simple decision rules inferred from the data features. Like any other tree representation, it has a root node, internal nodes, and leaf nodes. The internal node represents condition on attributes, the branches represent the results of the condition and the leaf node represents the class label.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn DecisionTreeRegressor interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for DecisionTreeRegressor on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

7.3.1 Detailed Description

7.3.1.1 1. DecisionTreeRegressor()

Parameters

criterion: A string object parameter that specifies the function to measure the quality of a split.

Currently, supported criteria is 'mse'. The mean squared error (mse), is equal to variance reduction as a feature selection criterion and minimizes the L2 loss using the mean of each terminal node. (Default: 'mse')

splitter: An unused parameter. (Default: 'best')

max_depth: A positive integer parameter that specifies the maximum depth of the tree. (Default: None) When it is None (not specified explicitly), then 'max_depth' is set to 5.

min_samples_split: An unused parameter. (Default: 2)

min_samples_leaf: A positive integer or float value that specifies the minimum number of samples required to be at a leaf node. A split point at any depth will only be considered if it leaves at least 'min_samples_leaf' training samples in each of the left and right branches. This may have the effect of smoothing the model, especially in regression. (Default: 1)

- If it is an integer, then 'min_samples_leaf' should be greater than or equal to 1.

- If it is float, then 'min_samples_leaf' should be in range **(0,0.5]**.

min_weight_fraction_leaf: An unused parameter. (Default: 0.0)

max_features: An unused parameter. (Default: None)

random_state: An unused parameter. (Default: None)

max_leaf_nodes: An unused parameter. (Default: None)

min_impurity_decrease: A positive float parameter. A node will be split if this split induces a decrease of the impurity greater than or equal to this value. (Default: 0.0)

presort: An unused parameter. (Default: 'deprecated')

ccp_alpha: An unused parameter. (Default: 0.0)

max_bins: A positive integer parameter that specifies the maximum number of bins created by ordered splits. (Default: 32)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

categorical_info: A dictionary that specifies categorical features information. Here, it gives column indices of categorical features and the number of categories for those features.

For example, a dictionary `{ {0, 2}, {4, 5} }` means that the feature[0] takes values 0 or 1 (binary) and the feature[4] has five categories (values 0, 1, 2, 3 or 4). Note that feature indices and category assignments are 0-based. (Default: `{}`)

Attributes

n_features: An integer value specifying the number of features when fitting the estimator.

Purpose

It initializes a DecisionTreeRegressor object with the given parameters.

The parameters: “splitter”, “min_samples_split”, “min_weight_fraction_leaf”, “max_features”, “random_state”, “max_leaf_nodes”, “presort” and “ccp_alpha” are simply kept in to make the interface uniform to the Scikit-learn DecisionTreeRegressor module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

7.3.1.2 2. fit(X, y)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisColmajorMatrix for dense data. It has shape **(n_samples, n_features)**.

y: Any python array-like object or an instance of FrovedisDvector containing the target values for X. It has shape **(n_samples,)**.

Purpose

It builds a decision tree regressor from the training data X and labels y.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_boston
mat, lbl = load_boston(return_X_y = True)

# fitting input matrix and label on DecisionTreeRegressor object
from frovedis.mllib.tree import DecisionTreeRegressor
dtr = DecisionTreeRegressor(max_depth = 5)
dtr.fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_boston
mat, lbl = load_boston(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)
```

```
# DecisionTreeRegressor with pre-constructed frovedis-like inputs
from frovedis.mllib.tree import DecisionTreeRegressor
dtr = DecisionTreeRegressor(max_depth = 5)
dtr.fit(cmat,dlbl)
```

Return Value

It simply returns “self” reference.

7.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

Predict regression value for X.

For a regression model, the predicted value based on X is returned.

For example,

```
# predicting on decision tree regressor model
dtr.predict(mat)
```

Output

```
[24.64166667 24.64166667 34.80666667 ... 27.83636364 27.83636364 20.98461538]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on decision tree regressor model using pre-constructed input
dtr.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[24.64166667 24.64166667 34.80666667 ... 27.83636364 27.83636364 20.98461538]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**,) containing the predicted values.

7.3.1.4 4. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by DecisionTreeRegressor. It is used to get parameters and their values of DecisionTreeRegressor class.

For example,

```
print(dtr.get_params())
```

Output

```
{'categorical_info': {}, 'ccp_alpha': 0.0, 'criterion': 'MSE', 'max_bins': 32,
'max_depth': 5, 'max_features': None, 'max_leaf_nodes': None, 'min_impurity_decrease': 0.0,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'presort': 'deprecated', 'random_state': None, 'splitter': 'best', 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

7.3.1.5 5. `set_params(**params)`

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by DecisionTreeRegressor, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(dtr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
dtr.set_params(max_depth = 4)
print("get parameters after setting:")
print(dtr.get_params())
```

Output

```
get parameters before setting:
{'categorical_info': {}, 'ccp_alpha': 0.0, 'criterion': 'MSE', 'max_bins': 32,
'max_depth': 5, 'max_features': None, 'max_leaf_nodes': None, 'min_impurity_decrease': 0.0,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'presort': 'deprecated', 'random_state': None, 'splitter': 'best', 'verbose': 0}
get parameters after setting:
{'categorical_info': {}, 'ccp_alpha': 0.0, 'criterion': 'MSE', 'max_bins': 32,
'max_depth': 4, 'max_features': None, 'max_leaf_nodes': None, 'min_impurity_decrease': 0.0,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'presort': 'deprecated', 'random_state': None, 'splitter': 'best', 'verbose': 0}
```

Return Value

It simply returns “self” reference.

7.3.1.6 6. `load(fname, dtype = None)`

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
dtr.load("./out/MyDecisionTreeRegressorModel")
```

Return Value

It simply returns “self” reference.

7.3.1.7 7. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the decision tree regressor model
dtr.save("./out/MyDecisionTreeRegressorModel")
```

This will save the decision tree regressor model on the path “/out/MyDecisionTreeRegressorModel”. It would raise exception if the directory already exists with same name.

The ‘MyDecisionTreeRegressorModel’ directory has

MyDecisionTreeRegressorModel

```
|—metadata
|—model
```

The ‘metadata’ file contains the information on model kind and input datatype used for trained model.

The ‘model’ file contains the decision tree model saved in binary format.

Return Value

It returns nothing.

7.3.1.8 8. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object or an instance of FrovedisDvector containing the true values for X. It has shape (**n_samples**,).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of self.predict(X) wrt. y.

The coefficient ‘R2’ is defined as $(1 - (u/v))$,

where ‘u’ is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and,

‘v’ is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
# calculate R2 score on given test data and labels
dtr.score(mat, lbl)
```

Output

0.9109

Return Value

It returns an R2 score of float type.

7.3.1.9 9. debug_print()

Purpose

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
dtr.debug_print()
```

Output

```
----- Regression Tree:: -----
# of nodes: 51, height: 5
<1> Split: feature[12] < 9.6575, IG: 37.2288
 \_ <2> Split: feature[5] < 7.14906, IG: 37.6417
  | \_ <4> Split: feature[7] < 1.47325, IG: 17.8515
  | | \_ (8) Predict: 50
  | | \_ <9> Split: feature[5] < 6.69962, IG: 9.19518
  | | | \_ <18> Split: feature[5] < 6.10819, IG: 2.41396
  | | | | \_ (36) Predict: 20.9846
  | | | | \_ (37) Predict: 24.6417
  | | | \_ <19> Split: feature[12] < 5.27, IG: 6.57304
  | | | | \_ (38) Predict: 32.9652
  | | | | \_ (39) Predict: 27.8364
  | | \_ <5> Split: feature[5] < 7.45969, IG: 25.6196
  | | | \_ <10> Split: feature[5] < 7.43531, IG: 4.71838
  | | | | \_ <20> Split: feature[8] < 7.125, IG: 1.42815
  | | | | | \_ (40) Predict: 34.8067
  | | | | | \_ (41) Predict: 31.6
  | | | | \_ (21) Predict: 44
  | | | \_ <11> Split: feature[0] < 2.1766, IG: 19.28
  | | | | \_ <22> Split: feature[10] < 17.9563, IG: 5.68288
  | | | | | \_ (44) Predict: 46.9375
  | | | | | \_ (45) Predict: 40.125
  | | | | \_ (23) Predict: 21.9
  | \_ <3> Split: feature[12] < 14.9763, IG: 9.38095
  | | \_ <6> Split: feature[5] < 6.56925, IG: 1.74534
  | | | \_ <12> Split: feature[11] < 101.175, IG: 0.921292
  | | | | \_ <24> Split: feature[4] < 0.5995, IG: 6.02083
  | | | | | \_ (48) Predict: 16.5667
  | | | | | \_ (49) Predict: 10.9
  | | | | \_ <25> Split: feature[2] < 2.57187, IG: 0.586878
  | | | | | \_ (50) Predict: 25.3667
  | | | | | \_ (51) Predict: 20.4598
  | | \_ <13> Split: feature[0] < 4.94925, IG: 10.4783
  | | | \_ <26> Split: feature[4] < 0.59225, IG: 7.70469
```

```

|      | \_ (52) Predict: 24.4429
|      | \_ (53) Predict: 30.5
|      \_ (27) Predict: 15
\_ <7> Split: feature[0] < 5.64819, IG: 5.7942
  \_ <14> Split: feature[4] < 0.52725, IG: 2.64736
    | \_ <28> Split: feature[12] < 29.8562, IG: 2.20522
    | | \_ (56) Predict: 20.4
    | | \_ (57) Predict: 15.45
    | \_ <29> Split: feature[12] < 18.0497, IG: 1.97256
    | \_ (58) Predict: 17.5667
    | \_ (59) Predict: 14.7563
  \_ <15> Split: feature[12] < 20.0091, IG: 3.36396
    \_ <30> Split: feature[5] < 5.92387, IG: 2.19683
      | \_ (60) Predict: 11.4714
      | \_ (61) Predict: 15.137
    \_ <31> Split: feature[4] < 0.67175, IG: 4.84222
      \_ (62) Predict: 13.88
      \_ (63) Predict: 9.15938

```

This output will be visible on server side. It displays the decision tree having maximum depth of 5 and total 51 nodes in the tree.

No such output will be visible on client side.

Return Value

It returns nothing.

7.3.1.10 10. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
dtr.release()
```

This will remove the trained model, model-id present on server, along with releasing server side memory.

Return Value

It returns nothing.

7.3.1.11 11. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

7.4 SEE ALSO

- [Introduction to FrovedisRowmajorMatrix](#)
- [Introduction to FrovedisColmajorMatrix](#)

- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Decision Tree Classifier in Frovedis**

Chapter 8

FactorizationMachineClassifier

8.1 NAME

FactorizationMachineClassifier - A factorization machine is a general-purpose supervised learning algorithm that can be used for classification tasks. It is an extension of a linear model that is designed to capture interactions between features within high dimensional sparse datasets.

8.2 SYNOPSIS

```
class frovedis.mllib.fm.FactorizationMachineClassifier(iteration=100, init_stdev=0.1,
                                                       init_learn_rate=0.01, optimizer="SGD",
                                                       dim=(True,True, 8), reg=(0, 0, 0),
                                                       batch_size_pernode=100, verbose=0)
```

8.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
load(fname, dtype = None)
score(X, y, sample_weight = None)
save(fname) get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

8.3 DESCRIPTION

The FactorizationMachineClassifier (fmc) is a general predictor like SVMs but is also able to estimate reliable parameters under very high sparsity. The factorization machine models all nested variable interactions (comparable to a polynomial kernel in SVM), but uses a factorized parameterization instead of a dense parametrization like in SVMs. We show that the model equation of fmcs can be computed in linear time and that it depends only on a linear number of parameters. This allows direct optimization and storage of model

parameters without the need of storing any training data (e.g. support vectors) for prediction. **Frovedis supports both binary and multinomial labels.**

During training, the input \mathbf{X} is the training data and \mathbf{y} are their corresponding label values (Frovedis supports any values as for labels, but internally it encodes the input binary labels to -1 and 1, before training at Frovedis server) which we want to predict.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as libFM. libFM is a software implementation for factorization machines that features stochastic gradient descent (SGD) and alternating least squares (ALS) optimization as well as Bayesian inference using Markov Chain Monte Carlo (MCMC). In this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for FactorizationMachineClassifier on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When predict-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

8.3.1 Detailed Description

8.3.1.1 1. FactorizationMachineClassifier()

Parameters

iteration: A positive integer parameter, specifying the maximum number of iteration count. (Default: 100)

init_stddev: A positive double parameter specifying the standard deviation which is used to initialize the model parameter of 2-way factors. (Default: 0.1)

init_learn_rate: A double parameter containing the learning rate for SGD optimizer. (Default: 0.01)

It should be in range from 0.00001 to 1.0.

optimizer: A string object parameter that specifies which algorithms minimize or maximize a Loss function $E(x)$ using its gradient values with respect to the parameters. (Default: 'SGD')

Only 'SGD' is supported.

dim: A tuple that specifies three important parameters with default values- (True, True, 8):

- **global_bias:** A boolean value that represents a switch to use bias. Currently, this parameter is not used in Frovedis implementation.

- **dim_one_interactions :** A boolean value represents a switch to use 1-way interaction.

- **dim_factors_no :** A positive integer that represents the dimension of 2-way interaction or number of factors that are used for pairwise interactions.

When any of the three is None (not specified explicitly), then user experiences an Error.

reg: A tuple that specifies three important parameters with default values- (Default: (0, 0, 0))

- **regularization_intercept :** A positive integer that represents the regularization parameters of intercept or bias regularization.

- **regularization_one_interactions :** A positive integer that represents the switch to use 1-way regularization.

- **regularization_factors_no :** A positive integer that represents the dimension of 2-way interaction or number of factors that are used for pairwise regularization.

When any of the three is None (not specified explicitly), then user experiences an Error.

batch_size_pernode: A positive integer parameter specifies the size of minibatch processed by one node. (Default: 100)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO

mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Attributes

classes: A numpy array of long (int64) type value that specifies unique labels given to the classifier during training. It has shape **(n_classes,)**, where n_classes is the unique number of classes.

Purpose

It initializes a FactorizationMachineClassifier object with the given parameters.

Return Value

It simply returns “self” reference.

8.3.1.2 2. fit(X, y, sample_weight=None)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix of float or double (float64) type. It has shape **(n_samples, n_features)**.

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape **(n_samples,)**.

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of **(n_samples,)**. (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It accepts the training matrix (X) with labels (y) and trains a FactorizationMachineClassifier model.

For example,

```
# loading sample data
row = np.array([0, 0, 0, 1, 1, 1, 1, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 5, 5])
col = np.array([0, 1, 4, 0, 1, 2, 4, 1, 2, 3, 2,3,4,5, 0, 1, 3, 4, 3, 5])
data = np.asarray([ 2.0, 1.0,1.0, 1.0, 3.0,1.0, 1.0, 1.0, 2.0,1.0, 1.0, 3.0,
                    1.0,1.0,1.0, 1.0, 1.0, 3.0,1.0, 1.0])
csr = csr_matrix((data, (row, col)), shape = (6, 6))
lbl = [10.0, 10.0, 10.0, 20.0, 10.0, 20.0]

# fitting input data on FactorizationMachineClassifier object
from frovedis.mllib.fm import FactorizationMachineClassifier
fmc = FactorizationMachineClassifier()
fmc.fit(csr, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample data
row = np.array([0, 0, 0, 1, 1, 1, 1, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 5, 5])
col = np.array([0, 1, 4, 0, 1, 2, 4, 1, 2, 3, 2,3,4,5, 0, 1, 3, 4, 3, 5])
data = np.asarray([ 2.0, 1.0,1.0, 1.0, 3.0,1.0, 1.0, 1.0, 2.0,1.0, 1.0, 3.0,
                    1.0,1.0,1.0, 1.0, 1.0, 3.0,1.0, 1.0])
csr = csr_matrix((data, (row, col)), shape = (6, 6))
lbl = [10.0, 10.0, 10.0, 20.0, 10.0, 20.0]

# Since "csr" is scipy sparse data, we have created FrovedisCRSMatrix.
from frovedis.matrix.crs import FrovedisCRSMatrix
```

```

from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisCRSMatrix(csr)
dlbl = FrovedisDvector(lbl)

# fitting input data on FactorizationMachineClassifier object
from frovedis.mllib.fm import FactorizationMachineClassifier
fmc = FactorizationMachineClassifier()
fmc.fit(cmat, dlbl)

```

Return Value

It simply returns “self” reference.

8.3.1.3 3. predict(X)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix of float or double (float64) type. It has shape (n_samples, n_features).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```

# predicting on FactorizationMachineClassifier model
fmc.predict(csr)

```

Output

```
[20. 20. 20. 20. 20. 20.]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```

# Since "csr" is scipy sparse data, we have created FrovedisCRSMatrix.
from frovedis.matrix.crs import FrovedisCRSMatrix
cmat = FrovedisCRSMatrix(csr)

# predicting on FactorizationMachineClassifier model using pre-constructed input
fmc.predict(cmat)

```

Output

```
[20. 20. 20. 20. 20. 20.]
```

Return Value

It returns a numpy array of double (float64) type containing the predicted outputs. It is of shape (n_samples,).

8.3.1.4 4. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

Currently, this method is not supported for FactorizationMachineClassifier. It is simply kept in FactorizationMachineClassifier module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an AttributeError.

8.3.1.5 5. score(X, y, sample_weight = None)**Parameters**

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing class labels. It is of shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of self.predict(X) wrt. y.

For example,

```
fmc.score(csr, lbl)
```

Output

```
0.83
```

Return Value

It returns an accuracy score of double (float64) type.

8.3.1.6 6. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information(label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the FactorizationMachineClassifier model
fmc.save("./out/FMCMModel")
```

The FMCMModel contains below directory structure:

FMCMModel

```
|—label_map
|—metadata
|—model
```

‘label_map’ contains information about labels mapped with their encoded value.

‘metadata’ represents the detail about model_kind and datatype of training vector.

Here, the model file contains information about trained model in binary format.

This will save the FactorizationMachineClassifier model on the path ‘/out/FMCMModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

8.3.1.7 7. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by FactorizationMachineClassifier. It is used to get parameters and their values of FactorizationMachineClassifier class.

For example,

```
print(fmc.get_params())
```

Output

```
{'batch_size_pernode': 100, 'dim': (True, True, 8), 'init_learn_rate': 0.01,
'init_stddev': 0.1, 'iteration': 100, 'optimizer': 'SGD', 'reg': (0, 0, 0),
'verbos': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

8.3.1.8 8. `set_params(**params)`

Parameters

****params:** All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by FactorizationMachineClassifier, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(fmc.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
fmc.set_params(iteration = 200)
print("Get parameters after setting:")
print(fmc.get_params())
```

Output

```
Get parameters before setting:
{'batch_size_pernode': 100, 'dim': (True, True, 8), 'init_learn_rate': 0.01,
'init_stddev': 0.1, 'iteration': 100, 'optimizer': 'SGD', 'reg': (0, 0, 0),
'verbos': 0}
Get parameters after setting:
{'batch_size_pernode': 100, 'dim': (True, True, 8), 'init_learn_rate': 0.01,
'init_stddev': 0.1, 'iteration': 200, 'optimizer': 'SGD', 'reg': (0, 0, 0),
'verbos': 0}
```

Return Value

It simply returns “self” reference.

8.3.1.9 9. debug_print()**Purpose**

Currently, this method is not supported for FactorizationMachineClassifier. It is simply kept in FactorizationMachineClassifier module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an AttributeError.

8.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
fmc.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

8.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

8.4 SEE ALSO

- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Factorization Machine Regressor in Frovedis**

Chapter 9

FactorizationMachineRegressor

9.1 NAME

FactorizationMachineRegressor - A factorization machine is a general-purpose supervised learning algorithm that can be used for regression tasks. It is an extension of a linear model that is designed to capture interactions between features within high dimensional sparse datasets.

9.2 SYNOPSIS

```
class frovedis.mllib.fm.FactorizationMachineRegressor(iteration=100, init_stdev=0.1,
                                                    init_learn_rate=0.01, optimizer="SGD",
                                                    dim=(True,True, 8), reg=(0, 0, 0),
                                                    batch_size_pernode=100, verbose=0)
```

9.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
load(fname, dtype = None)
save(fname)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

9.3 DESCRIPTION

The FactorizationMachineRegressor (fmr) is a general predictor like SVMs but is also able to estimate reliable parameters under very high sparsity. The factorization machine models all nested variable interactions (comparable to a polynomial kernel in SVM), but uses a factorized parameterization instead of a dense parametrization like in SVMs. We show that the model equation of fmrs can be computed in linear time and

that it depends only on a linear number of parameters. This allows direct optimization and storage of model parameters without the need of storing any training data (e.g. support vectors) for prediction.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as libFM. libFM is a software implementation for factorization machines that features stochastic gradient descent (SGD) and alternating least squares (ALS) optimization as well as Bayesian inference using Markov Chain Monte Carlo (MCMC). In this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for FactorizationMachineRegressor on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When predict-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

9.3.1 Detailed Description

9.3.1.1 1. FactorizationMachineRegressor()

Parameters

iteration: A positive integer parameter, specifying the maximum number of iteration count. (Default: 100)

init_stddev: A positive double parameter specifying the standard deviation which is used to initialize the model parameter of 2-way factors (Default: 0.1)

init_learn_rate: A double parameter containing the learning rate for SGD optimizer. (Default: 0.01)

It should be in range from 0.00001 to 1.0.

optimizer: A string object parameter that specifies which algorithms minimize or maximize a Loss function $E(x)$ using its Gradient values with respect to the parameters. (Default: 'SGD')

Only 'SGD' is supported.

dim: A tuple that specifies three important parameters with default values- (True, True, 8):

- **global_bias:** A boolean value that represents a switch to use bias. Currently, this parameter is not used in Frovedis implementation.

- **dim_one_interactions** : A boolean value that represents a switch to use 1-way interaction.

- **dim_factors_no** : A positive integer that represents the dimension of 2-way interaction or number of factors that are used for pairwise interactions.

When any of the three is None (not specified explicitly), then user experiences an Error.

reg: An tuple of values that specifies three important parameters with default values- (Default: (0, 0, 0))

- **regularization_intercept** : A positive integer that represents the regularization parameters of intercept or bias regularization.

- **regularization_one_interactions** : A positive integer that represents the switch to use 1-way regularization.

- **regularization_factors_no** : A positive integer that represents the dimension of 2-way interaction or number of factors that are used for pairwise regularization.

When any of the three is None (not specified explicitly), then user experiences an Error.

batch_size_pernode: A positive integer parameter specifies the size of minibatch processed by one node. (Default: 100)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Purpose

It initializes a FactorizationMachineRegressor object with the given parameters.

Return Value

It simply returns “self” reference.

9.3.1.2 2. fit(X, y, sample_weight=None)**Parameters**

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It accepts the training matrix (X) with labels (y) and trains a FactorizationMachineRegressor model.

For example,

```
# loading sample data
row = np.array([0, 0, 0, 1, 1, 1, 1, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 5, 5])
col = np.array([0, 1, 4, 0, 1, 2, 4, 1, 2, 3, 2,3,4,5, 0, 1, 3, 4, 3, 5])
data = np.asarray([ 2.0, 1.0,1.0, 1.0, 3.0,1.0, 1.0, 1.0, 2.0,1.0, 1.0, 3.0,
                    1.0,1.0,1.0, 1.0, 1.0, 3.0,1.0, 1.0])
csr = csr_matrix((data, (row, col)), shape = (6, 6))
lbl = [10.0, 10.0, 10.0, 20.0, 10.0, 20.0]

# fitting input data on FactorizationMachineRegressor object
from frovedis.mllib.fm import FactorizationMachineRegressor
fmr = FactorizationMachineRegressor()
fmr.fit(csr, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample data
row = np.array([0, 0, 0, 1, 1, 1, 1, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 5, 5])
col = np.array([0, 1, 4, 0, 1, 2, 4, 1, 2, 3, 2,3,4,5, 0, 1, 3, 4, 3, 5])
data = np.asarray([ 2.0, 1.0,1.0, 1.0, 3.0,1.0, 1.0, 1.0, 2.0,1.0, 1.0, 3.0,
                    1.0,1.0,1.0, 1.0, 1.0, 3.0,1.0, 1.0])
csr = csr_matrix((data, (row, col)), shape = (6, 6))
lbl = [10.0, 10.0, 10.0, 20.0, 10.0, 20.0]

# Since "csr" is scipy sparse data, we have created FrovedisCRSMatrix.
from frovedis.matrix.crs import FrovedisCRSMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisCRSMatrix(csr)
dlbl = FrovedisDvector(lbl)

# fitting input data on FactorizationMachineRegressor object
from frovedis.mllib.fm import FactorizationMachineRegressor
```

```
fmr = FactorizationMachineRegressor()
fmr.fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

9.3.1.3 3. predict(X)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix of float or double (float64) type. It has shape (n_samples, n_features).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
# predicting on FactorizationMachineRegressor model
fmr.predict(csr)
```

Output

```
[10.00277618 20.6134937 13.81870647 26.75876098 33.13530746 3.03828379]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "csr" is scipy sparse data, we have created FrovedisCRSMatrix.
from frovedis.matrix.crs import FrovedisCRSMatrix
cmat = FrovedisCRSMatrix(csr)

# predicting on FactorizationMachineRegressor model using pre-constructed input
fmr.predict(cmat)
```

Output

```
[10.00277618 20.6134937 13.81870647 26.75876098 33.13530746 3.03828379]
```

Return Value

It returns a numpy array of double (float64) type containing the predicted outputs. It is of shape (n_samples,).

9.3.1.4 4. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

Currently, this method is not supported for FactorizationMachineRegressor. It is simply kept in FactorizationMachineRegressor module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an AttributeError.

9.3.1.5 5. score(X, y, sample_weight = None)**Parameters**

X: A scipy sparse matrix or an instance of FrovedisCRSMMatrix of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the target values. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of self.predict(X) wrt. y.

The coefficient 'R2' is defined as $(1 - (u/v))$,

where 'u' is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and

'v' is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
fmr.score(csr, lbl)
```

Output

```
-0.10
```

Return Value

It returns an accuracy score of double (float64) type.

9.3.1.6 6. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information(metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the FactorizationMachineRegressor model
fmr.save("./out/FMRModel")
```

The FMRModel contains below directory structure:

FMRModel

```
|— metadata
|— model
```

'metadata' represents the detail about model_kind and datatype of training vector.

Here, the model file contains information about trained model in binary format.

This will save the FactorizationMachineRegressor model on the path '/out/FMRModel'. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

9.3.1.7 7. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by FactorizationMachineRegressor. It is used to get parameters and their values of FactorizationMachineRegressor class.

For example,

```
print(fmr.get_params())
```

Output

```
{'batch_size_pernode': 100, 'dim': (True, True, 8), 'init_learn_rate': 0.01,
'init_stddev': 0.1, 'iteration': 100, 'optimizer': 'SGD', 'reg': (False, False, 0),
'verbos': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

9.3.1.8 8. `set_params(**params)`

Parameters

****params:** All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by FactorizationMachineRegressor, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(fmr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
fmr.set_params(iteration=200)
print("Get parameters after setting:")
print(fmr.get_params())
```

Output

```
Get parameters before setting: {'batch_size_pernode': 100, 'dim': (True, True, 8),
'init_learn_rate': 0.01, 'init_stddev': 0.1, 'iteration': 100, 'optimizer': 'SGD',
'reg': (False, False, 0), 'verbos': 0}
Get parameters before setting: {'batch_size_pernode': 100, 'dim': (True, True, 8),
'init_learn_rate': 0.01, 'init_stddev': 0.1, 'iteration': 200, 'optimizer': 'SGD',
'reg': (0, 0, 0), 'verbos': 0}
```

Return Value

It simply returns “self” reference.

9.3.1.9 9. `debug_print()`

Purpose

Currently, this method is not supported for FactorizationMachineRegressor. It is simply kept in Factoriza-

tionMachineRegressor module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an `AttributeError`.

9.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
fmr.release()
```

This will reset the after-fit populated attributes to `None`, along with releasing server side memory.

Return Value

It returns nothing.

9.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, `predict()` is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns `'True'`, if the model is already fitted otherwise, it returns `'False'`.

9.4 SEE ALSO

- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Factorization Machine Classifier in Frovedis**

Chapter 10

FPGrowth

10.1 NAME

FPGrowth - A frequent pattern mining algorithm supported by Frovedis.

10.2 SYNOPSIS

```
class frovedis.mllib.fpm.FPGrowth(minSupport=0.3, minConfidence=0.8, itemsCol='items',
                                   predictionCol='prediction', numPartitions=None,
                                   tree_depth=None, compression_point=4, mem_opt_level=0,
                                   verbose=0, encode_string_input=False)
```

10.2.1 Public Member Functions

```
fit(data)
generate_rules(confidence = None)
transform(data)
load(fname)
save(fname)
debug_print()
release()
is_fitted()
```

10.3 DESCRIPTION

FPGrowth is an algorithm for discovering frequent itemsets in a transaction database. The input of FPGrowth is a set of transactions called transaction database. Each transaction is a set of items. **Frovedis supports numeric and non-numeric values for transaction data.**

For example, consider the following transaction database. It contains 4 transactions (t1, t2, t3, t4) and 4 items (1, 2, 3, 4). The first transaction represents the set of items 1, 2, 3 and 4.

Transaction id	Items
t1	{1, 2, 3, 4}
t2	{1, 2, 3}

```
t3          {1, 2}
t4          {1}
```

It is important to note that repetition of item in any given transaction needs to be avoided. It would raise exception in that case.

Now, if FPGrowth is run on the above transaction database with a minSupport of 40% and a tree_depth of 5 levels,

FPGrowth produces the following result:

items	freq
[1]	4
[2]	3
[3]	2
[2, 1]	3
[3, 1]	2
[3, 2]	2
[3, 2, 1]	2

In the results, each itemset is annotated with its corresponding frequency.

This module provides a client-server implementation, where the client application is a normal python program. In this implementation, a python client can interact with a frovedis server by sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for FPGrowth on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

10.3.1 Detailed Description

10.3.1.1 1. FPGrowth()

Parameters

minSupport: A positive double (float64) type value that specifies the minimum support level of frequent itemsets. Its value must be within 0 to 1. (Default: 0.3)

minConfidence: A positive double (float64) type value that specifies the minimal confidence for generating association rules. It will not affect the mining for frequent itemsets, but will affect the association rules generation. Its value must be within 0 to 1. (Default: 0.8)

itemsCol: An unused parameter. (Default: 'items')

predictionCol: An unused parameter. (Default: 'prediction')

numPartitions: An unused parameter. (Default: None)

tree_depth: A positive integer parameter specifying the maximum number of levels for tree construction. Its value must be greater than 1. (Default: None)

When it is None (not specified explicitly), the tree is constructed to its maximum depth according to the data. Since transaction databases tend to be very large, there may be a scenario wherein entire FP tree cannot be contained in memory. In those cases, the size of FP tree may be limited by using this parameter.

compression_point: A positive integer parameter. This is an internal memory optimisation strategy which helps when working with large transaction databases. Its value must be greater than or equal to 2. No compression will be performed till the level specified by this parameter is reached. (Default: 4)

mem_opt_level: An integer value which must be either 0 (memory optimisation OFF) or 1 (memory optimisation ON). If switched On, it will lead to removal of redundant tree data residing in memory at server

side. It should only be used where systems have memory constraints. By default, it is 0, but in case of memory constraints, execution should be attempted keeping this value as 1, it will help in reducing memory footprint. However, when it is 1, it might still cause memory issue in case data is too big. In this case, data may be spilled onto disk if this environment variable has been set. This will degrade performance (execution time).

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

encode_string_input: A boolean parameter when set to True, encodes the non-numeric (like strings) itemset values. It first internally encodes the named-items to an encoded numbered-items and the encoded dataframe is used for further training at frovedis server. It helps to train the encoded fpgrowth model faster with non-numeric itemsets since data present with server is two column dataframe, operations like join(), etc are little on slower on non-numeric columns than the numeric columns. (Default: False)

For example,

```
# let the data be some non-numeric transaction database
data = [['banana','apple','mango','cake'],
        ['cake','banana','apple'],
        ['bread','banana'],
        ['banana']]

# Using FPGrowth object with memory optimization parameters for training
# Here, disabling the parameter encode_string_input = False by default
from frovedis.mllib.fpm import FPGrowth
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1)
fpm.fit(data)
```

Frequent itemsets generation time: 0.0361 sec

And, when enabling 'encode_string_input' with the same non-numeric data,

```
# Using FPGrowth object with memory optimization parameters for training
# Here, parameter encode_string_input = True
from frovedis.mllib.fpm import FPGrowth
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1, encode_string_input = True)
fpm.fit(data)
```

Frequent itemsets generation time: 0.0315 sec

Attributes

freqItemsets: A pandas dataframe having two fields, 'items' and 'freq', where 'items' is an array whereas 'freq' is double (float64) type value. It contains the itemsets along with their frequency values. Here, the frequency of an itemset signifies as to how many times the itemset appears in the transaction database.

associationRules: A pandas dataframe having six fields, 'antecedent', 'consequent', 'confidence', 'lift', 'support' and 'conviction'.

Every association rule is composed of two parts: an antecedent (if) and a consequent (then).

An 'antecedent' is an item found within the data. A 'consequent' is an item found in combination with the 'antecedent'. Both are itemsets (arrays).

For measuring the effectiveness of association rule, 'confidence', 'lift', 'support' and 'conviction' are used. All are double (float64) type values.

'confidence' refers to the amount of times a given rule turns out to be true in practice.

'support' is an indication of how frequently the itemset appears in the dataset.

'lift' is the ratio of confidence to support. If the rule has a lift of 1, it would imply that the probability of occurrence of the 'antecedent' and that of the 'consequent' are independent of each other. When two events are independent of each other, no rule can be drawn involving those two events. If the lift is greater than 1, that lets us know the degree to which those two occurrences are dependent on one another, and makes those

rules potentially useful for predicting the consequent in future data sets. If the lift is less than 1, that lets us know the items are substitute to each other. This means that presence of one item has negative effect on presence of other item and vice versa.

‘conviction’ compares the probability that X appears without Y if they were dependent with the actual frequency of the appearance of X without Y. If it equals 1, then they are completely unrelated.

count: A positive integer value which specifies the frequent itemsets count.

encode_logic: A python dictionary having transaction items with a corresponding encoded number as key-value pairs. It is only available when transaction items are have string values and ‘encode_string_input’ parameter is set to True. Otherwise it is None. This is used internally to perform auto decoding of items during ‘freqItemsets’ construction.

For example,

```
# let data be some non-numeric transaction database
data = [['banana', 'apple', 'mango', 'cake'],
        ['cake', 'banana', 'apple'],
        ['bread', 'banana'],
        ['banana']]

# creating a pandas dataframe
import pandas as pd
dataDF = pd.DataFrame(data)

# Using FPGrowth object with memory optimization parameters and enabling encoding on non-numeric
# inputs and training
from frovedis.mllib.fpm import FPGrowth
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1, encode_string_input = True)
fpm.fit(data)
print("logic: ", fpm.encode_logic)
```

Output,

```
logic: {1: 'apple', 2: 'banana', 3: 'bread', 4: 'cake', 5: 'mango'}
```

Purpose

It initializes an FPGrowth object with the given parameters.

The parameters: “itemsCol”, “predictionCol” and “numPartitions” are simply kept in to make the interface uniform to the PySpark FPGrowth module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

10.3.1.2 2. fit(data)

Parameters

data: A python iterable or a pandas dataframe (manually constructed or loaded from file) or frovedis-two column dataframe containing the transaction data. Frovedis supports numeric and non-numeric values in transaction dataset.

Purpose

It accepts the training data and trains the fpgrowth model with specified minimum support value and tree depth value for construction of tree.

For pandas dataframe, if it has 2 columns, the second column would be treated as items and needs to be an array-like input.

For example,

```
# creating a pandas dataframe
import pandas as pd
data = [['banana', 'apple', 'mango', 'cake'],
        ['cake', 'banana', 'apple'],
        ['bread', 'banana'],
        ['banana']]
dataDF = pd.DataFrame(data)

# fitting input dataframe on FPGrowth object
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1).fit(dataDF)
# Now, to print the frequent itemsets table after training is completed
print(fpm.freqItemsets)
# to print table with all the association rules
print(fpm.associationRules)
```

Output,

frequent itemsets:

	items	freq
0	[banana]	4.0
1	[apple]	2.0
2	[cake]	2.0
3	[bread]	1.0
4	[mango]	1.0
5	[cake, apple]	2.0
6	[mango, apple]	1.0
7	[apple, banana]	2.0
8	[bread, banana]	1.0
9	[cake, banana]	2.0
10	[mango, banana]	1.0
...		
16	[mango, cake, apple, banana]	1.0

association rules:

	antecedent	consequent	confidence	lift	support	conviction
0	[banana]	[apple]	0.5	1.0	0.50	1.0
1	[apple]	[cake]	1.0	2.0	0.50	NaN
2	[banana]	[cake]	0.5	1.0	0.50	1.0
3	[apple]	[mango]	0.5	2.0	0.25	1.5
4	[cake]	[mango]	0.5	2.0	0.25	1.5
5	[apple]	[banana]	1.0	1.0	0.50	NaN
6	[bread]	[banana]	1.0	1.0	0.25	NaN
7	[cake]	[apple]	1.0	2.0	0.50	NaN
8	[cake]	[banana]	1.0	1.0	0.50	NaN
9	[mango]	[apple]	1.0	2.0	0.25	NaN
10	[mango]	[banana]	1.0	1.0	0.25	NaN
...						
27	[mango, cake, apple]	[banana]	1.0	1.0	0.25	NaN

If it has more than 2 columns, all columns would be treated as individual item and missing items in any given transaction needs to be NaN.

For example,

FILE: groceries.csv

```

Item(s),Item 1,Item 2,Item 3,Item 4,Item 5,Item 6,Item 7, ... , Item 30,Item 31,Item 32
4,citrus fruit,semi-finished bread,margarine,ready soups,,,,,,,,,,,,,,,,,,,,,
3,tropical fruit,yogurt,coffee,,,,,,,,,,,,,,,,,,,,,
1,whole milk,,,,,,,,,,,,,,,,,,,,,
4,pip fruit,yogurt,cream cheese,meat spreads,,,,,,,,,,,,,,,,,,,,,
...
10,chicken,citrus fruit,other vegetables,butter,yogurt, ..., cling film/bags,,,,,,,,,,,,,
4,semi-finished bread,bottled water,soda,bottled beer,,,,,,,,,,,,,,,,,,,,,
5,chicken,tropical fruit,other vegetables,vinegar,shopping bags,,,,,,,,,,,,,,,,,,,,,

# reading data from csv file
import pandas as pd
dataDF = pd.read_csv("./input/groceries.csv", dtype=str).drop(['Item(s)'], axis=1)

# fitting input dataframe on FPGrowth object
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1).fit(dataDF)
# Now, to print the frequent itemsets table after training is completed
print(fpm.freqItemsets)
# to print table with all the association rules
print(fpm.associationRules)

```

Output

frequent itemsets:

	items	freq
0	[whole milk]	2513.0
1	[other vegetables]	1903.0
2	[rolls/buns]	1809.0
3	[soda]	1715.0
4	[yogurt]	1372.0
..
328	[whipped/sour cream, yogurt, whole milk]	107.0
329	[yogurt, other vegetables, whole milk]	219.0
330	[yogurt, rolls/buns, other vegetables]	113.0
331	[yogurt, rolls/buns, whole milk]	153.0
332	[yogurt, soda, whole milk]	103.0

association rules:

	antecedent	consequent	confidence	lift	support	conviction
0	[yogurt, other vegetables]	[whole milk]	0.512881	2.007235	0.022267	1.528340
1	[whipped/sour cream, yogurt]	[whole milk]	0.524510	2.052747	0.010880	1.565719
2	[whipped/sour cream, other vegetables]	[whole milk]	0.507042	1.984385	0.014642	1.510239
3	[tropical fruit, yogurt]	[whole milk]	0.517361	2.024770	0.015150	1.542528
4	[tropical fruit, root vegetables]	[whole milk]	0.570048	2.230969	0.011998	1.731553
5	[root vegetables, yogurt]	[whole milk]	0.562992	2.203354	0.014540	1.703594
6	[root vegetables, rolls/buns]	[whole milk]	0.523013	2.046888	0.012710	1.560804
7	[pip fruit, other vegetables]	[whole milk]	0.517510	2.025351	0.013523	1.543003
8	[domestic eggs, other vegetables]	[whole milk]	0.552511	2.162336	0.012303	1.663694
9	[curd, yogurt]	[whole milk]	0.582353	2.279125	0.010066	1.782567
10	[butter, other vegetables]	[whole milk]	0.573604	2.244885	0.011490	1.745992
11	[tropical fruit, root vegetables]	[other vegetables]	0.584541	3.020999	0.012303	1.941244
12	[root vegetables, yogurt]	[other vegetables]	0.500000	2.584078	0.012913	1.613015
13	[root vegetables, rolls/buns]	[other vegetables]	0.502092	2.594890	0.012201	1.619792
14	[citrus fruit, root vegetables]	[other vegetables]	0.586207	3.029608	0.010371	1.949059

When native python iterable or a pandas dataframe is provided, it is converted to frovedis dataframe and sent

to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions. The storage representation when creating frovedis dataframe is slightly different than pandas dataframe as shown below :

```
# creating a pandas dataframe
import pandas as pd
data = [['banana','apple','mango','cake'],
        ['cake','banana','apple'],
        ['bread','banana'],
        ['banana']]
dataDF = pd.DataFrame(data)

# Creating a two column frovedis dataframe
import frovedis.dataframe as fpd
frovDF = fpd.DataFrame(dataDF)

# fitting frovedis-like input dataframe on FPGrowth object
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1).fit(frovDF)
# Now, to print the frequent itemsets table after training is completed
print(fpm.freqItemsets)
# to print table with all the association rules
print(fpm.associationRules)
```

Output,

frequent itemsets:

	items	freq
0	[banana]	4.0
1	[apple]	2.0
2	[cake]	2.0
3	[bread]	1.0
4	[mango]	1.0
5	[cake, apple]	2.0
6	[mango, apple]	1.0
7	[apple, banana]	2.0
8	[bread, banana]	1.0
9	[cake, banana]	2.0
10	[mango, banana]	1.0

...

16 [mango, cake, apple, banana] 1.0

association rules:

	antecedent	consequent	confidence	lift	support	conviction
0	[banana]	[apple]	0.5	1.0	0.50	1.0
1	[apple]	[cake]	1.0	2.0	0.50	NaN
2	[banana]	[cake]	0.5	1.0	0.50	1.0
3	[apple]	[mango]	0.5	2.0	0.25	1.5
4	[cake]	[mango]	0.5	2.0	0.25	1.5
5	[apple]	[banana]	1.0	1.0	0.50	NaN
6	[bread]	[banana]	1.0	1.0	0.25	NaN
7	[cake]	[apple]	1.0	2.0	0.50	NaN
8	[cake]	[banana]	1.0	1.0	0.50	NaN
9	[mango]	[apple]	1.0	2.0	0.25	NaN
10	[mango]	[banana]	1.0	1.0	0.25	NaN

...

```
27      [mango, cake, apple]      [banana]          1.0    1.0    0.25      NaN
```

Another way to work with frovedis dataframe is by loading it from a file. Since in real world, large transaction database will be used for rule mining, such files needs to have two columns named “trans_id” and “item”, where “trans_id” column will have respective transaction id and “item” column will have the individual items in that transaction. If other names are present for the columns in the file, then it would raise exception.

For example:

FILE: trans.csv

```
trans_id,item 1,1
1,2
1,3
1,4
2,1
2,2
2,3
3,1
3,2
4,1
```

The above data can be loaded and passed to frovedis FPGrowth as follows:

```
# loadinf frovedis dataframe from a csv file
import frovedis.dataframe as fdf
frovDF = fdf.read_csv("trans.csv")

# fitting frovedis-like input dataframe on FPGrowth object
fpm = FPGrowth(minSupport = 0.01, minConfidence = 0.5, compression_point = 4,
               mem_opt_level = 1).fit(frovDF)
```

Return Value

It simply returns “self” reference.

10.3.1.3 3. generate_rules(confidence = None)

Parameters

confidence: A double (float64) type parameter indicating the minimum confidence value. (Default: None) When it is None (not specified explicitly), then it will use confidence value used during FPGrowth object creation.

Purpose

It accepts the minimum confidence value to trim the rules during the generation of association rules.

For example,

```
# generating rules with minimum confidence value of 0.2
fp_rules = fpm.generate_rules(0.2)
```

This will generate tables containing rules at server side.

To print these generated rules, debug_print() may be used as show below:

```
fp_rules.debug_print()
```

This will show all tables of different antecedent length at server side. Here, encoding was disabled during rule generation.

Output,

--- rule[0] ---

antecedent1	consequent	confidence	lift	support	conviction
apple	cake	1	2	0.5	NULL
apple	mango	0.5	2	0.25	1.5
banana	apple	0.5	1	0.5	1
banana	cake	0.5	1	0.5	1
cake	mango	0.5	2	0.25	1.5

--- rule[1] ---

antecedent1	consequent	confidence	lift	support	conviction
cake	apple	1	2	0.5	NULL
mango	apple	1	2	0.25	NULL
apple	banana	1	1	0.5	NULL
bread	banana	1	1	0.25	NULL
cake	banana	1	1	0.5	NULL
mango	banana	1	1	0.25	NULL
mango	cake	1	2	0.25	NULL

--- rule[2] ---

antecedent1	antecedent2	consequent	confidence	lift	support	conviction
apple	banana	cake	1	2	0.5	NULL
cake	banana	mango	0.5	2	0.25	1.5
apple	banana	mango	0.5	2	0.25	1.5
cake	apple	mango	0.5	2	0.25	1.5

--- rule[3] ---

antecedent1	antecedent2	consequent	confidence	lift	support	conviction
mango	banana	apple	1	2	0.25	NULL
cake	banana	apple	1	2	0.5	NULL
mango	banana	cake	1	2	0.25	NULL
mango	apple	cake	1	2	0.25	NULL

--- rule[4] ---

antecedent1	antecedent2	consequent	confidence	lift	support	conviction
mango	cake	banana	1	1	0.25	NULL
mango	apple	banana	1	1	0.25	NULL
cake	apple	banana	1	1	0.5	NULL
mango	cake	apple	1	2	0.25	NULL

--- rule[5] ---

antecedent1	antecedent2	antecedent3	consequent	confidence	lift	support	conviction
cake	apple	banana	mango	0.5	2	0.25	1.5

--- rule[6] ---

antecedent1	antecedent2	antecedent3	consequent	confidence	lift	support	conviction
mango	apple	banana	cake	1	2	0.25	NULL

--- rule[7] ---

antecedent1	antecedent2	antecedent3	consequent	confidence	lift	support	conviction
mango	cake	banana	apple	1	2	0.25	NULL

--- rule[8] ---

antecedent1	antecedent2	antecedent3	consequent	confidence	lift	support	conviction
mango	cake	apple	banana	1	1	0.25	NULL

This output will be visible on server side. No such output will be visible on client side.

These above generated rules can also be saved and loaded separately as shown below:

```
rule.save("./out/FPRule")
```

It saves the rules in 'FPRule' directory.

It would raise exception if the directory already exists with same name.

The 'FPRule' directory has

FPRule

```
|—— encode_logic
|—— rule_0
|—— rule_1
|—— rule_2
|—— rule_3
|—— rule_4
|—— rule_5
|—— rule_6
|—— rule_7
|—— rule_8
```

The 'encode_logic' file is created only when 'encode_string_input = True' while training.

Other directories are created according to number of rules created which were constructed during training.

Each rule based directory contains information about antecedent, consequent, confidence, lift, support, conviction.

For loading the already saved rules, following should be done:

```
rule.load("./out/FPRule")
```

Return Value

It returns an FPRules object.

10.3.1.4 4. transform(data)

Parameters

data: A python iterable or a pandas dataframe or frovedis-like dataframe containing the transaction data. Frovedis supports numeric and non-numeric values as for transaction data.

Purpose

It gives the prediction of list of items for each of the corresponding transaction items.

For example,

```
print(fpm.transform(data))
```

Output

	items	prediction
0	[banana, apple, mango, cake]	[]
1	[cake, banana, apple]	[mango]
2	[bread, banana]	[apple, cake]
3	[banana]	[apple, cake]

In case no prediction is made for any list of items, then it will give an empty list, just like in pandas dataframe as well.

Return Value

It returns a pandas dataframe having two fields ‘items’ and ‘prediction’. Here ‘items’ is a list of transaction items and ‘prediction’ is a corresponding list of predictions for these transaction items.

10.3.1.5 5. load(fname)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

Purpose

It loads the model from the specified file path.

For example,

```
fpm.load("./out/FPModel")
```

Return Value

It simply returns “self” reference.

10.3.1.6 6. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information in the specified file path. Otherwise, it throws an exception.

For example,

```
# To save the FPGrowth model
fpm.save("./out/FPModel")
```

This will save the fp-growth model on the path ‘/out/FPModel’.

It would raise exception if the directory already exists with same name.

The ‘FPModel’ directory has

FPModel

```
|—— encode_logic
|—— metadata
|—— tree_0
|—— tree_1
|—— tree_2
|—— tree_3
```

The ‘encode_logic’ file is created only when ‘encode_string_input = True’ while training.

The metadata file contains the number of transactions and rule frequent itemsets count.

Rest of the directories are created according to tree levels which were constructed during training.

Each tree based directory contains information about items and their frequency count on each tree level.

Return Value

It returns nothing

10.3.1.7 7. debug_print()**Purpose**

It shows the target model information (frequent itemsets, generated fprules, frequent itemsets count) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
fpm.debug_print()
```

Output

```
--- item_support ---
item    item_support
banana  1
apple   0.5
cake    0.5
bread   0.25
mango   0.25

--- tree[0] ---
item    count
banana  4
apple   2
cake    2
bread   1
mango   1

--- tree_info[0] ---

--- tree[1] ---
item    item1  count
cake    apple  2
mango   apple  1
apple   banana  2
bread   banana  1
cake    banana  2
mango   banana  1
mango   cake   1

--- tree_info[1] ---

--- tree[2] ---
item    item1  item2  count
mango   cake   apple  1
cake    apple  banana  2
mango   apple  banana  1
mango   cake   banana  1

--- tree_info[2] ---

--- tree[3] ---
item    item1  item2  item3  count
mango   cake   apple  banana  1

--- tree_info[3] ---

total #FIS: 17
```

This output will be visible on server side. It displays the in memory frequent itemsets, generated fprules, frequent itemsets count which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing

10.3.1.8 8. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
fpm.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing

10.3.1.9 9. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

10.4 SEE ALSO

- Introduction to frovedis DataFrame

Chapter 11

GaussianMixture

11.1 NAME

GaussianMixture - It is a representation of a Gaussian mixture model probability distribution. This class allows to estimate the parameters of a Gaussian mixture distribution.

11.2 SYNOPSIS

```
class frovedis.mllib.mixture.gmm.GaussianMixture(n_components=1, covariance_type='full',
                                                  tol=1e-3, reg_covar=1e-6, max_iter=100,
                                                  n_init=1, init_params='kmeans',
                                                  weights_init=None, means_init=None,
                                                  precisions_init=None, random_state=None,
                                                  warm_start=False, verbose=0,
                                                  verbose_interval=10)
```

11.2.1 Public Member Functions

```
fit(X, y = None)
fit__predict(X, y = None)
predict(X)
predict__proba(X)
sample(n_samples = 1)
score(X, y = None)
score__samples(X)
get__params(deep = True)
set__params(**params)
load(fname, dtype = None)
save(fname)
bic(X)
aic(X)
debug__print()
release()
is__fitted()
```

11.3 DESCRIPTION

A Gaussian mixture model is a probabilistic model that assumes all the data points are generated from a mixture of a finite number of Gaussian distributions with unknown parameters. One can think of mixture models as generalizing k-means clustering to incorporate information about the covariance structure of the data as well as the centers of the latent Gaussians.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn GaussianMixture interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for GaussianMixture on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, the python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

11.3.1 Detailed Description

11.3.1.1 1. GaussianMixture()

Parameters

n_components: A positive integer parameter specifying the number of mixture components. (Default: 1)

covariance_type: A string object parameter specifying the type of covariance parameters to use.

Currently, 'full' is supported in frovedis. In 'full' covariance type, each component has its own general covariance matrix. (Default: 'full')

tol: Zero or a positive double (float64) parameter specifying the convergence tolerance. EM (expectation-maximization) iterations will stop when the lower bound average gain is below this threshold. (Default: 1e-3)

reg_covar: An unused parameter. (Default: 1e-6)

max_iter: A positive integer parameter specifying the number of EM (expectation-maximization) iterations to perform. (Default: 100)

n_init: A positive integer parameter specifying the number of initializations to perform. (Default: 1)

If it is None (not specified explicitly), it will be set as 1.

init_params: A string object parameter specifying the method used to initialize the weights, the means and the precisions. (Default: 'kmeans')

Must be one of the following:

- 'kmeans' : responsibilities are initialized using kmeans.
- 'random' : responsibilities are initialized randomly.

weights_init: An unused parameter. (Default: None)

means_init: An unused parameter. (Default: None)

precisions_init: An unused parameter. (Default: None)

random_state: An integer, float parameter that controls the random seed given to the method chosen to initialize the parameters. It does not support RandomState instance. (Default: None)

If it is None (not specified explicitly) or a RandomState instance, it will be set as 0.

warm_start: An unused parameter. (Default: False)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

verbose_interval: An unused parameter. (Default: 10)

Attribute

weights_: It is a python ndarray, containing double (float64) typed values and has shape (**n_components**,). It stores the weights of each mixture components.

covariances_: It is a python ndarray, containing double (float64) typed values. It stores the covariance of each mixture component. The shape depends on **covariance_type**:

- if ‘full’, then the shape is (**n_components**, **n_features**, **n_features**).

means_: It is a python ndarray, containing double (float64) typed values and has shape (**n_components**, **n_features**). It stores the mean of each mixture component.

converged_: A boolean value. If True, then convergence was reached in fit(), otherwise it will be False.

n_iter_int: An integer value specifying the number of step used by the best fit of EM to reach the convergence.

lower_bound_: A float value specifying the lower bound value on the log-likelihood (of the training data with respect to the model) of the best fit of EM.

Purpose

It initializes a GaussianMixture object with the given parameters.

The parameters: “reg_covar”, “weights_init”, “means_init”, “precisions_init”, “warm_start”, and “verbose_interval” are simply kept to make the interface uniform to Scikit-learn GaussianMixture module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

11.3.1.2 2. fit(X, y = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). During training, **n_samples** \geq **n_components**.

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It estimates the model parameters with the EM algorithm.

The method fits the model ‘**n_init**’ times and sets the parameters with which the model has the largest likelihood or lower bound. Within each trial, the method iterates between E-step and M-step for ‘**max_iter**’ times until the change of likelihood or lower bound is less than ‘**tol**’.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
                        [1., 4.],
                        [1., 0.],
                        [10., 2.],
                        [10., 4.],
                        [10., 0.]])

# fitting input matrix on GaussianMixture object
from frovedis.mllib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2)
gmm_model.fit(train_mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training

time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
#                       [1., 4.],
#                       [1., 0.],
#                       [10., 2.],
#                       [10., 4.],
#                       [10., 0.]])

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# fitting GaussianMixture object with pre-constructed input
from frovedis.ml.lib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2)
gmm_model.fit(rmat)
```

Return Value

It simply returns “self” reference.

11.3.1.3 3. fit_predict(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). During training, **n_samples** \geq **n_components**.

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It estimates the model parameters using X and predict the labels for X.

The method fits the model ‘**n_init**’ times and sets the parameters with which the model has the largest likelihood or lower bound. Within each trial, the method iterates between E-step and M-step for ‘**max_iter**’ times until the change of likelihood or lower bound is less than ‘**tol**’. After fitting, it predicts the most probable label for the input data points.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
#                       [1., 4.],
#                       [1., 0.],
#                       [10., 2.],
#                       [10., 4.],
#                       [10., 0.]])

# fitting input matrix on GaussianMixture object and perform predictions
from frovedis.ml.lib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2)
print(gmm_model.fit_predict(train_mat))
```

Output

```
[0 0 0 1 1 1]
```

Like in `fit()` frovedis-like input can be used to speed-up training at server side.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
                        [1., 4.],
                        [1., 0.],
                        [10., 2.],
                        [10., 4.],
                        [10., 0.]])

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# fitting GaussianMixture object with pre-constructed input and perform predictions
from frovedis.ml.lib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2)
print(gmm_model.fit_predict(rmat))
```

Output

```
[0 0 0 1 1 1]
```

Return Value

It returns a numpy array of int64 type values containing the components labels. It has a shape **(n_samples,)**.

11.3.1.4 4. `predict(X)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data. It has shape **(n_samples, n_features)**.

Purpose

It predict the labels for the data samples in X using trained model.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
                        [1., 4.],
                        [1., 0.],
                        [10., 2.],
                        [10., 4.],
                        [10., 0.]])

# predicting on GaussianMixture model
from frovedis.ml.lib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2).fit(train_mat)
print(gmm_model.predict(train_mat))
```

Output

```
[0 0 0 1 1 1]
```

Like in `fit()`, frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
                        [1., 4.],
                        [1., 0.],
                        [10., 2.],
                        [10., 4.],
                        [10., 0.]])

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# predicting on GaussianMixture model using pre-constructed input
from frovedis.mllib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2).fit(rmat)
print(gmm_model.predict(rmat))
```

Output

```
[0 0 0 1 1 1]
```

Return Value

It returns a numpy array of int64 type values containing the components labels. It has a shape **(n_samples,)**.

11.3.1.5 5. `predict_proba(X)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data. It has shape **(n_samples, n_features)**.

Purpose

It predict the labels for the data samples in X using trained model.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
                        [1., 4.],
                        [1., 0.],
                        [10., 2.],
                        [10., 4.],
                        [10., 0.]])

# predicting on GaussianMixture model
from frovedis.mllib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2).fit(train_mat)
print(gmm_model.predict_proba(train_mat))
```

Output

```
[[1. 0.]
```

```
[1. 0.]
[1. 0.]
[0. 1.]
[0. 1.]
[0. 1.]]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# loading sample matrix dense data
# train_mat = np.array([[1., 2.],
                        [1., 4.],
                        [1., 0.],
                        [10., 2.],
                        [10., 4.],
                        [10., 0.]])

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# predicting on GaussianMixture model using pre-constructed input
from frovedis.mllib.mixture import GaussianMixture
gmm_model = GaussianMixture(n_components = 2).fit(rmat)
print(gmm_model.predict_proba(rmat))
```

Output

```
[[1. 0.]
 [1. 0.]
 [1. 0.]
 [0. 1.]
 [0. 1.]
 [0. 1.]]
```

Return Value

It returns a numpy array of double (float64) type values containing the density of each gaussian component for each sample in `X`. It has a shape **(n_samples, n_components)**.

11.3.1.6 6. `sample(n_samples = 1)`

Parameters

n_samples: A positive integer value that specifies the number of samples to generate. (Default: 1)

Purpose

Currently this method is not supported for `GaussianMixture` in `frovedis`.

Return Value

It simply raises a `NotImplementedError`.

11.3.1.7 7. `score(X)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCSRMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data. It has shape **(n_samples, n_features)**.

Purpose

It computes the per-sample average log-likelihood of the given data X.

For example,

```
# calculate log-likelihood on given test data X
gmm_model.score(train_mat)
```

Output

```
3.386
```

Return Value

It returns a score of double (float64) type.

11.3.1.8 8. score_samples(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It computes the log-likelihood of each sample.

For example,

```
gmm_model.score_samples(train_mat)
```

Output

```
[3.88631622 3.1363165  3.1363165  3.88631622 3.1363165  3.1363165 ]
```

Return Value

It returns a numpy array of double (float64) type containing log-likelihood of each sample in 'X' under the current model. It has a shape (**n_samples**,).

11.3.1.9 9. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by GaussianMixture. It is used to get parameters and their values of GaussianMixture class.

For example,

```
print(gmm_model.get_params())
```

Output

```
{'covariance_type': 'full', 'init_params': 'kmeans', 'max_iter': 100, 'means_init': None,
'n_components': 2, 'n_init': 1, 'precisions_init': None, 'random_state': None,
'reg_covar': None, 'tol': 0.001, 'verbose': 0, 'verbose_interval': None, 'warm_start': None,
'weights_init': None}
```

Return Value

A dictionary of parameter names mapped to their values.

11.3.1.10 10. set_params(params)****Parameters**

****params:** All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by GaussianMixture, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(gmm_model.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
gmm_model.set_params(n_components = 4)
print("get parameters after setting:")
print(gmm_model.get_params())
```

Output

```
get parameters before setting:
{'covariance_type': 'full', 'init_params': 'kmeans', 'max_iter': 100, 'means_init': None,
'n_components': 2, 'n_init': 1, 'precisions_init': None, 'random_state': None,
'reg_covar': None, 'tol': 0.001, 'verbose': 0, 'verbose_interval': None, 'warm_start': None,
'weights_init': None}
get parameters after setting:
{'covariance_type': 'full', 'init_params': 'kmeans', 'max_iter': 100, 'means_init': None,
'n_components': 4, 'n_init': 1, 'precisions_init': None, 'random_state': None,
'reg_covar': None, 'tol': 0.001, 'verbose': 0, 'verbose_interval': None, 'warm_start': None,
'weights_init': None}
```

Return Value

It simply returns “self” reference.

11.3.1.11 11. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the same model
gmm_model.load("./out/MyGmmModel",dtype = np.float64)
```

Return Value

It simply returns “self” reference.

11.3.1.12 12. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
gmm_model.save("./out/MyGmmModel")
```

This will save the random forest classifier model on the path “/out/MyGmmModel”. It would raise exception if the directory already exists with same name.

The MyGmmModel contains below directory structure:

MyGmmModel

```
|— metadata
|— model
```

‘metadata’ represents the detail about `n_components`, `n_features`, `converged__`, `n_iter__`, `lower_bound__`, `model_kind` and datatype of training vector.

Here, the ‘model’ file contains information about gaussian mixture model in binary format.

Return Value

It returns nothing.

11.3.1.13 13. bic(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It is the Bayesian information criterion for the current model on the input X.

For example,

```
gmm_model.bic(train_mat)
```

Output

```
-20.926
```

Return Value

It returns a bayesian information criterion of double (float64) type.

11.3.1.14 14. aic(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It is the Akaike information criterion for the current model on the input X.

For example,

```
gmm_model.aic(train_mat)
```

Output

```
-18.636
```

Return Value

It returns an akaike information criterion of double (float64) type.

11.3.1.15 15. debug_print()**Purpose**

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
gmm_model.debug_print()
```

Output

Gaussian Mixture Model:

Means:

```
node = 0, local_num_row = 2, local_num_col = 2, val = 1 2 10 2
```

Weights:

```
node = 0, local_num_row = 2, local_num_col = 1, val = 0.5 0.5
```

Covariances:

```
node = 0, local_num_row = 2, local_num_col = 4, val = 0 0 0 2.66667 0 0 0 2.66667
```

This output will be visible on server side. It displays the information on the trained model such as means, weights, covariances which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

11.3.1.16 16. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
gmm_model.release()
```

This will reset the after-fit populated attributes (like weights_, means_, etc.) to None, along with releasing server side memory.

Return Value

It returns nothing.

11.3.1.17 17. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

11.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **KMeans in Frovedis**

Chapter 12

GradientBoostingClassifier

12.1 NAME

GradientBoostingClassifier - It is a machine learning algorithm, used for binary classification problems. It works on the principle that many weak learners (multiple decision trees) can together make a more accurate predictor.

12.2 SYNOPSIS

```
class frovedis.mllib.ensemble.gbtree.GradientBoostingClassifier(loss="deviance",
                                                                learning_rate=0.1,
                                                                n_estimators=100,
                                                                subsample=1.0,
                                                                criterion="friedman_mse",
                                                                min_samples_split=2,
                                                                min_samples_leaf=1,
                                                                min_weight_fraction_leaf=0.,
                                                                max_depth=3,
                                                                min_impurity_decrease=0.,
                                                                min_impurity_split=None,
                                                                init=None, random_state=None,
                                                                max_features=None, verbose=0,
                                                                max_leaf_nodes=None,
                                                                warm_start=False,
                                                                presort="deprecated",
                                                                validation_fraction=0.1,
                                                                n_iter_no_change=None,
                                                                tol=1e-4, ccp_alpha=0.0,
                                                                max_bins=32)
```

12.2.1 Public Member Functions

```
fit(X, y)
predict(X)
get_params(deep = True)
```

```

set_params(**params)
load(fname, dtype = None)
score(X, y, sample_weight = None)
save(fname)
debug_print()
release()
is_fitted()

```

12.3 DESCRIPTION

GradientBoostingClassifier is a supervised machine learning algorithm used for classification using decision trees. In gradient boosting decision trees, we combine many weak learners to come up with one strong learner. The weak learners here are the individual decision trees. All the trees are connected in series and each tree tries to minimise the error of the previous tree. Due to this sequential connection, boosting algorithms are usually slow to learn, but also highly accurate. **Currently, frovedis supports only binary gradient boosting classification algorithm.**

During training, the input **X** is the training data and **y** is the corresponding label values (Frovedis supports any values for labels, but internally it encodes the input binary labels to -1 and 1) before training at Frovedis server) which we want to predict.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn GradientBoostingClassifier interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for GradientBoostingClassifier on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

12.3.1 Detailed Description

12.3.1.1 1. GradientBoostingClassifier()

Parameters

loss: A string object parameter that specifies the loss function to be optimized. Currently, it supports only 'deviance' loss (logloss as in logistic regression) for classification. (Default: 'deviance')

learning_rate: A positive double (float64) parameter that shrinks the contribution of each tree by 'learning_rate' value provided. (Default: 0.1)

n_estimators: A positive integer parameter that specifies the number of boosting stages to perform. (Default: 100)

subsample: A positive double (float64) parameter that specifies the fraction of samples to be used for fitting the individual base learners. It must be in range (0, 1.0]. (Default: 1.0)

criterion: A string object parameter that specifies the function to measure the quality of a split. (Default: 'friedman_mse')

Currently, supported criteria are 'friedman_mse', 'mse' and 'mae'.

- 'friedman_mse': the mean squared error with improvement score by Friedman.

- ‘mse’: the mean squared error, which is equal to variance reduction as feature selection criterion.
- ‘mae’: the mean absolute error uses reduction in Poisson deviance to find splits.

min_samples_split: An unused parameter. (Default: 2)

min_samples_leaf: An unused parameter. (Default: 1)

min_weight_fraction_leaf: An unused parameter. (Default: 0.0)

max_depth: A positive integer parameter that specifies the maximum depth of the individual estimators. (Default: 3)

min_impurity_decrease: A positive double (float64) parameter. A node will be split if this split induces a decrease of the impurity greater than or equal to this value. (Default: 0.0)

min_impurity_split: An unused parameter. (Default: None)

init: An unused parameter. (Default: None)

random_state: An integer parameter that controls the random seed given to each tree estimator at each boosting iteration. In addition, it controls the random permutation of the features at each split. (Default: None)

If it is None (not specified explicitly), then ‘random_state’ is set as -1.

max_features: A string object parameter that specifies the number of features to consider when looking for the best split:

- If it is an integer, then it will be set as $(\text{max_features} * 1.0) / \text{n_features_}$.
- If it is float, then it will be ‘max_features’ number of features at each split.
- If it is ‘auto’, then it will be set as $\text{sqrt}(\text{n_features_})$.
- If ‘sqrt’, then it will be set as $\text{sqrt}(\text{n_features_})$ (same as ‘auto’).
- If ‘log2’, then it will be set as $\text{log2}(\text{n_features_})$.
- If None, then it will be set as n_features_ . (Default: ‘None’)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

max_leaf_nodes: An unused parameter. (Default: None)

warm_start: An unused parameter. (Default: False)

presort: An unused parameter. (Default: ‘deprecated’)

validation_fraction: An unused parameter. (Default: 0.1)

n_iter_no_change: An unused parameter. (Default: None)

tol: A double (float64) parameter that specifies the tolerance value for the early stopping. (Default: 1e-4)

ccp_alpha: An unused parameter. (Default: 0.0)

max_bins: A positive integer parameter that specifies the maximum number of bins created by ordered splits. (Default: 32)

Attributes

classes: It is a python ndarray (any type) of unique labels given to the classifier during training. It has shape (n_classes,).

n_estimators: An integer value specifying the ‘n_estimators’ value.

n_features: An integer value specifying the number of features when fitting the estimator.

Purpose

It initializes a GradientBoostingClassifier object with the given parameters.

The parameters: “min_samples_split”, “min_samples_leaf”, “min_weight_fraction_leaf”, “min_impurity_split”, “init”, “max_leaf_nodes”, “warm_start”, “presort”, “validation_fraction”, “n_iter_no_change” and “ccp_alpha” are simply kept in to make the interface uniform to the Scikit-learn GradientBoostingClassifier module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

12.3.1.2 2. fit(X, y)

Parameters

X: A numpy dense or any python array-like object or an instance of FrovedisColmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**,).

Purpose

It fits the gradient boosting model.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]])
lbl = np.array([1, 0, 1, 0])

# fitting input matrix and label on GradientBoostingClassifier object
from frovedis.mllib.ensemble import GradientBoostingClassifier
gbc = GradientBoostingClassifier(n_estimators = 2)
gbc.fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]])
lbl = np.array([1, 0, 1, 0])

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# fitting input matrix and label on GradientBoostingClassifier object
from frovedis.mllib.ensemble import GradientBoostingClassifier
gbc = GradientBoostingClassifier(n_estimators = 2)
gbc.fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

12.3.1.3 3. predict(X)**Parameters**

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

Purpose

Predict class for X.

For example,

```
# predicting on gradient boosting classifier model
gbc.predict(mat)
```

Output

```
[1 0 1 0]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on gradient boosting classifier model
gbc.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[1 0 1 0]
```

Return Value

It returns a numpy array of int64 type and of shape (**n_samples**,) containing the predicted classes.

12.3.1.4 4. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by GradientBoostingClassifier. It is used to get parameters and their values of GradientBoostingClassifier class.

For example,

```
print(gbc.get_params())
```

Output

```
{'ccp_alpha': 0.0, 'criterion': 'friedman_mse', 'init': None, 'learning_rate': 0.1,
'loss': 'deviance', 'max_bins': 32, 'max_depth': 3, 'max_features': None,
'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'n_estimators': 2, 'n_iter_no_change': None, 'presort': 'deprecated', 'random_state': -1,
'subsample': 1.0, 'tol': 0.0001, 'validation_fraction': 0.1, 'verbose': 0,
'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

12.3.1.5 5. `set_params(**params)`

Parameters

****params:** All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the `BaseEstimator` class inherited by `GradientBoostingClassifier`, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(gbc.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
gbc.set_params(n_estimators = 5)
print("get parameters after setting:")
print(gbc.get_params())
```

Output

```
get parameters before setting:
{'ccp_alpha': 0.0, 'criterion': 'friedman_mse', 'init': None, 'learning_rate': 0.1,
'loss': 'deviance', 'max_bins': 32, 'max_depth': 3, 'max_features': None,
'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'n_estimators': 2, 'n_iter_no_change': None, 'presort': 'deprecated', 'random_state': -1,
'subsample': 1.0, 'tol': 0.0001, 'validation_fraction': 0.1, 'verbose': 0,
'warm_start': False}
get parameters after setting:
{'ccp_alpha': 0.0, 'criterion': 'friedman_mse', 'init': None, 'learning_rate': 0.1,
'loss': 'deviance', 'max_bins': 32, 'max_depth': 3, 'max_features': None,
'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'n_estimators': 5, 'n_iter_no_change': None, 'presort': 'deprecated', 'random_state': -1,
'subsample': 1.0, 'tol': 0.0001, 'validation_fraction': 0.1, 'verbose': 0,
'warm_start': False}
```

Return Value

It simply returns “self” reference.

12.3.1.6 6. `load(fname, dtype = None)`

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
gbc.load("./out/gbt_classifier_model")
```

Return Value

It simply returns “self” reference.

12.3.1.7 7. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the gradient boosting classifier model
gbc.save("./out/gbt_classifier_model")
```

This will save the gradient boosted classifier model on the path “/out/gbt_classifier_model”. It would raise exception if the directory already exists with same name.

The ‘gbt_classifier_model’ directory has

gbt_classifier_model

```
|—label_map
|—metadata
|—model
```

‘label_map’ file contains information about labels mapped with their encoded value.

The ‘metadata’ file contains the number of classes, model kind and input datatype used for trained model.

The ‘model’ file contains the gradient boosting model saved in binary format.

Return Value

It returns nothing.

12.3.1.8 8. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object containing the true labels for X. It has shape (**n_samples**).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of self.predict(X) wrt. y.

For example,

```
# calculate mean accuracy score on given test data and labels
gbc.score(mat, lbl)
```

Output

```
1.00
```

Return Value

It returns an accuracy score of float type.

12.3.1.9 9. debug_print()**Purpose**

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
gbc.debug_print()
```

Output

```
----- Classification Trees (GBTs):: -----
# of trees: 2
---- [0] ----
  # of nodes: 3, height: 1
  <1> Split: feature[3] < 0.25, IG: 1
    \_ (2) Predict: 1
    \_ (3) Predict: -1
---- [1] ----
  # of nodes: 3, height: 1
  <1> Split: feature[3] < 0.25, IG: 0.227349
    \_ (2) Predict: 0.476812
    \_ (3) Predict: -0.476812
```

This output will be visible on server side. It displays the gradient boosting tree having maximum depth of 4 and total 2 decision trees.

No such output will be visible on client side.

Return Value

It returns nothing.

12.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
gbc.release()
```

This will reset the after-fit populated attributes (like `classes_`, `n_features_`) to `None`, along with releasing server side memory.

Return Value

It returns nothing.

12.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

12.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Gradient Boosting Regressor in Frovedis**
- **Decision Tree Classifier in Frovedis**

Chapter 13

GradientBoostingRegressor

13.1 NAME

GradientBoostingRegressor - It is a machine learning algorithm, used for regression. It works on the principle that many weak learners (multiple decision trees) can together make a more accurate predictor.

13.2 SYNOPSIS

```
class frovedis.mllib.ensemble.gbtree.GradientBoostingRegressor(loss="ls",
                                                                learning_rate=0.1,
                                                                n_estimators=100,
                                                                subsample=1.0,
                                                                criterion="friedman_mse",
                                                                min_samples_split=2,
                                                                min_samples_leaf=1,
                                                                min_weight_fraction_leaf=0.,
                                                                max_depth=3,
                                                                min_impurity_decrease=0.,
                                                                min_impurity_split=None,
                                                                init=None, random_state=None,
                                                                max_features=None, alpha=0.9,
                                                                verbose=0,
                                                                max_leaf_nodes=None,
                                                                warm_start=False,
                                                                presort="deprecated",
                                                                validation_fraction=0.1,
                                                                n_iter_no_change=None,
                                                                tol=1e-4, ccp_alpha=0.0,
                                                                max_bins=32)
```

13.2.1 Public Member Functions

```
fit(X, y)
predict(X)
get_params(deep = True)
```

```

set_params(**params)
load(fname, dtype = None)
score(X, y, sample_weight = None)
save(fname)
debug_print()
release()
is_fitted()

```

13.3 DESCRIPTION

GradientBoostingRegressor is a supervised machine learning algorithm used for regression using decision trees. In gradient boosting decision trees, we combine many weak learners to come up with one strong learner. The weak learners here are the individual decision trees. All the trees are connected in series and each tree tries to minimise the error of the previous tree. Due to this sequential connection, boosting algorithms are usually slow to learn, but also highly accurate.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn GradientBoostingRegressor interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for GradientBoostingRegressor on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

13.3.1 Detailed Description

13.3.1.1 1. GradientBoostingRegressor()

Parameters

loss: A string object parameter that specifies the loss function to be optimized. Currently, it supports 'ls' and 'lad' loss. (Default: 'ls')

- 'ls': refers to least squares regression.

- 'lad' (**least absolute deviation**): it is a highly robust loss function solely based on order information of the input variables.

learning_rate: A positive double (float64) parameter that shrinks the contribution of each tree by 'learning_rate' value provided. (Default: 0.1)

n_estimators: A positive integer parameter that specifies the number of boosting stages to perform. (Default: 100)

subsample: A positive double (float64) parameter that specifies the fraction of samples to be used for fitting the individual base learners. It must be in range (0, 1.0]. (Default: 1.0)

criterion: A string object parameter that specifies the function to measure the quality of a split. (Default: 'friedman_mse')

Currently, supported criteria are 'friedman_mse', 'mse' and 'mae'.

- 'friedman_mse': the mean squared error with improvement score by Friedman.

- ‘mse’: the mean squared error, which is equal to variance reduction as feature selection criterion.
- ‘mae’: the mean absolute error uses reduction in Poisson deviance to find splits.

min_samples_split: An unused parameter. (Default: 2)

min_samples_leaf: An unused parameter. (Default: 1)

min_weight_fraction_leaf: An unused parameter. (Default: 0.0)

max_depth: A positive integer parameter that specifies the maximum depth of the individual regression estimators. It limits the number of nodes in the tree. (Default: 3)

min_impurity_decrease: A positive double (float64) parameter. A node will be split if this split induces a decrease of the impurity greater than or equal to this value. (Default: 0.0)

min_impurity_split: An unused parameter. (Default: None)

init: An unused parameter. (Default: None)

random_state: An integer parameter that controls the random seed given to each tree estimator at each boosting iteration. In addition, it controls the random permutation of the features at each split. (Default: None)

If it is None (not specified explicitly), then ‘random_state’ is set as -1.

max_features: A string object parameter that specifies the number of features to consider when looking for the best split:

- If it is an integer, then it will be set as $(\text{max_features} * 1.0) / \text{n_features_}$.
- If it is float, then it will be ‘max_features’ number of features at each split.
- If it is ‘auto’, then it will be set as $\text{sqrt}(\text{n_features_})$.
- If ‘sqrt’, then it will be set as $\text{sqrt}(\text{n_features_})$ (same as ‘auto’).
- If ‘log2’, then it will be set as $\text{log2}(\text{n_features_})$.
- If None, then it will be set as n_features_ . (Default: ‘None’)

alpha: An unused parameter. (Default: 0.9)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

max_leaf_nodes: An unused parameter. (Default: None)

warm_start: An unused parameter. (Default: False)

presort: An unused parameter. (Default: ‘deprecated’)

validation_fraction: An unused parameter. (Default: 0.1)

n_iter_no_change: An unused parameter. (Default: None)

tol: A double (float64) parameter that specifies the tolerance value for the early stopping. (Default: 1e-4)

ccp_alpha: An unused parameter. (Default: 0.0)

max_bins: A positive integer parameter that specifies the maximum number of bins created by ordered splits. (Default: 32)

Attributes

n_estimators: An integer value specifying the ‘n_estimators’ value.

n_features: An integer value specifying the number of features when fitting the estimator.

Purpose

It initializes a GradientBoostingRegressor object with the given parameters.

The parameters: “min_samples_split”, “min_samples_leaf”, “min_weight_fraction_leaf”, “min_impurity_split”, “init”, “alpha”, “max_leaf_nodes”, “warm_start”, “presort”, “validation_fraction”, “n_iter_no_change” and “ccp_alpha” are simply kept in to make the interface uniform to the Scikit-learn GradientBoostingRegressor module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

13.3.1.2 2. fit(X, y)

Parameters

X: A numpy dense or any python array-like object or an instance of FrovedisColmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**,).

Purpose

It fits the gradient boosting model.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]])
lbl = np.array([1.2, 0.3, 1.1, 1.9])

# fitting input matrix and label on GradientBoostingRegressor object
from frovedis.mllib.ensemble import GradientBoostingRegressor
gbr = GradientBoostingRegressor(n_estimators = 2)
gbr.fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]])
lbl = np.array([1.2, 0.3, 1.1, 1.9])

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# fitting input matrix and label on GradientBoostingRegressor object
from frovedis.mllib.ensemble import GradientBoostingRegressor
gbr = GradientBoostingRegressor(n_estimators = 2)
gbr.fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

13.3.1.3 3. predict(X)**Parameters**

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

Purpose

Predict regression target for X.

For example,

```
# predicting on gradient boosting regressor model
gbr.predict(mat)
```

Output

```
[1.2 0.3 1.1 1.9]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on gradient boosting regressor model
gbr.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[1.2 0.3 1.1 1.9]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**,) containing the predicted values.

13.3.1.4 4. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by GradientBoostingRegressor. It is used to get parameters and their values of GradientBoostingRegressor class.

For example,

```
print(gbr.get_params())
```

Output

```
{'alpha': 0.9, 'ccp_alpha': 0.0, 'criterion': 'friedman_mse', 'init': None,
'learning_rate': 0.1, 'loss': 'ls', 'max_bins': 32, 'max_depth': 3, 'max_features': None,
'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'n_estimators': 2, 'n_iter_no_change': None, 'presort': 'deprecated', 'random_state': -1,
'subsample': 1.0, 'tol': 0.0001, 'validation_fraction': 0.1, 'verbose': 0,
'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

13.3.1.5 5. `set_params(**params)`

Parameters

****params:** All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the `BaseEstimator` class inherited by `GradientBoostingRegressor`, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(gbr.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
gbr.set_params(n_estimators = 5)
print("get parameters after setting:")
print(gbr.get_params())
```

Output

```
get parameters before setting:
{'alpha': 0.9, 'ccp_alpha': 0.0, 'criterion': 'friedman_mse', 'init': None,
 'learning_rate': 0.1, 'loss': 'ls', 'max_bins': 32, 'max_depth': 3, 'max_features': None,
 'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
 'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 2, 'n_iter_no_change': None, 'presort': 'deprecated', 'random_state': -1,
 'subsample': 1.0, 'tol': 0.0001, 'validation_fraction': 0.1, 'verbose': 0,
 'warm_start': False}
get parameters after setting:
{'alpha': 0.9, 'ccp_alpha': 0.0, 'criterion': 'friedman_mse', 'init': None,
 'learning_rate': 0.1, 'loss': 'ls', 'max_bins': 32, 'max_depth': 3, 'max_features': None,
 'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
 'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 5, 'n_iter_no_change': None, 'presort': 'deprecated', 'random_state': -1,
 'subsample': 1.0, 'tol': 0.0001, 'validation_fraction': 0.1, 'verbose': 0,
 'warm_start': False}
```

Return Value

It simply returns “self” reference.

13.3.1.6 6. `load(fname, dtype = None)`

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
gbr.load("./out/gbt_regressor_model")
```

Return Value

It simply returns “self” reference.

13.3.1.7 7. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the gradient boosting regressor model
gbr.save("./out/gbt_regressor_model")
```

This will save the gradient boosted regressor model on the path “/out/gbt_regressor_model”. It would raise exception if the directory already exists with same name.

The ‘gbt_regressor_model’ directory has

```
gbt_regressor_model
|—metadata
|—model
```

The ‘metadata’ file contains the model kind and input datatype used for trained model.

The ‘model’ file contains the gradient boosting model saved in binary format.

Return Value

It returns nothing.

13.3.1.8 8. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object containing the true values for X. It has shape (**n_samples**).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. $R2(r\text{-squared})$ of `self.predict(X)` wrt. `y`.

The coefficient ‘R2’ is defined as $(1 - (u/v))$,

where ‘u’ is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and,

‘v’ is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of `y`, disregarding the input features, would get a R2 score of 0.0.

For example,

```
# calculate R2 score on given test data and labels
gbr.score(mat, lbl)
```

Output

```
1.00
```

Return Value

It returns an R2 score of float type.

13.3.1.9 9. debug_print()**Purpose**

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
gbr.debug_print()
```

Output

```
----- Regression Trees (GBTs):: -----
# of trees: 2
---- [0] ----
  # of nodes: 7, height: 2
  <1> Split: feature[1] < 0.25, IG: 0.180625
    \_ <2> Split: feature[0] < 5.5, IG: 0.1225
      | \_ (4) Predict: 1.9
      | \_ (5) Predict: 1.2
      \_ <3> Split: feature[3] < 0.5, IG: 0.16
        \_ (6) Predict: 1.1
        \_ (7) Predict: 0.3
---- [1] ----
  # of nodes: 1, height: 0
  (1) Predict: 0
```

This output will be visible on server side. It displays the gradient boosting tree having maximum depth of 4 and total 2 decision trees.

No such output will be visible on client side.

Return Value

It returns nothing.

13.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
gbr.release()
```

This will reset the after-fit populated attributes (like `n_features__`) to None, along with releasing server side memory.

Return Value

It returns nothing.

13.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns ‘True’, if the model is already fitted, otherwise, it returns ‘False’.

13.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Gradient Boosting Classifier in Frovedis**
- **Decision Tree Regressor in Frovedis**

Chapter 14

KMeans Clustering

14.1 NAME

KMeans - is a clustering algorithm commonly used in EDA (exploratory data analysis).

14.2 SYNOPSIS

```
class frovedis.mllib.cluster.KMeans(n_clusters=8, init='random', n_init=10, max_iter=300,
                                     tol=1e-4, precompute_distances='auto', verbose=0,
                                     random_state=None, copy_x=True, n_jobs=1,
                                     algorithm='auto', use_shrink=False)
```

14.2.1 Public Member Functions

```
fit(X, y = None, sample_weight = None)
fit_predict(X, y = None, sample_weight = None)
fit_transform(X, y = None, sample_weight = None)
transform(X)
predict(X, sample_weight = None)
score(X, y = None, sample_weight = None)
load(fname, dtype = None)
save(fname)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

14.3 DESCRIPTION

Clustering is an unsupervised learning problem whereby we aim to group subsets of entities with one another based on some notion of similarity. Kmeans is one of the most commonly used clustering algorithms that clusters the data points into a predefined number of clusters.

Under unsupervised learning, there are two clustering methods- ‘k-means’ and ‘k-means++’. The main difference between these two lies in the selection of the centroids (we assume centroid is the center of the cluster) around which the clustering takes place.

Frovedis supports only k-means clustering method (i.e. `init = ‘random’`) which will randomly initialize the data points called centroid. Further, each data point is clustered to its nearest centroid and after every iteration the centroid is updated for each cluster. This cycle continues for a given number of repetitions and after that we have our final clusters.

This module provides a client-server implementation, where the client application is a normal python program. Frovedis is almost same as Scikit-learn clustering module providing kmeans support, but it doesn’t have any dependency with Scikit-learn. It can be used simply even if the system doesn’t have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Kmeans on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, the python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

14.3.1 Detailed Description

14.3.1.1 1. KMeans()

Parameters

`n_clusters`: An integer parameter specifying the number of clusters. The number of clusters should be greater than zero and less than `n_samples`. (Default: 8)

When it is None (specified explicitly), then it will be set as `min(8, n_samples)`.

`init`: A string object parameter specifies the method of initialization. (Default: ‘random’)

Unlike Scikit-learn, currently it only supports ‘random’ initialization.

`n_init`: A positive integer specifying the number of times the kmeans algorithm will be run with different centroid seeds. (Default: 10)

When it is None (specified explicitly), then it will be set as default 10.

`max_iter`: A positive integer parameter specifying the maximum iteration count. (Default: 300)

`tol`: Zero or a positive double (float64) parameter specifying the convergence tolerance. (Default: 1e-4)

`precompute_distances`: A string object parameter. (unused)

`verbose`: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

`random_state`: A zero or positive integer parameter. When it is None (not specified explicitly), it will be set as 0. (unused)

`copy_x`: A boolean parameter. (unused)

`n_jobs`: An integer parameter. (unused)

`algorithm`: A string object parameter, specifies the kmeans algorithm to use. (Default: auto)

When it is ‘auto’, it will be set as ‘full’. Unlike Scikit-learn, currently it supports only ‘full’.

`use_shrink`: A boolean parameter applicable only for “sparse” input (X). When set to True for sparse input, it can improve training performance by reducing communication overhead across participating processes. (Default: False)

Attribute

`cluster_centers_`: It is a python ndarray, containing float or double (float64) typed values and has shape (`n_clusters`, `n_features`). These are the coordinates of cluster centers.

labels_: A python ndarray of int64 values and has shape (**n_clusters**,). It contains predicted cluster labels for each point.

inertia_: A float parameter specifies the sum of squared distances of samples to their closest cluster center, weighted by the sample weights if provided.

n_iter_: An integer parameter specifies the number of iterations to run.

Purpose

It initializes a Kmeans object with the given parameters.

The parameters: “precompute_distances”, “random_state”, “copy_x” and “n_jobs” are simply kept to make the interface uniform to Scikit-learn cluster module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

14.3.1.2 2. fit(X, y = None, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: An unused parameter whose default value is None. It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It clusters the given data points (X) into a predefined number of clusters (**n_clusters**).

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on kmeans object
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1).fit(train_mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# KMeans with pre-constructed frovedis-like inputs
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1).fit(rmat)
```

Return Value

It simply returns “self” reference.

14.3.1.3 3. fit_predict(X, y = None, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: An unused parameter whose default value is None. It is simply ignored in frovedis implementation.

Purpose

It clusters the given data points (X) into a predefined number of clusters (**n_clusters**) and predicts the cluster index for each sample.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on KMeans object
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1)
print(kmeans.fit_predict(train_mat))
```

Output

```
[0 0 1 1 1]
```

Like in fit() frovedis-like input can be used to speed-up training at server side.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1)
print(kmeans.fit_predict(rmat))
```

Output

```
[0 0 1 1 1]
```

Return Value

It returns a numpy array of int64 type containing the cluster labels. It has a shape (**n_samples**,).

14.3.1.4 4. fit_transform(X, y = None, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: An unused parameter whose default value is None and simply ignored in frovedis implementation.

Purpose

It computes clustering and transforms training data (X) to cluster-distance space.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on KMeans object
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1)
print(kmeans.fit_transform(train_mat))
```

Output

```
[[ 0.08660254 15.58845727]
 [ 0.08660254 15.41525219]
 [15.32864965  0.17320508]
 [15.50185473  0.         ]
 [15.67505981  0.17320508]]
```

If training data (X) is a numpy array or a scipy sparse matrix, it will return a new numpy dense array.

Like in fit() frovedis-like input can be used to speed-up training at server side.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("sample_data.txt")

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1)
# it returns a FrovedisRowmajorMatrix object
kmeans.fit_transform(rmat)
```

If training data (X) is a frovedis-like input, it will return a FrovedisRowmajorMatrix object.

Return Value

- **When training data is native-python data:**

Then the output will be a numpy array containing the transformed matrix.

- **When training data is frovedis-like data:**

Then the output will be a FrovedisRowmajorMatrix.

In both cases output would be of float or double (float64) type (depending upon input dtype) and of shape (**n_samples**, **n_clusters**).

Note that even if training data (X) is sparse, the output would typically be dense.

14.3.1.5 5. transform(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

Purpose

It transforms the test data (X) to a cluster-distance space.

For example,

```
# loading sample matrix data
test_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on kmeans object
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1).fit(test_mat)
print(kmeans.transform(test_mat))
```

Output

```
[[ 0.08660254 15.58845727]
 [ 0.08660254 15.41525219]
 [15.32864965  0.17320508]
 [15.50185473  0.         ]
 [15.67505981  0.17320508]]
```

If test data (X) is a numpy array or a scipy sparse matrix, it will return a new numpy dense array.

Like in fit() frovedis-like input can be used to speed-up training the test data at server side.

For example,

```
# loading sample matrix data
test_mat = np.loadtxt("sample_data.txt")

# Since "test_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
tr_mat = FrovedisRowmajorMatrix(test_mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1).fit(tr_mat)
kmeans.transform(tr_mat)
```

If test data (X) is a frovedis-like input, it will return FrovedisRowmajorMatrix object.

Return Value

- If native-python data is input:

Then it returns a numpy array containing the transformed matrix.

- If frovedis-like data is input:

Then it returns a FrovedisRowmajorMatrix.

In both cases output would be of float or double (float64) type (depending upon input dtype) and of shape (**n_samples**, **n_clusters**).

Note that even if test data (X) is sparse, the output would typically be dense.

14.3.1.6 6. `predict(X, sample_weight = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

sample_weight: None or any python array-like object containing the intended weights for each input samples. It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It accepts the test data points (X) and returns the closest cluster each sample in X belongs to.

For example,

```
# loading sample matrix data
test_mat = np.loadtxt("sample_data.txt")

# fitting input matrix on KMeans object
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1).fit(test_mat)
print(kmeans.predict(test_mat))
```

Output

```
[0 0 1 1 1]
```

Like in fit() frovedis-like input can be used to speed-up prediction on the test data at server side.

For example,

```
# loading sample matrix data
test_mat = np.loadtxt("sample_data.txt")

# Since "test_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
tr_mat = FrovedisRowmajorMatrix(test_mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import KMeans
kmeans = KMeans(n_clusters = 2, n_init = 1).fit(tr_mat)
print(kmeans.predict(tr_mat))
```

Output

```
[0 0 1 1 1]
```

Return Value

It returns a numpy array of int32 type containing the centroid values. It has a shape (**n_samples**,).

14.3.1.7 7. `score(X, y = None, sample_weight = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix

for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: None or any python array-like object containing the intended weights for each input samples. It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It is calculated as “-1.0 * inertia”, which an indication of how far the points are from the centroids. Bad scores will return a large negative number, whereas good scores return a value close to zero.

For example,

```
kmeans.score(test_mat)
```

Output

```
-0.07499999552965164
```

Return Value

It returns a score of float type.

14.3.1.8 8. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the same model
kmeans.load("./out/MyKMeansModel",dtype=np.float64)
```

Return Value

It simply returns “self” instance.

14.3.1.9 9. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information(metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
kmeans.save("./out/MyKMeansModel")
```

The MyKMeansModel contains below directory structure:

MyKMeansModel

```
|— metadata
|— model
```


‘metadata’ represents the detail about model_id, model_kind and datatype of training vector. Here, the **model** directory contains information about n_clusters_, n_features, model_kind and datatype of training vector.

This will save the Kmeans model on the path ‘/out/MyKMeansModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

14.3.1.10 10. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by Kmeans. It is used to get parameters and their values of Kmeans class.

For example,

```
print(kmeans.get_params())
```

Output

```
{'algorithm': 'auto', 'copy_x': True, 'init': 'random', 'max_iter': 300, 'n_clusters': 2, 'n_init': 1, 'n_jobs': 1, 'precompute_distances': 'auto', 'random_state': None, 'tol': 0.0001, 'use_shrink': False, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

14.3.1.11 11. set_params(**params)

Parameters

****params**: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by Kmeans, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(kmeans.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
kmeans.set_params(n_clusters = 4, n_init = 5)
print("Get parameters after setting:")
print(kmeans.get_params())
```

Output

```
Get parameters before setting:
{'algorithm': 'auto', 'copy_x': True, 'init': 'random', 'max_iter': 300, 'n_clusters': 2, 'n_init': 1, 'n_jobs': 1, 'precompute_distances': 'auto', 'random_state': None, 'tol': 0.0001, 'use_shrink': False, 'verbose': 0}
Get parameters after setting: {'algorithm': 'auto', 'copy_x': True,
```

```
'init': 'random', 'max_iter': 300, 'n_clusters': 4, 'n_init': 5,
'n_jobs': 1, 'precompute_distances': 'auto', 'random_state': None,
'tol': 0.0001, 'use_shrink': False, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

14.3.1.12 12. debug_print()

Purpose

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
kmeans.debug_print()
```

Output

```
centroid:
node = 0, local_num_row = 2, local_num_col = 3, val = 0.15 0.15 0.15 9.1 9.1 9.1
```

This output will be visible on server side. It displays the centroid information on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

14.3.1.13 13. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
kmeans.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

14.3.1.14 14. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the clustering model first.

Return Value

It returns ‘True’, if the model is already fitted otherwise, it returns ‘False’.

14.4 SEE ALSO

- Introduction to FrovedisRowmajorMatrix

- **Introduction to FrovedisCRSMatrix**
- **Agglomerative Clustering in Frovedis**
- **Spectral Clustering in Frovedis**
- **DBSCAN in Frovedis**

Chapter 15

KNeighborsClassifier

15.1 NAME

KNeighborsClassifier - Classifier implementing the k-nearest neighbors vote.

15.2 SYNOPSIS

```
class frovedis.mllib.neighbors.KNeighborsClassifier(n_neighbors=5, weights='uniform',
                                                    algorithm='auto', leaf_size=30, p=2,
                                                    metric='euclidean', metric_params=None,
                                                    n_jobs=None, verbose=0, chunk_size=1.0,
                                                    batch_fraction=None)
```

15.2.1 Public Member Functions

```
fit(X, y)
kneighbors(X = None, n_neighbors = None, return_distance = True)
kneighbors_graph(X = None, n_neighbors = None, mode = 'connectivity')
save(fname)
load(fname)
predict(X, save_proba = False)
predict_proba(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

15.3 DESCRIPTION

Neighbors-based classification is a type of instance-based learning or non-generalizing learning. It does not attempt to construct a general internal model, but simply stores instances of the training data. Classification is computed from a simple majority vote of the nearest neighbors of each point- a query point is assigned the

data class which has the most representatives within the nearest neighbors of the point. **Frovedis supports both binary and multinomial labels.**

The k-neighbors classification in KNeighborsClassifier is a commonly used technique. The optimal choice of the value is highly data-dependent. In general, a larger k suppresses the effects of noise, but makes the classification boundaries less distinct.

During training, the input **X** is the training data and **y** are their corresponding label values (Frovedis supports any values as for labels, but internally it encodes the input binary labels to -1 and 1, before training at Frovedis server) which we want to predict.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn KNeighborsClassifier interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for KNeighborsClassifier on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When kneighbors-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

15.3.1 Detailed Description

15.3.1.1 1. KNeighborsClassifier()

Parameters

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for 'kneighbors' queries. It must be within the range of 0 and n_samples. (Default: 5)

weights: An unused parameter. (Default: uniform)

algorithm: A string object parameter, specifying the algorithm used to compute the nearest neighbors. (Default: auto)

When it is 'auto', it will be set as 'brute'(brute-force search approach). Unlike Scikit-learn, currently it supports only 'brute'.

leaf_size: An unused parameter. (Default: 30)

p: An unused parameter. (Default: 2)

metric: A string object parameter specifying the distance metric to use for the tree. (Default: 'euclidean') Currently it only supports 'euclidean', 'seuclidean' and 'cosine' distance.

metric_params: An unused parameter. (Default: None)

n_jobs: An unused parameter. (Default: None)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

chunk_size: A positive float parameter specifying the amount of data (in megabytes) to be processed in one time. (Default: 1.0)

batch_fraction: A positive double (float64) parameter used to calculate the batches of specific size. These batches are used to construct the distance matrix. It must be within the range of 0.0 to 1.0. (Default: None) When it is None (not specified explicitly), it will be set as np.finfo(np.float64).max value.

Attributes

classes_: A numpy array of int64 type value that specifies unique labels given to the classifier during training. It has shape (n_samples,).

Purpose

It initializes a KNeighborsClassifier object with the given parameters.

The parameters: “weights”, “leaf_size”, “p”, “metric_params” and “n_jobs” are simply kept in to make the interface uniform to the Scikit-learn KNeighborsClassifier module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

15.3.1.2 2. fit(X, y)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (n_samples, n_features).

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape (n_samples,).

Purpose

It accepts the training matrix (X) with labels (y) and trains a KNeighborsClassifier model.

For example,

```
# loading sample data
samples = np.loadtxt("./input/knc_data.txt", dtype = np.float64)
lbl = [10, 10, 10, 20, 10, 20]

# fitting input data on KNeighborsClassifier object
from frovedis.mllib.neighbors import KNeighborsClassifier
knc = KNeighborsClassifier(n_neighbors = 3)
knc.fit(samples, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample data
samples = np.loadtxt("./input/knc_data.txt", dtype = np.float64)
lbl = [10, 10, 10, 20, 10, 20]

# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
rmat = FrovedisRowmajorMatrix(samples)
dlbl = FrovedisDvector(lbl)

# fitting input data on KNeighborsClassifier object
from frovedis.mllib.neighbors import KNeighborsClassifier
knc = KNeighborsClassifier(n_neighbors = 3)
knc.fit(rmat, dlbl)
```

Return Value

It simply returns “self” reference.

15.3.1.3 3. kneighbors(X = None, n_neighbors = None, return_distance = True)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (n_queries, n_features), where 'n_queries' is the number of rows in the test data. (Default: None)

When it is None (not specified explicitly), it will be training data (X) used as input in fit().

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for 'kneighbors' queries. It must be within the range of 0 and n_samples. (Default: None)

When it is None (not specified explicitly), it will be 'n_neighbors' value used during KNeighborsClassifier object creation.

return_distance: A boolean parameter specifying whether or not to return the distances. (Default: True) If set to False, it will not return distances. Then, only indices are returned by this method.

Purpose

Finds the k-Neighbors of a point and returns the indices of neighbors and distances to the neighbors of each point.

For example,

```
distances, indices = knc.kneighbors(samples)
print('distances')
print(distances)
print('indices')
print(indices)
```

Output

```
distances
[[0.          1.          2.23606798]
 [0.          1.          1.41421356]
 [0.          1.41421356  2.23606798]
 [0.          1.          2.23606798]
 [0.          1.          1.41421356]
 [0.          1.41421356  2.23606798]]
indices
[[0 1 2]
 [1 0 2]
 [2 1 0]
 [3 4 5]
 [4 3 5]
 [5 4 3]]
```

Like in fit(), frovedis-like input can be used to speed-up the computation of indices and distances at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# fitting input data on KNeighborsClassifier object
from frovedis.mllib.neighbors import KNeighborsClassifier
knc = KNeighborsClassifier(n_neighbors = 3)
distances, indices = knc.kneighbors(rmat)
```



```
# Here FrovedisRowmajorMatrix().debug_print() is used
print('distances')
distances.debug_print()
# Here FrovedisRowmajorMatrix().debug_print() is used
print('indices')
indices.debug_print()
```

Output

```
distances
matrix:
num_row = 6, num_col = 3
node 0
node = 0, local_num_row = 6, local_num_col = 3, val = 0 1 2.23607 0 1 1.41421 0 1.41421
2.23607 0 1 2.23607 0 1 1.41421 0 1.41421 2.23607
indices
matrix:
num_row = 6, num_col = 3
node 0
node = 0, local_num_row = 6, local_num_col = 3, val = 0 1 2 1 0 2 2 1 0 3 4 5 4 3 5 5 4 3
```

It returns distances and indices as FrovedisRowmajorMatrix objects.

Return Value

1. When test data and training data used by fitted model are python native input:

- **distances**: A numpy array of float or double (float64) type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data. It is only returned by `kneighbors()`, if `return_distance = True`.
- **indices**: A numpy array of int64 type values. It has shape (**n_queries**, **n_neighbors**).

2. When either test data or training data used by fitted model is frovedis-like input:

- **distances**: A FrovedisRowmajorMatrix of float or double (float64) type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data. It is only returned by `kneighbors()`, if `return_distance = True`.
- **indices**: A FrovedisRowmajorMatrix of int64 type values. It has shape (**n_queries**, **n_neighbors**).

15.3.1.4 4. `kneighbors_graph(X = None, n_neighbors = None, mode = 'connectivity')`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCSRMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_queries**, **n_features**), where 'n_queries' is the number of rows in the test data. (Default: None)

When it is None (not specified explicitly), it will be training data (X) used as input in `fit()`.

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for 'kneighbors' queries. It must be within the range of 0 and `n_samples`. (Default: None)

When it is None (not specified explicitly), it will be 'n_queries' value used during NearestNeighbors object creation.

mode: A string object parameter which can be either 'connectivity' or 'distance'. It specifies the type of returned matrix.

For 'connectivity', it will return the connectivity matrix with ones and zeros, whereas for 'distance', the edges are euclidean distance between points, type of distance depends on the selected 'metric' value in KNeighborsClassifier class. (Default: 'connectivity')

Purpose

It computes the (weighted) graph of k-Neighbors for points in X.

For example,

```
# Here 'mode = connectivity' by default
graph = knc.kneighbors_graph(samples)
print('kneighbors graph')
print(graph)
```

Output

```
kneighbors graph
(0, 0)      1.0
(0, 1)      1.0
(0, 2)      1.0
(1, 1)      1.0
(1, 0)      1.0
(1, 2)      1.0
(2, 2)      1.0
(2, 1)      1.0
(2, 0)      1.0
(3, 3)      1.0
(3, 4)      1.0
(3, 5)      1.0
(4, 4)      1.0
(4, 3)      1.0
(4, 5)      1.0
(5, 5)      1.0
(5, 4)      1.0
(5, 3)      1.0
```

For example, when mode = 'distance'

```
# Here 'mode = distance'
graph = knc.kneighbors_graph(samples, mode = 'distance')
print('kneighbors graph')
print(graph)
```

Output

```
(0, 0)      0.0
(0, 1)      1.0
(0, 2)      2.23606797749979
(1, 1)      0.0
(1, 0)      1.0
(1, 2)      1.4142135623730951
(2, 2)      0.0
(2, 1)      1.4142135623730951
(2, 0)      2.23606797749979
(3, 3)      0.0
(3, 4)      1.0
(3, 5)      2.23606797749979
(4, 4)      0.0
(4, 3)      1.0
(4, 5)      1.4142135623730951
(5, 5)      0.0
(5, 4)      1.4142135623730951
```

(5, 3) 2.23606797749979

Like in `fit()`, `frovedis`-like input can be used to speed-up the graph making at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# fitting input data on KNeighborsClassifier object
from frovedis.mllib.neighbors import KNeighborsClassifier
knc = KNeighborsClassifier(n_neighbors = 3)
# Here 'mode = connectivity' by default
graph = knc.kneighbors_graph(rmat)

# Here FrovedisCRSMatrix().debug_print() is used
print('graph')
graph.debug_print()
```

Output

```
matrix:
num_row = 6, num_col = 6
node 0
local_num_row = 6, local_num_col = 6
val : 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
idx : 0 1 2 1 0 2 2 1 0 3 4 5 4 3 5 4 3
off : 0 3 6 9 12 15 18
```

It returns a `FrovedisCRSMatrix` object.

Return Value

- **When test data and training data used by fitted model are python native input:**

It returns a scipy sparse matrix of float or double (float64) type values. It has shape (`n_queries`, `n_samples_fit`), where '`n_queries`' is the number of rows in the test data and '`n_samples_fit`' is the number of samples in the fitted data.

- **When either test data or training data used by fitted model is `frovedis`-like input:**

It returns a `FrovedisCRSMatrix` of float or double (float64) type values. It has shape (`n_queries`, `n_samples_fit`), where '`n_queries`' is the number of rows in the test data and '`n_samples_fit`' is the number of samples in the fitted data.

15.3.1.5 5. `save(fname)`

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

Currently, this method is not supported for `KNeighborsClassifier`. It is simply kept in `KNeighborsClassifier` module to maintain uniform interface like other estimators in `frovedis`.

Return Value

It simply raises an `AttributeError`.

15.3.1.6 6. load(fname)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

Purpose

Currently, this method is not supported for KNeighborsClassifier. It is simply kept in KNeighborsClassifier module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an AttributeError.

15.3.1.7 7. predict(X, save_proba = False)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_queries**, **n_features**), where 'n_queries' is the number of rows in the test data.

save_proba: A boolean parameter specifies whether to save the predicted probability or not.

If it is set to 'True', a matrix with the name 'probability_matrix' would be generated in the current execution directory, which can be used for inspecting the probability of individual classes (just for debug purpose).

Purpose

Predict the class labels for the provided data.

For example,

```
# predicting on KNeighborsClassifier model
knc.predict(samples)
```

Output

```
[10 10 10 20 20 20]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)
```

```
# predicting on KNeighborsClassifier model using pre-constructed input
knc.predict(rmat)
```

Output

```
[10 10 10 20 20 20]
```

Return Value

It returns a numpy array of long (int64) type containing the predicted outputs. It is of shape (**n_queries**), where 'n_queries' is the number of rows in the test data.

15.3.1.8 8. predict_proba(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix

for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_queries**, **n_features**), where 'n_queries' is the number of rows in the test data.

Purpose

Calculates probability for the test data X.

For example,

```
knc.predict_proba(samples)
```

Output

```
[[1.      0.      ]
 [1.      0.      ]
 [1.      0.      ]
 [0.33333333 0.66666667]
 [0.33333333 0.66666667]
 [0.33333333 0.66666667]]
```

Like in fit(), frovedis-like input can be used to speed-up the computation of probability at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCSRMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# fitting input data on KNeighborsClassifier object
from frovedis.ml.lib.neighbors import KNeighborsClassifier
knc = KNeighborsClassifier(n_neighbors = 3)
proba = knc.predict_proba(rmat)

# Here FrovedisRowmajorMatrix().debug_print() is used
print(proba)
proba.debug_print()
```

Output

```
matrix:
num_row = 6, num_col = 2
node 0
node = 0, local_num_row = 6, local_num_col = 2, val = 1 0 1 0 1 0 0.333333 0.666667
0.333333 0.666667 0.333333 0.666667
```

It returns a FrovedisRowmajorMatrix object.

Return Value

- **When test data and training data used by fitted model are python native input :**

A numpy array of float or double (float64) type values. It has shape (**n_queries**, **n_neighbors**).

- **When either test data or training data used by fitted model is frovedis-like input:**

A FrovedisRowmajorMatrix of float or double (float64) type values. It has shape (**n_queries**, **n_neighbors**).

15.3.1.9 9. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCSRMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the target labels. It has shape (**n_samples**,).

sample_weight: An unused parameter whose default value is None. It is simply ignored in frovedis implementation, just like in Scikit-learn.

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of `self.predict(X)` wrt. `y`.

For example,

```
knc.score(samples, lbl)
```

Output

```
0.89
```

Return Value

It returns an accuracy score of double (float64) type.

15.3.1.10 9. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by KNeighborsClassifier. It is used to get parameters and their values of KNeighborsClassifier class.

For example,

```
print(knc.get_params())
```

Output

```
{'algorithm': 'auto', 'batch_fraction': None, 'chunk_size': 1.0, 'leaf_size': 30,
'metric': 'euclidean', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 3,
'p': 2, 'verbose': 0, 'weights': 'uniform'}
```

Return Value

A dictionary of parameter names mapped to their values.

15.3.1.11 10. `set_params(**params)`

Parameters

****params**: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by KNeighborsClassifier, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(knc.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
knc.set_params(n_neighbors = 5)
print("Get parameters after setting:")
print(knc.get_params())
```

Output

Get parameters before setting:

```
{'algorithm': 'auto', 'batch_fraction': None, 'chunk_size': 1.0,
'leaf_size': 30, 'metric': 'euclidean', 'metric_params': None, 'n_jobs': None,
'n_neighbors': 3, 'p': 2, 'verbose': 0, 'weights': 'uniform'}
```

Get parameters after setting:

```
{'algorithm': 'auto', 'batch_fraction': None, 'chunk_size': 1.0, 'leaf_size': 30,
'metric': 'euclidean', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 5,
'p': 2, 'verbose': 0, 'weights': 'uniform'}
```

Return Value

It simply returns “self” reference.

15.3.1.12 11. debug_print()**Purpose**

Currently, this method is not supported for KNeighborsClassifier. It is simply kept in KNeighborsClassifier module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an AttributeError.

15.3.1.13 12. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
knc.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

15.3.1.14 13. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns ‘True’, if the model is already fitted otherwise, it returns ‘False’.

15.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMMatrix**
- **Introduction to FrovedisDvector**
- **KNeighbors Regressor in Frovedis**

Chapter 16

KNeighborsRegressor

16.1 NAME

KNeighborsRegressor - Regression based on k-nearest neighbors.

16.2 SYNOPSIS

```
class frovedis.mllib.neighbors.KNeighborsRegressor(n_neighbors=5, weights='uniform',
                                                    algorithm='auto', leaf_size=30, p=2,
                                                    metric='euclidean', metric_params=None,
                                                    n_jobs=None, verbose=0,
                                                    chunk_size=1.0, batch_fraction=None)
```

16.2.1 Public Member Functions

```
fit(X, y)
kneighbors(X = None, n_neighbors = None, return_distance = True)
kneighbors_graph(X = None, n_neighbors = None, mode = 'connectivity')
save(fname)
load(fname)
predict(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

16.3 DESCRIPTION

Neighbors-based regression can be used in cases where the data labels are continuous rather than discrete variables. The label assigned to a query point is computed based on the mean of the labels of its nearest neighbors.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn KNeighborsRegressor interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for KNeighborsRegressor on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When kneighbors-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

16.3.1 Detailed Description

16.3.1.1 1. KNeighborsRegressor()

Parameters

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for 'kneighbors' queries. It must be within the range of 0 and n_samples. (Default: 5)

weights: An unused parameter. (Default: uniform)

algorithm: A string object parameter, specifying the algorithm used to compute the nearest neighbors. (Default: auto)

When it is 'auto', it will be set as 'brute'(brute-force search approach). Unlike Scikit-learn, currently it supports only 'brute'.

leaf_size: An unused parameter. (Default: 30)

p: An unused parameter. (Default: 2)

metric: A string object parameter specifying the distance metric to use for the tree. (Default: 'euclidean') Currently it only supports 'euclidean', 'seuclidean' and 'cosine' distance.

metric_params: An unused parameter. (Default: None)

n_jobs: An unused parameter. (Default: None)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

chunk_size: A positive float parameter specifying the amount of data (in megabytes) to be processed in one time. (Default: 1.0)

batch_fraction: A positive double (float64) parameter used to calculate the batches of specific size. These batches are used to construct the distance matrix. It must be within the range of 0.0 to 1.0. (Default: None) When it is None (not specified explicitly), it will be set as `np.finfo(np.float64).max` value.

Purpose

It initializes a KNeighborsRegressor object with the given parameters.

The parameters: "weights", "leaf_size", "p", "metric_params" and "n_jobs" are simply kept in to make the interface uniform to the Scikit-learn KNeighborsRegressor module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns "self" reference.

16.3.1.2 2. fit(X, y)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (n_samples, n_features).

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape (n_samples,).

Purpose

It accepts the training matrix (X) with labels (y) and trains a KNeighborsRegressor model.

For example,

```
# loading sample data
samples = np.loadtxt("./input/knr_data.txt", dtype = np.float64)
lbl = [10, 10, 10, 20, 10, 20]

# fitting input data on KNeighborsRegressor object
from frovedis.mllib.neighbors import KNeighborsRegressor
knr = KNeighborsRegressor(n_neighbors = 3)
knr.fit(samples, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample data
samples = np.loadtxt("./input/knr_data.txt", dtype = np.float64)
lbl = [10, 10, 10, 20, 10, 20]

# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
rmat = FrovedisRowmajorMatrix(samples)
dlbl = FrovedisDvector(lbl)

# fitting input data on KNeighborsRegressor object
from frovedis.mllib.neighbors import KNeighborsRegressor
knr = KNeighborsRegressor(n_neighbors = 3)
knr.fit(rmat, dlbl)
```

Return Value

It simply returns “self” reference.

16.3.1.3 3. kneighbors(X = None, n_neighbors = None, return_distance = True)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (n_queries, n_features), where ‘n_queries’ is the number of rows in the test data. (Default: None)

When it is None (not specified explicitly), it will be training data (X) used as input in fit().

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for ‘kneighbors’ queries. It must be within the range of 0 and n_samples. (Default: None)

When it is None (not specified explicitly), it will be 'n_neighbors' value used during KNeighborsRegressor object creation.

return_distance: A boolean parameter specifying whether or not to return the distances. (Default: True) If set to False, it will not return distances. Then, only indices are returned by this method.

Purpose

Finds the k-Neighbors of a point and returns the indices of neighbors and distances to the neighbors of each point.

For example,

```
distances, indices = knr.kneighbors(samples)
print('distances')
print(distances)
print('indices')
print(indices)
```

Output

```
distances
[[0.          1.          2.23606798]
 [0.          1.          1.41421356]
 [0.          1.41421356  2.23606798]
 [0.          1.          2.23606798]
 [0.          1.          1.41421356]
 [0.          1.41421356  2.23606798]]
indices
[[0 1 2]
 [1 0 2]
 [2 1 0]
 [3 4 5]
 [4 3 5]
 [5 4 3]]
```

Like in fit(), frovedis-like input can be used to speed-up the computation of indices and distances at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCSRMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# fitting input data on KNeighborsRegressor object
from frovedis.mllib.neighbors import KNeighborsRegressor
knr = KNeighborsRegressor(n_neighbors = 3)
distances, indices = knr.kneighbors(rmat)

# Here FrovedisRowmajorMatrix().debug_print() is used
print('distances')
distances.debug_print()

# Here FrovedisRowmajorMatrix().debug_print() is used
print('indices')
indices.debug_print()
```

Output

```

distances
matrix:
num_row = 6, num_col = 3
node 0
node = 0, local_num_row = 6, local_num_col = 3, val = 0 1 2.23607 0 1 1.41421 0 1.41421
2.23607 0 1 2.23607 0 1 1.41421 0 1.41421 2.23607
indices
matrix:
num_row = 6, num_col = 3
node 0
node = 0, local_num_row = 6, local_num_col = 3, val = 0 1 2 1 0 2 2 1 0 3 4 5 4 3 5 5 4 3

```

It returns distances and indices as FrovedisRowmajorMatrix objects.

Return Value

1. When test data and training data used by fitted model are python native input :
 - **distances**: A numpy array of float or double (float64) type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data. It is only returned by kneighbors(), if return_distance = True.
 - **indices**: A numpy array of int64 type values. It has shape (**n_queries**, **n_neighbors**).
2. When test data and training data used by fitted model is frovedis-like input::
 - **distances**: A FrovedisRowmajorMatrix of float or double (float64) type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data. It is only returned by kneighbors(), if return_distance = True.
 - **indices**: A FrovedisRowmajorMatrix of int64 type values. It has shape (**n_queries**, **n_neighbors**).

16.3.1.4 4. kneighbors_graph(X = None, n_neighbors = None, mode = 'connectivity')

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_queries**, **n_features**), where 'n_queries' is the number of rows in the test data. (Default: None)

When it is None (not specified explicitly), it will be training data (X) used as input in fit().

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for 'kneighbors' queries. It must be within the range of 0 and n_samples. (Default: None)

When it is None (not specified explicitly), it will be 'n_neighbors' value used during KNeighborsRegressor object creation.

mode: A string object parameter which can be either 'connectivity' or 'distance'. It specifies the type of returned matrix.

For 'connectivity', it will return the connectivity matrix with ones and zeros, whereas for 'distance', the edges are euclidean distance between points, type of distance depends on the selected 'metric' value in KNeighborsRegressor class. (Default: 'connectivity')

Purpose

It computes the (weighted) graph of k-Neighbors for points in X.

For example,

```

# Here 'mode = connectivity' by default
graph = knr.kneighbors_graph(samples)
print('kneighbors graph')
print(graph)

```

Output

```
kneighbors graph
(0, 0)      1.0
(0, 1)      1.0
(0, 2)      1.0
(1, 1)      1.0
(1, 0)      1.0
(1, 2)      1.0
(2, 2)      1.0
(2, 1)      1.0
(2, 0)      1.0
(3, 3)      1.0
(3, 4)      1.0
(3, 5)      1.0
(4, 4)      1.0
(4, 3)      1.0
(4, 5)      1.0
(5, 5)      1.0
(5, 4)      1.0
(5, 3)      1.0
```

For example, when mode = 'distance'

```
# Here 'mode = distance'
graph = knr.kneighbors_graph(samples, mode = 'distance')
print('kneighbors graph')
print(graph)
```

Output

```
kneighbors graph
(0, 0)      0.0
(0, 1)      1.0
(0, 2)      2.23606797749979
(1, 1)      0.0
(1, 0)      1.0
(1, 2)      1.4142135623730951
(2, 2)      0.0
(2, 1)      1.4142135623730951
(2, 0)      2.23606797749979
(3, 3)      0.0
(3, 4)      1.0
(3, 5)      2.23606797749979
(4, 4)      0.0
(4, 3)      1.0
(4, 5)      1.4142135623730951
(5, 5)      0.0
(5, 4)      1.4142135623730951
(5, 3)      2.23606797749979
```

Like in fit(), frovedis-like input can be used to speed-up the graph making at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)
```

```
# fitting input data on KNeighborsRegressor object
from frovedis.mllib.neighbors import KNeighborsRegressor
knr = KNeighborsRegressor(n_neighbors = 3)

# Here 'mode = connectivity' by default
graph = knr.kneighbors_graph(rmat)

# Here FrovedisCRSMatrix().debug_print() is used
print('graph')
graph.debug_print()
```

Output

```
graph
Active Elements:  18
matrix:
num_row = 6, num_col = 6
node 0
local_num_row = 6, local_num_col = 6
val : 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
idx : 0 1 2 1 0 2 2 1 0 3 4 5 4 3 5 4 3
off : 0 3 6 9 12 15 18
```

It returns a FrovedisCRSMatrix object for frovedis-like input.

Return Value

- **When test data and training data used by fitted model are python native input:**

It returns a scipy sparse matrix of float or double (float64) type values. It has shape (**n_queries**, **n_samples_fit**), where 'n_queries' is the number of rows in the test data and 'n_samples_fit' is the number of samples in the fitted data.

- **When either test data or training data used by fitted model is frovedis-like input:**

It returns a FrovedisCRSMatrix of float or double (float64) type values. It has shape (**n_queries**, **n_samples_fit**), where 'n_queries' is the number of rows in the test data and 'n_samples_fit' is the number of samples in the fitted data.

16.3.1.5 5. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

Currently, this method is not supported for KNeighborsRegressor. It is simply kept in KNeighborsRegressor module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an AttributeError.

16.3.1.6 6. load(fname)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

Purpose

Currently, this method is not supported for KNeighborsRegressor. It is simply kept in KNeighborsRegressor module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an `AttributeError`.

16.3.1.7 7. predict(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data of float or double (float64) type. It has shape `(n_queries, n_features)`, where 'n_queries' is the number of rows in the test data.

Purpose

Predict the class labels for the provided data.

For example,

```
# predicting on KNeighborsRegressor model
knr.predict(samples)
```

Output

```
[10.          10.          10.          16.66666667 16.66666667 16.66666667]
```

Like in `fit()`, frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)
```

```
# predicting on KNeighborsRegressor model using pre-constructed input
knr.predict(rmat)
```

Output

```
[10.          10.          10.          16.66666667 16.66666667 16.66666667]
```

Return Value

It returns a numpy array of long (int64) type containing the predicted outputs. It is of shape `(n_queries,)`, where 'n_queries' is the number of rows in the test data.

16.3.1.8 8. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data of float or double (float64) type. It has shape `(n_samples, n_features)`.

y: Any python array-like object or an instance of `FrovedisDvector` containing the target values. It has shape `(n_samples,)`.

sample_weight: An unused parameter whose default value is `None`. It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of `self.predict(X)` wrt. `y`.

The coefficient 'R2' is defined as $(1 - (u/v))$,

where 'u' is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and,

'v' is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
knr.score(samples, lbl)
```

Output

```
0.5
```

Return Value

It returns an R2 score of float type.

16.3.1.9 9. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by KNeighborsRegressor. It is used to get parameters and their values of KNeighborsRegressor class.

For example,

```
print(knr.get_params())
```

Output

```
{'algorithm': 'auto', 'batch_fraction': None, 'chunk_size': 1.0, 'leaf_size': 30,
'metric': 'euclidean', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 3,
'p': 2, 'verbose': 0, 'weights': 'uniform'}
```

Return Value

A dictionary of parameter names mapped to their values.

16.3.1.10 10. set_params(**params)

Parameters

****params:** All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by KNeighborsRegressor, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(knr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
knr.set_params(n_neighbors = 5)
print("Get parameters after setting:")
print(knr.get_params())
```

Output

```
Get parameters before setting:
{'algorithm': 'auto', 'batch_fraction': None, 'chunk_size': 1.0, 'leaf_size': 30,
'metric': 'euclidean', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 3,
'p': 2, 'verbose': 0, 'weights': 'uniform '}
Get parameters after setting:
{'algorithm': 'auto', 'batch_fraction': None, 'chunk_size': 1.0, 'leaf_size': 30,
'metric': 'euclidean', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 5,
'p': 2, 'verbose': 0, 'weights': 'uniform '}
```

Return Value

It simply returns “self” reference.

16.3.1.11 11. `debug_print()`

Purpose

Currently, this method is not supported for KNeighborsRegressor. It is simply kept in KNeighborsRegressor module to maintain uniform interface like other estimators in frovedis.

Return Value

It simply raises an `AttributeError`.

16.3.1.12 12. `release()`

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
knr.release()
```

This will remove the trained model, model-id present on server, along with releasing server side memory.

Return Value

It returns nothing.

16.3.1.13 13. `is_fitted()`

Purpose

It can be used to confirm if the model is already fitted or not.

Return Value

It returns ‘True’, if the model is already fitted otherwise, it returns ‘False’.

16.4 SEE ALSO

- [Introduction to FrovedisRowmajorMatrix](#)
- [Introduction to FrovedisCRSMatrix](#)
- [Introduction to FrovedisDvector](#)
- [KNeighbors Classifier in Frovedis](#)

Chapter 17

Lasso Regression

17.1 NAME

Lasso Regression - A regression algorithm used to predict the continuous output with L1 regularization.

17.2 SYNOPSIS

```
class frovedis.mllib.linear_model.Lasso(alpha=0.01, fit_intercept=True,
                                         normalize=False, precompute=False,
                                         copy_X=True, max_iter=1000,
                                         tol=1e-4, warm_start=False,
                                         positive=False, random_state=None,
                                         selection='cyclic', lr_rate=1e-8,
                                         verbose=0, solver='sag')
```

17.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

17.3 DESCRIPTION

Linear least squares is the most common formulation for regression problems. It is a linear method with the loss function given by the **squared loss**:

$$L(\mathbf{w}; \mathbf{x}, y) := 1/2(\mathbf{w}^T \mathbf{x} - y)^2$$

During training, the input \mathbf{X} is the training data and \mathbf{y} is the corresponding label values which we want to predict. \mathbf{w} is the linear model (also known as weight) which uses a single weighted sum of features to make a prediction. The method is called linear since it can be expressed as a function of $\mathbf{w}^T\mathbf{x}$ and \mathbf{y} . **Lasso regression uses L1 regularization to address the overfit problem.**

The gradient of the squared loss is: $(\mathbf{w}^T\mathbf{x}-\mathbf{y})\cdot\mathbf{x}$

The gradient of the regularizer is: $\text{sign}(\mathbf{w})$

Frovedis provides implementation of lasso regression with two different optimizers:

- (1) stochastic gradient descent with minibatch
- (2) LBFGS optimizer

The simplest method to solve optimization problems of the form $\min \mathbf{f}(\mathbf{w})$ is gradient descent. Such first-order optimization methods well-suited for large-scale and distributed computation. Whereas, L-BFGS is an optimization algorithm in the family of quasi-Newton methods to solve the optimization problems of the similar form.

Like the original BFGS, L-BFGS (Limited Memory BFGS) uses an estimation to the inverse Hessian matrix to steer its search through feature space, but where BFGS stores a dense $n \times n$ approximation to the inverse Hessian (n being the number of features in the problem), L-BFGS stores only a few vectors that represent the approximation implicitly. L-BFGS often achieves rapider convergence compared with other first-order optimization.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn Lasso interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Lasso on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

17.3.1 Detailed Description

17.3.1.1 1. Lasso()

Parameters

alpha: A constant that multiplies the L1 term. It must be a positive value of double (float64) type . (Default: 0.01)

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the decision function. (Default: True)

normalize: An unused parameter. (Default: False)

precompute: An unused parameter. (Default: False)

copy_X: An unused parameter. (Default: True)

max_iter: A positive integer value used to set the maximum number of iterations. (Default: 1000)

tol: Zero or a positive value of double (float64) type specifying the convergence tolerance value. (Default: 1e-4)

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

positive: An unused parameter. (Default: False)

random_state: An unused parameter. (Default: None)

selection: An unused parameter. (Default: cyclic)

lr_rate: Zero or a positive value of double (float64) type containing the learning rate. (Default: 1e-8)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

solver: A string object specifying the solver to use. (Default: 'sag')

It can be "sag" for frovedis side stochastic gradient descent or "lbfgs" for frovedis side LBFGS optimizer when optimizing the lasso regression model. **Both "sag" and "lbfgs" can handle L1 penalty.**

Attributes

coef_: It is a python ndarray (containing float or double (float64) typed values depending on data-type of input matrix (X)) of estimated coefficients for the lasso regression problem. It has shape (**n_features**).

intercept_(bias): It is a python ndarray (containing float or double (float64) typed values depending on data-type of input matrix (X)). If fit_intercept is set to False, the intercept is set to zero. It has shape (**1**).

n_iter_: A positive integer value used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a Lasso object with the given parameters.

The parameters: "normalize", "precompute", "copy_X", "positive", "random_state" and "selection" are simply kept in to make the interface uniform to the Scikit-learn Lasso module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns "self" reference.

17.3.1.2 2. fit(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisColmajorMatrix for dense data.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a lasso regression model with L1 regularization with those data at frovedis server.

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used as well to speed up the training time, especially when same data would be used for multiple executions.

Return Value

It simply returns "self" reference.

17.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix as for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

In case pre-constructed frovedis-like training data such as FrovedisColmajorMatrix (X) is provided during prediction, then “X.to_frovedis_rowmatrix()” will be used for prediction.

Return Value

It returns a numpy array of float or double (float64) type and has shape (**n_samples**,) containing the predicted outputs.

17.3.1.4 4. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: Any python array-like object containing true values for X. It has shape (**n_samples**,).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of self.predict(X) wrt. y.

The coefficient ‘ R^2 ’ is defined as $(1 - (u/v))$,

where ‘u’ is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and

‘v’ is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R^2 score of 0.0.

Return Value

It returns an R^2 score of float type.

17.3.1.5 5. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by Lasso. It is used to get parameters and their values of Lasso class.

Return Value

A dictionary of parameter names mapped to their values.

17.3.1.6 6. set_params(**params)

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by Lasso, used to set parameter values.

Return Value

It simply returns “self” reference.

17.3.1.7 7. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

Return Value

It simply returns “self” reference.

17.3.1.8 8. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

Suppose ‘LassoModel’ directory is the model created, It will have

LassoModel

```
|—— metadata
|—— model
```

The metadata file contains the number of classes, model kind, input datatype used for trained model. Here, the model file contains information about weights, intercept and threshold.

It would raise exception if the ‘LassoModel’ directory already existed with same name.

Return Value

It returns nothing.

17.3.1.9 9. debug_print()**Purpose**

It shows the target model information like weight values, intercept on the server side user terminal. It is mainly used for debugging purpose.

No such output will be visible on client side.

Return Value

It returns nothing.

17.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server. With this, after-fit populated attributes are reset to None, along with releasing server side memory.

Return Value

It returns nothing.

17.3.1.11 11. `is_fitted()`**Purpose**

It can be used to confirm if the model is already fitted or not. In case, `predict()` is used before training the model, then it can prompt the user to train the lasso regression model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

17.4 SEE ALSO

- **Introduction to `FrovedisRowmajorMatrix`**
- **Introduction to `FrovedisColmajorMatrix`**
- **Introduction to `FrovedisCRSMatrix`**
- **Introduction to `FrovedisDvector`**
- **Linear Regression in `Frovedis`**
- **Ridge Regression in `Frovedis`**

Chapter 18

Latent Dirichlet Allocation

18.1 NAME

Latent Dirichlet Allocation(LDA) - It is a Natural Language Processing algorithm that allows a set of observations to be explained by unobserved groups. These are used to explain why some parts of the data are similar.

18.2 SYNOPSIS

```
class frovedis.mllib.decomposition.LatentDirichletAllocation(n_components=10,
                                                             doc_topic_prior=None,
                                                             topic_word_prior=None,
                                                             learning_method='batch',
                                                             learning_decay=.7,
                                                             learning_offset=10.,
                                                             max_iter=10, batch_size=128,
                                                             evaluate_every=-1,
                                                             total_samples=1e6,
                                                             perp_tol=1e-1,
                                                             mean_change_tol=1e-3,
                                                             max_doc_update_iter=100,
                                                             n_jobs=None, verbose=0,
                                                             random_state=None,
                                                             algorithm="original",
                                                             explore_iter=0)
```

18.2.1 Public Member Functions

```
fit(X, y = None)
transform(X)
perplexity(X, sub_sampling = False)
score(X, y = None)
get_params(deep = True)
set_params(**params)
fit_transform(X, y = None)
```

```
load(fname, dtype = None)
save(fname)
release()
is_fitted()
```

18.3 DESCRIPTION

Latent Dirichlet Allocation (LDA) is a widely used machine learning technique for topic modeling. The input is a corpus of documents and the output is per-document topic distribution and per-topic word distribution.

This implementation provides python wrapper interface to Vectorised LDA (VLDA), which accelerates LDA training by exploiting the data-level, and the thread-level parallelism using vector processors. The priority-aware scheduling approach is proposed to address the high memory requirement and workload imbalance issues with existing works.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn LatentDirichletAllocation interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for LatentDirichletAllocation on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When transform-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

18.3.1 Detailed Description

18.3.1.1 1. LatentDirichletAllocation()

Parameters

n_components: A positive integer parameter specifying the number of topics. (Default: 10)

doc_topic_prior: A float parameter which specifies the prior of document topic distribution theta. It is also called alpha. (Default: None)

If it is None (not specified explicitly), it will be set as $(1 / n_components)$.

topic_word_prior: A float parameter which specifies the prior of topic word distribution beta. It is also called beta. (Default: None)

If it is None (not specified explicitly), it will be set as $(1 / n_components)$.

learning_method: An unused parameter which specifies the method used to update 'components_'.

Although, this parameter is unused in frovedis but it must be 'batch' or 'online' update. This is simply done to keep the behavior consistent with Scikit-learn. (Default: batch)

learning_decay: An unused parameter. (Default: 0.7)

learning_offset: An unused parameter. Although, this parameter is unused in frovedis but it must be zero or a positive float value. This is simply done to keep the behavior consistent with Scikit-learn. (Default: 10.0)

max_iter: An integer parameter which specifies the maximum number of passes over the training data. (Default: 10)

batch_size: An unused parameter. (Default: 128)

evaluate_every: An integer parameter that specifies how often to evaluate perplexity. Perplexity is not

evaluated in training by default. If this parameter is greater than 0, perplexity will be evaluated, which can help in checking convergence in training process, but total training time will be increased. (Default: -1)

For example, when `evaluate_every < 0`

```
# loading sample csr matrix data representing the input corpus-vocabulary count
corpus = [
    'This is the first document.',
    'This document is the second document.',
    'And this is the third one.',
    'Is this the first document?',
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(corpus)

# fitting input matrix on lda object
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3, evaluate_every = -1)
lda.fit(csr_mat)
```

LDA training time: 0.0007402896881103516

For example, when `evaluate_every > 0`

```
# loading sample csr matrix data representing the input corpus-vocabulary count
corpus = [
    'This is the first document.',
    'This document is the second document.',
    'And this is the third one.',
    'Is this the first document?',
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(corpus)

# fitting input matrix on lda object
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3, evaluate_every = 2)
lda.fit(csr_mat)
```

LDA training time: 0.0008301734924316406

total_samples: A positive integer parameter which is unused in frovedis. This is simply kept to make the behavior consistent with Scikit-learn. (Default: 1e6)

perp_tol: An unused parameter. (Default: 1e-1)

mean_change_tol: An unused parameter. (Default: 1e-3)

max_doc_update_iter: An unused parameter. (Default: 100)

n_jobs: An unused parameter. (Default: None)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1(for DEBUG mode) or 2(for TRACE mode) for getting training time logs from frovedis server.

algorithm: A string object parameter which specifies the sampling technique to be used. (Default: 'original')

Frovedis `LatentDirichletAllocation(LDA)` supports either of two sampling techniques:

1. Collapsed Gibbs Sampling (CGS)

2. Metropolis Hastings

The default sampling algorithm is CGS (Default: 'original')

If Metropolis Hastings is to be used, then it is required to set the proposal types namely:

2a. document proposal: “dp” algo,

2b. word proposal: “wp” algo,

2c. cycle proposal: “cp” algo,

2d. sparse lda: “sparse”

`explore_iter`: An integer parameter that specifies the number of iterations to explore optimal hyperparams. (Default: 0)

Attributes

`components_`: A numpy ndarray of double (float64) type values and has shape (**`n_components`**, **`n_features`**).

Purpose

It initializes a `LatentDirichletAllocation` object with the given parameters.

The parameters: “`learning_method`”, “`learning_decay`”, “`learning_offset`”, “`batch_size`”, “`total_samples`”, “`perp_tol`”, “`mean_change_tol`”, “`max_doc_update_iter`” and “`n_jobs`” are simply kept in to make the interface uniform to the Scikit-learn `LatentDirichletAllocation` module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

18.3.1.2 2. `fit(X, y = None)`

Parameters

`X`: A scipy sparse matrix or an instance of `FrovedisCRSMatrix` for sparse data of int, float or double (float64) type. It has shape (**`n_samples`**, **`n_features`**).

`y`: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

Fit `LatentDirichletAllocation` model on training data X.

For example,

```
# loading sample csr matrix data representing the input corpus-vocabulary count
corpus = [
    'This is the first document.',
    'This document is the second document.',
    'And this is the third one.',
    'Is this the first document?',
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(corpus)

# fitting input matrix on lda object
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3)
lda.fit(csr_mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample csr matrix data representing the input corpus-vocabulary count
corpus = [
    'This is the first document.',
    'This document is the second document.',
    'And this is the third one.',
    'Is this the first document?',
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(corpus)

# Since "csr_mat" is scipy sparse data, we have created FrovedisCRSMatrix
# Also it only supports int64 as itype for input sparse data during training
from frovedis.matrix.crs import FrovedisCRSMatrix
cmat = FrovedisCRSMatrix(csr_mat, dtype = np.int32, itype = np.int64)

# fitting input matrix on lda object
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3)
lda.fit(cmat)
```

Return Value

It simply returns “self” reference.

18.3.1.3 3. transform(X)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix for sparse data of int, float or double (float64) type. It has shape (**n_samples**, **n_features**).

Purpose

It transforms input matrix X according to the trained model.

For example,

```
# loading sample csr matrix data representing the input corpus-vocabulary count
test_corpus = [
    'This is the first second third document.',
    'This This one third document.'
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(test_corpus)

# transform input according to trained model
from frovedis.mllib.decomposition import LatentDirichletAllocation
```

```
lda = LatentDirichletAllocation(n_components = 3)
lda.fit(csr_mat)
print(lda.transform(csr_mat))
```

Output

```
[[0.47619048 0.19047619 0.47619048]
 [0.26666667 0.26666667 0.66666667]]
```

Like in `fit()`, we can also provide frovedis-like input in `transform()` for faster computation.

For example,

```
# loading sample csr matrix data representing the input corpus-vocabulary count
test_corpus = [
    'This is the first second third document.',
    'This This one third document.'
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(test_corpus)

# Since "csr_mat" is scipy sparse data, we have created FrovedisCRSMatrix
# Also it only supports int64 as itype for input sparse data during training
from frovedis.matrix.crs import FrovedisCRSMatrix
cmat = FrovedisCRSMatrix(csr_mat, dtype = np.int32, itype = np.int64)

# transform input according to trained model
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3)
lda.fit(cmat)
lda.transform(cmat).debug_print()
```

Output

```
matrix:
num_row = 2, num_col = 3
node 0
node = 0, local_num_row = 2, local_num_col = 3, val = 0.47619 0.333333 0.333333 0.266667
0.0666667 0.866667
```

Return Value

- When X is python native input:

It returns a python ndarray of shape **(n_samples, n_components)** and double (float64) type values. It contains the document-wise topic distribution for input data X.

- **When X is frovedis-like input:**

It returns a FrovedisRowMajorMatrix of shape **(n_samples, n_components)** and double (float64) type values, containing document-wise topic distribution for input data X.

18.3.1.4 4. perplexity(X, sub_sampling = False)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix for sparse data of int, float, double (float64) type. It has shape **(n_samples, n_features)**.

sub_sampling: A boolean parameter that specifies whether to do sub-sampling or not. It is simply ignored in frovedis implementation. (Default: False)

Purpose

Calculate approximate perplexity for data X.

Perplexity is defined as $\exp(-1. * \text{log-likelihood per word})$.

Ideally, as the number of components increase, the perplexity of model should decrease.

For example,

```
perp = lda.perplexity(csr_mat)
print("perplexity: %.2f" % (perp))
```

Output

```
perplexity: 8.81
```

Return Value

It returns float type perplexity score.

18.3.1.5 5. score(X, y = None)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMatrix for sparse data of int, float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, just like in Scikit-learn.

Purpose

Calculate approximate log-likelihood as score for data X.

Likelihood is a measure of how plausible model parameters are, given the data. Taking a logarithm makes calculations easier. All values are negative: when $x < 1$, $\log(x) < 0$.

The idea is to search for the largest log-likelihood (good score will be close to 0).

For example,

```
print("score: %.2f" % (lda.score(csr_mat)))
```

Output

```
score: -2.18
```

Return Value

It returns a float value as likelihood score

18.3.1.6 6. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by LatentDirichletAllocation. It is used to get parameters and their values of LatentDirichletAllocation class.

For example,

```
print(lda.get_params())
```

Output

```
{'algorithm': 'original', 'batch_size': 128, 'doc_topic_prior': 0.3333333333333333,
'evaluate_every': -1, 'explore_iter': 0, 'learning_decay': 0.7, 'learning_method': 'batch',
'learning_offset': 10.0, 'max_doc_update_iter': 100, 'max_iter': 10,
'mean_change_tol': 0.001, 'n_components': 3, 'n_jobs': None, 'perp_tol': 0.1,
'random_state': None, 'topic_word_prior': 0.3333333333333333, 'total_samples': 1000000.0,
'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

18.3.1.7 7. set_params(**params)

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by LatentDirichletAllocation, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(lda.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
lda.set_params(n_components = 4, algorithm = 'dp')
print("get parameters after setting:")
print(lda.get_params())
```

Output

```
get parameters before setting:
{'algorithm': 'original', 'batch_size': 128, 'doc_topic_prior': 0.3333333333333333,
'evaluate_every': -1, 'explore_iter': 0, 'learning_decay': 0.7, 'learning_method': 'batch',
'learning_offset': 10.0, 'max_doc_update_iter': 100, 'max_iter': 10,
'mean_change_tol': 0.001, 'n_components': 3, 'n_jobs': None, 'perp_tol': 0.1,
'random_state': None, 'topic_word_prior': 0.3333333333333333, 'total_samples': 1000000.0,
'verbose': 0}
get parameters after setting:
{'algorithm': 'dp', 'batch_size': 128, 'doc_topic_prior': 0.3333333333333333,
'evaluate_every': -1, 'explore_iter': 0, 'learning_decay': 0.7, 'learning_method': 'batch',
'learning_offset': 10.0, 'max_doc_update_iter': 100, 'max_iter': 10,
'mean_change_tol': 0.001, 'n_components': 4, 'n_jobs': None, 'perp_tol': 0.1,
'random_state': None, 'topic_word_prior': 0.3333333333333333, 'total_samples': 1000000.0,
'verbose': 0}
```

Return Value

It simply returns “self” reference.

18.3.1.8 8. fit_transform(X, y = None)

Parameters

X: A scipy sparse matrix or an instance of FrovedisCRSMMatrix for sparse data of int, float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It performs `fit()` and `transform()` on `X` and `y` (unused), and returns a transformed version of `X`.

For example,

```
# loading sample csr matrix data representing the input corpus-vocabulary count
test_corpus = [
    'This is the first second third document.',
    'This This one third document.'
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(test_corpus)

# transform input after fitting
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3)
print(lda.fit_transform(csr_mat))
```

Output

```
[[0.04761905 0.61904762 0.47619048]
 [0.06666667 0.66666667 0.46666667]]
```

Like in `fit()` and `transform()`, we can also provide frovedis-like input in `fit_transform()` for faster computation.

For example,

```
# loading sample csr matrix data representing the input corpus-vocabulary count
test_corpus = [
    'This is the first second third document.',
    'This This one third document.'
]

# feature transformers CountVectorizer can be used for converting text to word count vectors.
from sklearn.feature_extraction.text import CountVectorizer
vectorizer = CountVectorizer()
csr_mat = vectorizer.fit_transform(test_corpus)

# Since "csr_mat" is scipy sparse data, we have created FrovedisCRSMatrix
# Also it only supports int64 as itype for input sparse data during training
from frovedis.matrix.crs import FrovedisCRSMatrix
cmat = FrovedisCRSMatrix(csr_mat, dtype = np.int32, itype = np.int64)

# transform input according to trained model
from frovedis.mllib.decomposition import LatentDirichletAllocation
lda = LatentDirichletAllocation(n_components = 3)
lda.fit_transform(cmat).debug_print()
```

Output

```
matrix:
num_row = 2, num_col = 3
node 0
```

```
node = 0, local_num_row = 2, local_num_col = 3, val = 0.333333 0.619048 0.190476
0.266667 0.666667 0.266667
```

Return Value

- When X is python native input:

It returns a numpy ndarray of shape (**n_samples**, **n_components**) and double (float64) type values. This is a transformed array X.

- **When X is frovedis-like input:**

It returns a FrovedisRowmajorMatrix of shape (**n_samples**, **n_components**) and double (float64) type values. This is a transformed array X.

18.3.1.9 9. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
lda.load("./out/trained_lda_model")
```

Return Value

It simply returns “self” reference.

18.3.1.10 10. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (corpus_topic, metadata and word_topic) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the lda model
lda.save("./out/trained_lda_model")
```

This will save the lda model on the path “/out/trained_lda_model”.

It would raise exception if the directory already exists with same name.

The ‘trained_lda_model’ directory has

trained_lda_model

```
|—corpus_topic
|—metadata
|—word_topic
```

Here, the **corpus_topic** directory contains the corpus topic count as sparse matrix in binary format.

The metadata file contains the information of model kind, input datatype used for trained model.

Here, the **word_topic** directory contains the word topic count as sparse matrix in binary format.

Return Value

It returns nothing.

18.3.1.11 11. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
lda.release()
```

This will reset the after-fit populated attributes (like `components__`) to `None`, along with releasing server side memory.

Return Value

It returns nothing.

18.3.1.12 12. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

18.4 SEE ALSO

- **Introduction to FrovedisCRSMatrix**

Chapter 19

Linear Regression

19.1 NAME

Linear Regression - A regression algorithm used to predict the continuous output without any regularization.

19.2 SYNOPSIS

```
class frovedis.mllib.linear_model.LinearRegression(fit_intercept=True, normalize=False,
                                                    copy_X=True, n_jobs=None,
                                                    max_iter=None, tol=0.0001,
                                                    lr_rate=1e-8, solver=None,
                                                    verbose=0, warm_start=False)
```

19.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

19.3 DESCRIPTION

Linear least squares is the most common formulation for regression problems. It is a linear method with the loss function given by the **squared loss**:

$$L(\mathbf{w}; \mathbf{x}, y) := 1/2(\mathbf{w}^T \mathbf{x} - y)^2$$

During training, the input \mathbf{X} is the training data and \mathbf{y} is the corresponding label values which we want to predict. \mathbf{w} is the linear model (also known as weight) which uses a single weighted sum of features to

make a prediction. The method is called linear since it can be expressed as a function of wTx and y . **Linear Regression does not use any regularizer.**

The gradient of the squared loss is: $(wTx - y) \cdot x$

Frovedis provides implementation of linear regression with the following optimizers:

- (1) stochastic gradient descent with minibatch
- (2) LBFGS optimizer
- (3) least-square

The simplest method to solve optimization problems of the form $\min f(w)$ is gradient descent. Such first-order optimization methods well-suited for large-scale and distributed computation. Whereas, L-BFGS is an optimization algorithm in the family of quasi-Newton methods to solve the optimization problems of the similar form.

Like the original BFGS, L-BFGS (Limited Memory BFGS) uses an estimation to the inverse Hessian matrix to steer its search through feature space, but where BFGS stores a dense $n \times n$ approximation to the inverse Hessian (n being the number of features in the problem), L-BFGS stores only a few vectors that represent the approximation implicitly. L-BFGS often achieves rapid convergence compared with other first-order optimization.

For least-square solver, we have used LAPACK routine “gelsd” and ScaLAPACK routine “gels” when input data is dense in nature. For the sparse-input we have provided a least-square implementation, similar to `scipy.sparse.linalg.lsqr`.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn Linear Regression interface, but it doesn’t have any dependency with Scikit-learn. It can be used simply even if the system doesn’t have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server by sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Linear Regression on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

19.3.1 Detailed Description

19.3.1.1 1. LinearRegression()

Parameters

fit_intercept: A boolean parameter specifying whether a constant(intercept) should be added to the decision function. (Default: True)

normalize: An unused parameter. (Default: False)

copy_X: An unused parameter. (Default: True)

n_jobs: An unused parameter. (Default: None)

max_iter: A positive integer parameter used to set the maximum number of iterations. When it is None(not specified explicitly), it will be set as 1000 for “sag”, “lbfgs”, “lapack” and “scalapack” solvers and for “sparse_lsqr” solver, it will be $2 * (n_features)$. (Default: None)

tol: Zero or a positive value of double (float64) type specifying the convergence tolerance value. (Default: 0.001)

lr_rate: A positive value of double (float64) type containing the learning rate. (Default: 1e-8)

solver: A string parameter specifying the solver to use. (Default: None).

When it is None (not explicitly specified), the value will be set to “lapack” when dense input matrix (X)

is provided and for sparse input matrix (X), it is set as “sparse_lsqr”. Frovedis supports “sag”, “lbfgs”, “lapack”, “scalapack”, “sparse_lsqr” solvers.

“lapack” and “scalapack” solvers can only work with dense data.

“sparse_lsqr” solver can only work with sparse data.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. Only supported by “sag” and “lbfgs” solvers. (Default: False)

Attributes

coef_: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)) of estimated coefficients for the linear regression problem. It has shape (**n_features**,).

rank_: An integer value used to store rank of matrix (X). It is only available when matrix (X) is dense and “lapack” solver is used.

singular_: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)) and has shape (**min(X,y)**,) which is used to store singular values of X. It is only available when X is dense and “lapack” solver is used.

intercept_ (bias): It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)). If fit_intercept is set to False, the intercept is set to zero. It has shape (**1**,).

n_iter_: A positive integer value used to get the actual iteration point at which the problem is converged. It is only available for “sag”, “lbfgs” and “sparse_lsqr” solvers.

Purpose

It initializes a Linear Regression object with the given parameters.

The parameters: “normalize”, “copy_X” and “n_jobs” are simply kept in to to make the interface uniform to the Scikit-learn Linear Regression module. They are not used anywhere within the frovedis implementation.

“solver” can be “sag” for frovedis side stochastic gradient descent, “lbfgs” for frovedis side LBFGS optimizer, “sparse_lsqr”, “lapack” and “scalapack” when optimizing the linear regression model.

“max_iter” can be used to set the maximum iterations to achieve the convergence. In case the convergence is not achieved, it displays a warning for the same.

Return Value

It simply returns “self” reference.

19.3.1.2 2. fit(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisColmajorMatrix for dense data.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**,).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a linear regression model with those data at frovedis server.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_boston
mat, label = load_boston(return_X_y = True)

# fitting input matrix and label on linear regression object
from frovedis.mllib.linear_model import LinearRegression
lr = LinearRegression(solver = 'sag').fit(mat,label)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_boston
mat, label = load_boston(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# Linear Regression with pre-constructed frovedis-like inputs
from frovedis.mllib.linear_model import LinearRegression
lr = LinearRegression(solver = 'sag').fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

19.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
# predicting on sag linear regression model
lr.predict(mat[:10])
```

Output

```
[30.00384338 25.02556238 30.56759672 28.60703649 27.94352423 25.25628446
 23.00180827 19.53598843 11.52363685 18.92026211]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
```



```
# predicting on sag linear regression model using pre-constructed input
lr.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[30.00384338 25.02556238 30.56759672 28.60703649 27.94352423 25.25628446
 23.00180827 19.53598843 11.52363685 18.92026211]
```

Return Value

It returns a numpy array of float or double (float64) type and has shape **(n_samples,)** containing the predicted outputs.

19.3.1.4 4. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: Any python array-like object containing the true values for X. It has shape **(n_samples,)**.

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of **(n_samples,)**.

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of self.predict(X) wrt. y.

The coefficient 'R2' is defined as $(1 - (u/v))$,

where 'u' is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and,

'v' is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
# calculate R2 score on given test data and labels
lr.score(mat[:10], label[:10])
```

Output

```
0.40
```

Return Value

It returns an R2 score of float type.

19.3.1.5 5. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by LinearRegression. It is used to get parameters and their values of LinearRegression class.

For example,

```
print(lr.get_params())
```

Output

```
{'copy_X': True, 'fit_intercept': True, 'lr_rate': 1e-08, 'max_iter': None, 'n_jobs': None,
'normalize': False, 'solver': 'lapack', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

19.3.1.6 6. `set__params(**params)`

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by LinearRegression, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(lr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
lr.set_params(solver='lbfgs', max_iter = 10000)
print("get parameters after setting:")
print(lr.get_params())
```

Output

```
get parameters before setting:
{'copy_X': True, 'fit_intercept': True, 'lr_rate': 1e-08, 'max_iter': None, 'n_jobs': None,
'normalize': False, 'solver': 'lapack', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
get parameters after setting:
{'copy_X': True, 'fit_intercept': True, 'lr_rate': 1e-08, 'max_iter': 10000, 'n_jobs': None,
'normalize': False, 'solver': 'lbfgs', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

19.3.1.7 7. `load(fname, dtype = None)`

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
lr.load("./out/LNRModel")
```

Return Value

It simply returns “self” reference.

19.3.1.8 8. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the linear regression model
lr.save("./out/LNRModel")
```

This will save the linear regression model on the path “/out/LNRModel”. It would raise exception if the directory already exists with same name.

The ‘LNRModel’ directory has

LNRModel

```
|—— metadata
|—— model
```

The metadata file contains the number of classes, model kind, input datatype used for trained model. Here, the model file contains information about weights, intercept.

Return Value

It returns nothing.

19.3.1.9 9. debug_print()**Purpose**

It shows the target model information (weight values, intercept) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
lr.debug_print()
```

Output

```
----- Weight Vector:: -----
-0.108011 0.0464205 0.0205586 2.68673 -17.7666 3.80987 0.000692225 -1.47557 0.306049
-0.0123346 -0.952747 0.00931168 -0.524758
Intercept:: 36.4595
```

This output will be visible on server side. It displays the weights and intercept values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

19.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
lr.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

19.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, `predict()` is used before training the model, then it can prompt the user to train the linear regression model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

19.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Lasso Regression in Frovedis**
- **Ridge Regression in Frovedis**

Chapter 20

LinearSVC

20.1 NAME

LinearSVC (Support Vector Classification) - A classification algorithm used to predict the binary output with hinge loss.

20.2 SYNOPSIS

```
class frovedis.mllib.svm.LinearSVC(penalty='l2', loss='hinge', dual=True, tol=1e-4,
                                   C=1.0, multi_class='ovr', fit_intercept=True,
                                   intercept_scaling=1, class_weight=None,
                                   verbose=0, random_state=None,
                                   max_iter=1000, lr_rate=0.01,
                                   solver='sag', warm_start=False)
```

20.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
load(fname, dtype = None)
save(fname)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

20.3 DESCRIPTION

Classification aims to divide items into categories. The most common classification type is binary classification, where there are two categories, usually named positive and negative. **Frovedis supports only binary Linear SVM classification algorithm.**

The Linear SVM is a standard method for large-scale classification tasks. It is a linear method with the loss function given by the **hinge loss**:

$$L(\mathbf{w}; \mathbf{x}, y) := \max\{0, 1 - y\mathbf{w}^T \mathbf{x}\}$$

During training, the input \mathbf{X} is the training data and \mathbf{y} are their corresponding label values (Frovedis supports any values as for labels, but internally it encodes the input binary labels to -1 and 1, before training at Frovedis server) which we want to predict. \mathbf{w} is the linear model (also known as weight) which uses a single weighted sum of features to make a prediction. Linear SVM supports ZERO, L1 and L2 regularization to address the overfit problem.

The gradient of the hinge loss is: $-y \cdot \mathbf{x}$, if $y\mathbf{w}^T \mathbf{x} < 1$, 0 otherwise.

The gradient of the L1 regularizer is: **sign(w)**

And The gradient of the L2 regularizer is: **w**

For binary classification problems, the algorithm outputs a binary svm model. Given a new data point, denoted by \mathbf{x} , the model makes predictions based on the value of $\mathbf{w}^T \mathbf{x}$.

By default (threshold=0), if $\mathbf{w}^T \mathbf{x} \geq 0$, then the response is positive (1), else the response is negative (0).

Frovedis provides implementation of linear SVM with **stochastic gradient descent with minibatch**.

The simplest method to solve optimization problems of the form **min f(w)** is gradient descent. Such first-order optimization methods well-suited for large-scale and distributed computation.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn LinearSVC (Support Vector Classification) interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for LinearSVC on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

20.3.1 Detailed Description

20.3.1.1 1. LinearSVC()

Parameters

penalty: A string object containing the regularizer type to use. Currently none, l1 and l2 are supported by Frovedis. (Default: 'l2')

loss: A string object containing the loss function type to use. Unlike Scikit-learn, currently it supports only hinge loss. (Default: 'hinge')

dual: An unused parameter. (Default: True)

tol: A double (float64) parameter specifying the convergence tolerance value. It must be zero or a positive value. (Default: 1e-4)

C: A positive float parameter, also called as inverse of regularization strength. (Default: 1.0)

multi_class: An unused parameter. (Default: 'ovr')

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the decision function. (Default: True)

intercept_scaling: An unused parameter. (Default: 1)

class_weight: An unused parameter. (Default: 'None')

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

random_state: An unused parameter. (Default: 'None')

max_iter: A positive integer parameter specifying maximum iteration count. (Default: 1000)

lr_rate: A double (float64) parameter containing the learning rate. (Default: 0.01)

solver: A string object specifying the solver to use. (Default: 'sag')

"sag" can handle L1, L2 or no penalty.

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

Attributes

coef_: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)). It is the weights assigned to the features. It has shape **(1, n_features)**.

classes_: It is a python ndarray(any type) of unique labels given to the classifier during training. It has shape **(n_classes,)**.

intercept_: It is a python ndarray(float or double (float64) values depending on input matrix data type) and has shape **(1,)**.

n_iter: It is a python ndarray of shape **(1,)** and has integer data. It is used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a LinearSVC object with the given parameters.

The parameters: "dual", "intercept_scaling", "class_weight", "multi_class" and "random_state" are simply kept to make the interface uniform to Scikit-learn LinearSVC module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns "self" reference.

20.3.1.2 2. fit(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisColmajorMatrix for dense data. It has shape **(n_samples, n_features)**.

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape **(n_samples,)**.

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of **(n_samples,)**.

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a linear svm model with specified regularization with those data at frovedis server.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# fitting input matrix and label on LinearSVC object
from frovedis.ml.lib.svm import LinearSVC
svm = LinearSVC().fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# and for scipy sparse data, FrovedisCRSMMatrix should be used.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)

# Linear SVC with pre-constructed frovedis-like inputs
from frovedis.ml.lib.svm import LinearSVC
svm = LinearSVC().fit(cmat,dlbl)
```

Return Value

It simply returns “self” reference.

20.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
svm.predict(mat)
```

Output:

```
[0 0 0 ... 0 0 1]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix

# predicting on LinearSVC using frovedis-like input
svm.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[0 0 0 ... 0 0 1]
```

Return Value

It returns a numpy array of double (float64) type containing the predicted outputs. It has shape (**n_samples**,).

20.3.1.4 4. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the LinearSVC model
svm.load("./out/SVMModel")
```

Return Value

It simply returns “self” instance.

20.3.1.5 5. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
svm.save("./out/SVMModel")
```

The SVMModel contains below directory structure:

SVMModel

```
|——label_map
|——metadata
|——model
```

‘label_map’ contains information about labels mapped with their encoded value.

‘metadata’ represents the detail about model_kind and datatype of training vector.

Here, the model file contains information about model_id, model_kind and datatype of training vector.

This will save the LinearSVC model on the path ‘/out/SVMModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

20.3.1.6 6. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the target labels. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of `self.predict(X)` wrt. `y`.

For example,

```
# calculate mean accuracy score on given test data and labels
svm.score(mat, lbl)
```

Output

0.63

Return Value

It returns an accuracy score of float type.

20.3.1.7 7. `get_params(deep = True)`**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by LinearSVC. It is used to get parameters and their values of LinearSVC class.

For example,

```
print(svm.get_params())
```

Output

```
{'C': 1.0, 'class_weight': None, 'dual': True, 'fit_intercept': True, 'intercept_scaling': 1,
'loss': 'hinge', 'lr_rate': 0.01, 'max_iter': 1000, 'multi_class': 'ovr', 'penalty': 'l2',
'random_state': None, 'solver': 'sag', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

20.3.1.8 8. `set_params(params)`****Parameters**

*****params***: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by LinearSVC, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(svm.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
svm.set_params( penalty='l1', dual=False)
print("Get parameters after setting:")
print(svm.get_params())
```

Output

Get parameters before setting:

```
{'C': 1.0, 'class_weight': None, 'dual': True, 'fit_intercept': True,
'intercept_scaling': 1, 'loss': 'hinge', 'lr_rate': 0.01, 'max_iter': 1000,
'multi_class': 'ovr', 'penalty': 'l2', 'random_state': None, 'solver': 'sag',
'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Get parameters after setting:

```
{'C': 1.0, 'class_weight': None, 'dual': False, 'fit_intercept': True,
'intercept_scaling': 1, 'loss': 'hinge', 'lr_rate': 0.01, 'max_iter': 1000,
'multi_class': 'ovr', 'penalty': 'l1', 'random_state': None, 'solver': 'sag',
'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

20.3.1.9 9. debug_print()

Purpose

It shows the target model information (weight values, intercept, etc.) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
svm.debug_print()
```

Output:

```
----- Weight Vector:: -----
83.7418 122.163 486.84 211.922 1.32991 0.287324 -0.867741 -0.0505454
2.04889 1.16388 0.750738 8.61861 -2.13628 -234.118 0.582984 0.445561
0.353854 0.519177 0.667717 0.547778 89.3196 157.824 499.367
-293.736 1.56023 -0.636429 -2.30027 -0.061839 2.66517 1.15244
Intercept:: 19.3242
Threshold:: 0
```

This output will be visible on server side. It displays the weights, intercept and threshold values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

20.3.1.10 10. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
svm.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

20.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

20.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **LinearSVR in Frovedis**
- **SVC in Frovedis**

Chapter 21

LinearSVR

21.1 NAME

LinearSVR (Support Vector Regression) - A regression algorithm used to predict the binary output with L1 and L2 loss.

21.2 SYNOPSIS

```
class frovedis.mllib.svm.LinearSVR(epsilon=0.0, tol=1e-4, C=1.0, loss='epsilon_insensitive',
                                    fit_intercept=True, intercept_scaling=1, dual=True,
                                    verbose=0, random_state=None, max_iter=1000,
                                    penalty='l2', lr_rate=0.01, solver='sag',
                                    warm_start=False)
```

21.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
load(fname, dtype = None)done
save(fname)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

21.3 DESCRIPTION

Based on support vector machines method, the Linear SVR is an algorithm to solve the regression problems. The Linear SVR algorithm applies linear kernel method and it works well with large datasets. L1 or L2 method can be specified as a loss function in this model.

The model produced by Support Vector Regression depends only on a subset of the training data, because the cost function ignores samples whose prediction is close to their target.

LinearSVR supports ZERO, L1 and L2 regularization to address the overfit problem.

Frovedis provides implementation of LinearSVR with **stochastic gradient descent with minibatch**.

The simplest method to solve optimization problems of the form $\min \mathbf{f}(\mathbf{w})$ is gradient descent. Such first-order optimization methods well-suited for large-scale and distributed computation.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn LinearSVR (Support Vector Regression) interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for LinearSVR on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

21.3.1 Detailed Description

21.3.1.1 1. LinearSVR()

Parameters

epsilon: A zero or positive double (float64) parameter used in the epsilon-insensitive loss function. (Default: 0.0)

tol: A double (float64) parameter specifying the convergence tolerance value. It must be zero or a positive value. (Default: 1e-4)

C: A positive float parameter, it is inversely proportional to regularization strength. (Default: 1.0)

loss: A string object containing the loss function type to use.

Currently, frovedis supports 'epsilon_insensitive' and 'squared_epsilon_insensitive' loss function.

The 'epsilon-insensitive' loss (standard SVR) is the L1 loss, while the squared epsilon-insensitive loss ('squared_epsilon_insensitive') is the L2 loss. (Default: 'epsilon_insensitive')

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the decision function. (Default: True)

intercept_scaling: An unused parameter. (Default: 1)

dual: An unused parameter. (Default: True)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

random_state: An unused parameter. (Default: None)

max_iter: A positive integer parameter specifying maximum iteration count. (Default: 1000)

penalty: A string object containing the regularizer type to use. Currently none, l1 and l2 are supported by Frovedis. (Default: 'l2')

If it is None (not specified explicitly), it will be set as 'ZERO' regularization type.

lr_rate: A positive double (float64) value of parameter containing the learning rate. (Default: 0.01)

solver: A string object specifying the solver to use. (Default: 'sag')

Currently, it only supports 'sag'.

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

Attributes

coef_: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)). It is the weights assigned to the features. It has shape **(1, n_features)**.

intercept_: It is a python ndarray(float or double (float64) values depending on input matrix data type) and has shape **(1,)**.

n_iter_: An integer value used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a LinearSVR object with the given parameters.

The parameters: “intercept_scaling”, “dual” and “random_state” are simply kept to make the interface uniform to Scikit-learn LinearSVR module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

21.3.1.2 2. fit(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisColmajorMatrix for dense data. It has shape **(n_samples, n_features)**.

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape **(n_samples,)**.

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of **(n_samples,)**.

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a LinearSVR model with specified regularization with those data at frovedis server.

For example,

```
# let a sample matrix be 'mat 'and labels be 'lbl'
mat = np.array([[ 5.5, 0.0, 6.5, 2.3],
                 [ 0.0, 7.2, 0.0, 8.4],
                 [ 8.5, 0.0, 6.5, 4.3],
                 [ 0.0, 8.5, 0.1, 9.1]], dtype=np.float64)
lbl = np.array([1.1, 6.2, 1.2, 5.9])
```

```
# fitting input matrix and label on LinearSVR object
from frovedis.mllib.svm import LinearSVR
svr = LinearSVR().fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
and for scipy sparse data, FrovedisCRSMatrix should be used.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)
```

```
# LinearSVR with pre-constructed frovedis-like inputs
from frovedis.mllib.svm import LinearSVR
svr = LinearSVR().fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

21.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
svr.predict(mat)
```

Output:

```
[-181.66076961 -162.62098062 -166.05339001 ... -170.80953572 -169.6636383 -171.76112166]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix

# predicting on LinearSVR using frovedis-like input
svr.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[-181.66076961 -162.62098062 -166.05339001 ... -170.80953572 -169.6636383 -171.76112166]
```

Return Value

It returns a numpy array of double (float64) type containing the predicted outputs. It has shape (**n_samples**,).

21.3.1.4 4. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the LinearSVR model
svr.load("./out/SVRModel")
```

Return Value

It simply returns “self” reference.

21.3.1.5 5. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
svr.save("./out/SVRModel")
```

The SVRModel contains below directory structure:

SVRModel

```
|— metadata
|— model
```

‘metadata’ represents the detail about model_kind and datatype of training vector.

Here, the model file contains information about model_id, model_kind and datatype of training vector.

This will save the LinearSVR model on the path ‘/out/SVRModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

21.3.1.6 6. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the target labels. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. $R2(r\text{-squared})$ of self.predict(X) wrt. y.

The coefficient ‘R2’ is defined as $(1 - (u/v))$,

where ‘u’ is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and ‘v’ is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
# calculate R2 score on given test data and labels
svr.score(mat, lbl)
```

Output

0.97

Return Value

It returns an R2 score of float type.

21.3.1.7 7. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by LinearSVR. It is used to get parameters and their values of LinearSVR class.

For example,

```
print(svr.get_params())
```

Output

```
{'C': 1.0, 'dual': True, 'epsilon': 0.0, 'fit_intercept': True, 'intercept_scaling': 1,
'loss': 'epsilon_insensitive', 'lr_rate': 0.01, 'max_iter': 1000, 'penalty': 'l2',
'random_state': None, 'solver': 'sag', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

21.3.1.8 8. set_params(params)****Parameters**

****params**: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by LinearSVR, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(svr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
svr.set_params( penalty = 'l1', dual = False)
print("Get parameters after setting:")
print(svr.get_params())
```

Output

```
Get parameters before setting:
{'C': 1.0, 'dual': True, 'epsilon': 0.0, 'fit_intercept': True, 'intercept_scaling': 1,
'loss': 'epsilon_insensitive', 'lr_rate': 0.01, 'max_iter': 1000, 'penalty': 'l2',
'random_state': None, 'solver': 'sag', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
Get parameters before setting:
{'C': 1.0, 'dual': False, 'epsilon': 0.0, 'fit_intercept': True, 'intercept_scaling': 1,
'loss': 'epsilon_insensitive', 'lr_rate': 0.01, 'max_iter': 1000, 'penalty': 'l1',
'random_state': None, 'solver': 'sag', 'tol': 0.0001, 'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

21.3.1.9 9. debug_print()**Purpose**

It shows the target model information(weight values and intercept) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
svr.debug_print()
```

Output:

```
----- Weight Vector:: -----
0.406715 0.413736 0.440404 0.539082 0.536528 0.51577 0.157345 0.520861
0.513352 -0.341082 0.460518 -0.292171 0.433433
Intercept:: 1.00832
```

This output will be visible on server side. It displays the weights and intercept values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

21.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
svr.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

21.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

21.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisCRSMatrix**

- **Introduction to FrovedisDvector**
- **LinearSVC in Frovedis**
- **SVC in Frovedis**

Chapter 22

Logistic Regression

22.1 NAME

Logistic Regression - A classification algorithm to predict the binary and multi-class output with logistic loss.

22.2 SYNOPSIS

```
class frovedis.mllib.linear_model.LogisticRegression(penalty='l2', dual=False, tol=1e-4,
                                                    C=100.0, fit_intercept=True,
                                                    intercept_scaling=1, class_weight=None,
                                                    random_state=None, solver='lbfgs',
                                                    max_iter=1000, multi_class='auto',
                                                    verbose=0, warm_start=False,
                                                    n_jobs=1, l1_ratio=None,
                                                    lr_rate=0.01, use_shrink=False)
```

22.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
predict_proba(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

22.3 DESCRIPTION

Classification aims to divide the items into categories. The most common classification type is binary classification, where there are two categories, usually named positive and negative. The other is multinomial

classification, where there are more than two categories. **Frovedis supports both binary and multinomial logistic regression algorithms.** For multinomial classification, it uses softmax probability.

Logistic regression is widely used to predict a binary response. It is a linear method with the loss function given by the **logistic loss**:

$$L(\mathbf{w}; \mathbf{x}, y) := \log(1 + \exp(-y\mathbf{w}^T\mathbf{x}))$$

During training, the input \mathbf{X} is the training data and \mathbf{y} is the corresponding label values (Frovedis supports any values as for labels, but internally it encodes the input binary labels to -1 and 1, and input multinomial labels to 0, 1, 2, ..., N-1 (where N is the no. of classes) before training at Frovedis server) which we want to predict. \mathbf{w} is the linear model (also called as weight) which uses a single weighted sum of features to make a prediction. **Frovedis Logistic Regression supports ZERO, L1 and L2 regularization to address the overfit problem. However, LBFGS solver supports only L2 regularization.**

The gradient of the logistic loss is: $-\mathbf{y}(1 - 1 / (1 + \exp(-y\mathbf{w}^T\mathbf{x}))).\mathbf{x}$

The gradient of the L1 regularizer is: $\text{sign}(\mathbf{w})$

And, the gradient of the L2 regularizer is: \mathbf{w}

For binary classification problems, the algorithm outputs a binary logistic regression model. Given a new data point, denoted by \mathbf{x} , the model makes predictions by applying the logistic function:

$$f(z) := 1 / (1 + \exp(-z))$$

Where $z = \mathbf{w}^T\mathbf{x}$. By default (threshold=0.5), if $f(\mathbf{w}^T\mathbf{x}) > 0.5$, the response is positive (1), else the response is negative (0).

In the multiclass case, the training algorithm uses the one-vs-rest (OvR) scheme. Currently the “multinomial” option is supported only by the “sag” solvers.

Frovedis provides implementation of logistic regression with two different optimizers:

(1) stochastic gradient descent with minibatch

(2) LBFGS optimizer

They can handle both dense and sparse input.

The simplest method to solve optimization problems of the form $\min \mathbf{f}(\mathbf{w})$ is gradient descent. Such first-order optimization methods well-suited for large-scale and distributed computation. Whereas, L-BFGS is an optimization algorithm in the family of quasi-Newton methods to solve the optimization problems of the similar form.

Like the original BFGS, L-BFGS (Limited Memory BFGS) uses an estimation to the inverse Hessian matrix to steer its search through feature space, but where BFGS stores a dense $n \times n$ approximation to the inverse Hessian (n being the number of features in the problem), L-BFGS stores only a few vectors that represent the approximation implicitly. L-BFGS often achieves rapider convergence compared with other first-order optimization.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn Logistic Regression interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Logistic Regression on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

22.3.1 Detailed Description

22.3.1.1 1. LogisticRegression()

Parameters

penalty: A string object containing the regularizer type to use. Currently none, l1 and l2 are supported by Frovedis. (Default: 'l2')

dual: A boolean parameter. (unused)

tol: A double (float64) type value specifying the convergence tolerance value. It must be zero or a positive value. (Default: 1e-4)

C: A positive float parameter, it is the inverse of regularization strength. Like in support vector machines, smaller values specify stronger regularization. (Default: 100.0)

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the decision function. (Default: True)

intercept_scaling: An unused parameter. (Default: 1)

class_weight: An unused parameter. (Default: None)

random_state: An unused parameter. (Default: None)

solver: A string object specifying the solver to use. (Default: 'lbfgs')

It can be "sag" for frovedis side stochastic gradient descent or "lbfgs" for frovedis side LBFGS optimizer when optimizing the logistic regression model.

"sag" can handle L1, L2 or no penalty.

"lbfgs" can handle only L2 penalty.

max_iter: A positive integer value specifying maximum iteration count. (Default: 1000)

multi_class: A string object specifying the type of classification. - If it is "auto" or "ovr", then a binary classification is selected when N = 2, otherwise multinomial classification is selected (where N is the no. of classes in training labels). - If it is "multinomial", then it always selects a multinomial problem (even when N = 2). Only "sag" solvers support multinomial classification currently. (Default: 'auto')

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not explicitly specified). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

n_jobs: An unused parameter. (Default: 1)

l1_ratio: An unused parameter. (Default: None)

lr_rate(alpha): A double (float64) parameter containing the learning rate. (Default: 0.01)

use_shrink: A boolean parameter applicable only for "sag" solver with "sparse" input (X). When set to True for sparse input, it can improve training performance by reducing communication overhead across participating processes. (Default: False)

Attributes

coef_: It is a python ndarray (float or double (float64) values depending on input matrix data type) of coefficient of the features in the decision function. It has shape (1, n_features) when the given problem is "binary" and (n_classes, n_features) when it is a "multinomial" problem.

intercept_(bias): It is a python ndarray(float or double (float64) values depending on input matrix data type) If fit_intercept is set to False, the intercept is set to zero. It has shape (1,) when the given problem is "binary" and (n_classes) when its "multinomial" problem.

classes_: It is a python ndarray (any type) of unique labels given to the classifier during training. It has shape (n_classes,).

n_iter_: It is a python ndarray of shape (1,) and has integer data. It is used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a Logistic Regression object with the given parameters.

The parameters: "dual", "intercept_scaling", "class_weight", "random_state", "n_jobs" and "l1_ratio" are

simply kept to make the interface uniform to the Scikit-learn Logistic Regression module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

22.3.1.2 2. fit(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisColmajorMatrix for dense data.

y: Any python array-like object or an instance of FrovedisDvector containing target labels. It has shape (n_samples,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (n_samples,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a logistic regression model with specified regularization with those data at frovedis server.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# fitting input matrix and label on LogisticRegression object
from frovedis.mllib.linear_model import LogisticRegression
lr = LogisticRegression(solver = 'lbfgs').fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# Logistic Regression with pre-constructed frovedis-like inputs
from frovedis.mllib.linear_model import LogisticRegression
lr = LogisticRegression(solver = 'lbfgs').fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

22.3.1.3 3. predict(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
# predicting on lbfgs logistic regression model
lr.predict(mat)
```

Output

```
[0 0 0 ... 0 0 1]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
```

```
# predicting on lbfgs logistic regression model using pre-constructed input
lr.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[0 0 0 ... 0 0 1]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**,) containing the predicted outputs.

22.3.1.4 4. predict_proba(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server. But unlike predict(), it returns the softmax probability matrix of shape (**n_samples**, **n_classes**) containing the probability of each class in each sample.

For example,

```
# finds the probability sample for each class in the model
lr.predict_proba(mat)
```

Output

```
[[1.46990588e-19 1.00000000e+00]
 [7.23344224e-10 9.99999999e-01]
 [8.43160984e-10 9.99999999e-01]
 ...
 [4.03499383e-04 9.99596501e-01]
 [3.03132738e-13 1.00000000e+00]]
```

```
[6.14030540e-03 9.93859695e-01]]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix

# finds the probability sample for each class in the model
lr.predict_proba(mat)
```

Output

```
[[1.46990588e-19 1.00000000e+00]
 [7.23344224e-10 9.99999999e-01]
 [8.43160984e-10 9.99999999e-01]
 ...
 [4.03499383e-04 9.99596501e-01]
 [3.03132738e-13 1.00000000e+00]
 [6.14030540e-03 9.93859695e-01]]
```

Return Value

It returns an ndarray of float or double (float64) type and of shape `(n_samples, n_classes)` containing the prediction probability values.

22.3.1.5 5. `score(X, y, sample_weight = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data.

y: Any python array-like object containing true labels for X. It has shape `(n_samples,)`.

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of `(n_samples,)`.

When it is `None` (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: `None`)

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of `self.predict(X)` wrt. `y`.

For example,

```
# calculate mean accuracy score on given test data and labels
lr.score(mat, lbl)
```

Output

```
0.96
```

Return Value

It returns an accuracy score of float type.

22.3.1.6 6. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If `True`, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: `True`)

Purpose

This method belongs to the BaseEstimator class inherited by LogisticRegression. It is used to get parameters and their values of LogisticRegression class.

For example,

```
print(lr.get_params())
```

Output

```
{'C': 100.0, 'class_weight': None, 'dual': False, 'fit_intercept': True,
'intercept_scaling': 1, 'l1_ratio': None, 'lr_rate': 0.01, 'max_iter': 1000,
'multi_class': 'auto', 'n_jobs': 1, 'penalty': 'l2', 'random_state': None,
'solver': 'sag', 'tol': 0.0001, 'use_shrink': False, 'verbose': 0,
'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

22.3.1.7 7. set_params(params)****Parameters**

****params:** All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by LogisticRegression, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(lr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
lr.set_params(solver='lbfgs', max_iter = 10000)
print("get parameters after setting:")
print(lr.get_params())
```

Output

```
get parameters before setting:
{'C': 100.0, 'class_weight': None, 'dual': False, 'fit_intercept': True,
'intercept_scaling': 1, 'l1_ratio': None, 'lr_rate': 0.01, 'max_iter': 1000,
'multi_class': 'auto', 'n_jobs': 1, 'penalty': 'l2', 'random_state': None,
'solver': 'sag', 'tol': 0.0001, 'use_shrink': False, 'verbose': 0,
'warm_start': False}
get parameters after setting:
{'C': 100.0, 'class_weight': None, 'dual': False, 'fit_intercept': True,
'intercept_scaling': 1, 'l1_ratio': None, 'lr_rate': 0.01, 'max_iter': 10000,
'multi_class': 'auto', 'n_jobs': 1, 'penalty': 'l2', 'random_state': None,
'solver': 'lbfgs', 'tol': 0.0001, 'use_shrink': False, 'verbose': 0,
'warm_start': False}
```

Return Value

It simply returns “self” reference.

22.3.1.8 8. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
lr.load("./out/LRModel")
```

Return Value

It simply returns “self” reference.

22.3.1.9 9. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the logistic regression model
lr.save("./out/LRModel")
```

This will save the logistic regression model on the path ‘/out/LRModel’. It would raise exception if the directory already exists with same name.

The ‘LRModel’ directory has

LRModel

```
|—label_map
|—metadata
|—model
   |—model
   |—type
```

‘label_map’ contains information about labels mapped with their encoded value.

The metadata file contains the number of classes, model kind, input datatype used for trained model.

Here, the **model** directory contains information about weights, intercept, threshold and thier datatype.

Return Value

It returns nothing.

22.3.1.10 10. debug_print()**Purpose**

It shows the target model information (weight values, intercept, etc.) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
lr.debug_print()
```

Output

```
----- Weight Vector:: -----
25.4745 47.8416 155.732 190.863 0.271114 0.0911008 -0.151433 -0.0785512 0.511576
0.203452 0.199293 3.8659 1.22203 -42.3556 0.0239707 0.0395711 0.0389786 0.017432
0.0647208 0.0105295 24.7162 60.7113 150.789 -148.921 0.354222 0.104251 -0.202345
-0.0363726 0.734499 0.22635
Intercept:: 60.7742
Threshold:: 0.5
```

This output will be visible on server side. It displays the weights, intercept, etc. values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

22.3.1.11 11. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
lr.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

22.3.1.12 12. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the logistic regression model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

22.4 SEE ALSO

- Introduction to FrovedisRowmajorMatrix
- Introduction to FrovedisColmajorMatrix
- Introduction to FrovedisCRSMatrix
- Introduction to FrovedisDvector
- Linear Regression in Frovedis

Chapter 23

Multinomial Naive Bayes

23.1 NAME

Multinomial Naive Bayes - One of the variations of Naive Bayes algorithm. It is a multinomial classification algorithm.

23.2 SYNOPSIS

```
class frovedis.mllib.naive_bayes.MultinomialNB(alpha=1.0, fit_prior=True, class_prior=None,
                                              verbose=0)
```

23.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
predict_proba(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
debug_print()
load(fname, dtype = None)
save(fname)
release()
is_fitted()
```

23.3 DESCRIPTION

Naive Bayes classifier for multinomial models. The multinomial Naive Bayes classifier is suitable for classification with discrete features (e.g., word counts for text classification).

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn MultinomialNB interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python

data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for MultinomialNB on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

23.3.1 Detailed Description

23.3.1.1 1. MultinomialNB()

Parameters

alpha: A positive double (float64) smoothing parameter. It must be greater than or equal to 1. (Default: 1.0)

fit_prior: A boolean parameter specifying whether to learn class prior probabilities or not. If False, a uniform prior will be used. (Default: True)

class_prior: A numpy ndarray of double (float64) type values and must be of shape (**n_classes**,). It gives prior probabilities of the classes. (Default: None)

When it is None (not specified explicitly), the priors are adjusted according to the data.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Attributes

class_log_prior: A python ndarray of double (float64) type values and has shape (**n_classes**,). It contains log probability of each class (smoothed).

feature_log_prob: A python ndarray of double (float64) type values and has shape (**n_classes**, **n_features**). It contains empirical log probability of features given a class.

class_count: A python ndarray of double (float64) type values and has shape (**n_classes**,). It contains the number of samples encountered for each class during fitting. This value is weighted by the sample weight when provided.

classes: A python ndarray of double (float64) type values and has shape (**n_classes**,). It contains the of unique labels given to the classifier during training.

feature_count: A python ndarray of double (float64) type values and has shape (**n_classes**, **n_features**). It contains the number of samples encountered for each (class, feature) during fitting. This value is weighted by the sample weight when provided.

coef: A python ndarray of double (float64) type values.

- If 'classes' is 2, then the shape (**1**, **n_features**)
- If 'classes' is more then 2, then the shape is (**n_classes**, **n_features**).

It mirrors 'feature_log_prob' for interpreting MultinomialNB as a linear model.

intercept: A python ndarray of double (float64) type values.

- If 'classes' are 2, then the shape (**1**,)
- If 'classes' is more then 2, then the shape is (**n_classes**,).

It mirrors 'class_log_prior' for interpreting MultinomialNB as a linear model.

Purpose

It initializes a MultinomialNB object with the given parameters.

Return Value

It simply returns "self" reference.

23.3.1.2 2. fit(X, y, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It accepts the training matrix (X) with labels (y) and trains a MultinomialNB model.

For example,

```
# loading sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# fitting input matrix and label on MultinomialNB object
from frovedis.mllib.naive_bayes import MultinomialNB
mnb = MultinomialNB().fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
rmat = FrovedisRowmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# MultinomialNB with pre-constructed frovedis-like inputs
from frovedis.mllib.naive_bayes import MultinomialNB
mnb = MultinomialNB().fit(rmat, dlbl)
```

Return Value

It simply returns “self” reference.

23.3.1.3 3. predict(X)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
# predicting on MultinomialNB model
mnb.predict(mat)
```

Output

```
[1 1 1 1 1]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# predicting on MultinomialNB model using pre-constructed input
mnb.predict(rmat)
```

Output

```
[1 1 1 1 1]
```

Return Value

It returns a numpy array of long (int64) type and of shape **(n_samples,)** containing the predicted outputs.

23.3.1.4 4. `predict_proba(X)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data. It has shape **(n_samples, n_features)**.

Purpose

Perform classification on an array and return probability estimates for the test vector **X**.

It accepts the test feature matrix (**X**) in order to make prediction on the trained model at `frovedis` server. Unlike `sklearn`, it performs the classification on an array and returns the probability estimates for the test feature matrix (**X**).

For example,

```
# finds the probability sample for each class in the model
mnb.predict_proba(mat)
```

Output

```
[[1.00000000e+00 1.31173349e-73]
 [1.00000000e+00 3.02940294e-46]
 [1.00000000e+00 1.60233859e-37]
 ...
 [9.99537526e-01 4.62474254e-04]
 [1.00000000e+00 2.74000590e-29]
 [3.87284789e-18 1.00000000e+00]]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# finds the probability sample for each class in the model
mnb.predict_proba(rmat)
```

Output

```
[[1.00000000e+00 1.31173349e-73]
 [1.00000000e+00 3.02940294e-46]
 [1.00000000e+00 1.60233859e-37]
 ...
 [9.99537526e-01 4.62474254e-04]
 [1.00000000e+00 2.74000590e-29]
 [3.87284789e-18 1.00000000e+00]]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**, **n_classes**) containing the predicted probability values.

23.3.1.5 5. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the target labels. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of self.predict(X) wrt. y.

For example,

```
mnb.score(mat, lbl)
```

Output

```
0.89
```

Return Value

It returns an accuracy score of double (float64) type.

23.3.1.6 6. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by MultinomialNB. It is used to get parameters and their values of MultinomialNB class.

For example,

```
print(mnb.get_params())
```

Output

```
{'alpha': 1.0, 'class_prior': None, 'fit_prior': True, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

23.3.1.7 7. set_params(**params)

Parameters

****params**: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by MultinomialNB, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(mnb.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
mnb.set_params(fit_prior = False)
print("Get parameters after setting:")
print(mnb.get_params())
```

Output

```
Get parameters before setting:
{'alpha': 1.0, 'class_prior': None, 'fit_prior': True, 'verbose': 0}
Get parameters after setting:
{'alpha': 1.0, 'class_prior': None, 'fit_prior': False, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

23.3.1.8 8. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information such as theta, cls_count, feature_count, label, pi, type to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model stored previously from the specified file (having little-endian binary data).

For example,

```
mnb.load("./out/MNBModel", dtype = np.float64)
```

Return Value

It simply returns “self” reference.

23.3.1.9 9. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information(label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the MultinomialNB model
mnb.save("./out/MNBModel")
```

The MNBModel contains below directory structure:

MNBModel

```
|—label_map
|—metadata
|—model
    |—cls_count
    |—feature_count
    |—label
    |—pi
    |—theta
    |—type
```

‘label_map’ contains information about labels mapped with their encoded value.

‘metadata’ represents the detail about model_kind and datatype of training vector.

Here, the **model** directory contains information about cls_count, feature_count, label, pi, **theta** and type.

This will save the naive bayes model on the path ‘/out/MNBModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

23.3.1.10 10. debug_print()**Purpose**

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
mnb.debug_print()
```

Output

```
model_type: multinomial
binarize: 0
feature_count: 3702.12 4580.24 24457.5 207416 21.8145 ... 65.2141 59.3469 26.5766 96.4778 28.3608
theta: node = 0, local_num_row = 2, local_num_col = 30, val = -5.08709 -4.8743 -3.19929 -1.06154
-10.1766
...
-8.83939 -8.93218 -9.71532 -8.45266 -9.65263
pi: -0.987294 -0.466145
label: 0 1
class count: 212 357
```

This output will be visible on server side. It displays the target model information like model_type, binarize, feature_count, theta, pi, etc. values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

23.3.1.11 11. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
mnb.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

23.3.1.12 12. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the naive bayes model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

23.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Bernoulli Naive Bayes in Frovedis**

Chapter 24

NearestNeighbors

24.1 NAME

NearestNeighbors - Unsupervised learner for implementing neighbor searches.

24.2 SYNOPSIS

```
class frovedis.mllib.neighbors.NearestNeighbors(n_neighbors=5, radius=1.0, algorithm='auto',
                                                leaf_size=30, metric='euclidean', p=2,
                                                metric_params=None, n_jobs=None, verbose=0,
                                                chunk_size=1.0, batch_fraction=None)
```

24.2.1 Public Member Functions

```
fit(X, y = None)
kneighbors(X = None, n_neighbors = None, return_distance = True)
kneighbors_graph(X = None, n_neighbors = None, mode = 'connectivity')
radius_neighbors(X = None, radius = None, return_distance = True)
radius_neighbors_graph(X = None, radius = None, mode = 'connectivity')
get_params(deep = True)
set_params(**params)
save(fname)
load(fname)
debug_print()
release()
is_fitted()
```

24.3 DESCRIPTION

Unsupervised nearest neighbors is the foundation of many other learning methods, notably manifold learning and spectral clustering.

The principle behind nearest neighbor methods is to find a predefined number of training samples closest in distance to the new point and predict the label from these. The number of samples can be a user-defined

constant (k-nearest neighbor learning), or vary based on the local density of points (radius-based neighbor learning). In general, the distance can be any metric measure: standard Euclidean distance is the most common choice. Neighbors-based methods are known as non-generalizing machine learning methods, since they simply “remember” all of its training data.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn NearestNeighbors interface, but it doesn’t have any dependency with Scikit-learn. It can be used simply even if the system doesn’t have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for NearestNeighbors on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When operations like `kneighbors()` will be required on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

24.3.1 Detailed Description

24.3.1.1 1. NearestNeighbors()

Parameters

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for ‘kneighbors’ queries. It must be within the range of 0 and `n_samples`. (Default: 5)

radius: A positive float parameter, specifying the range of parameter space to use by default for ‘radius_neighbors’ queries. (Default: 1.0)

algorithm: A string object parameter, specifying the algorithm used to compute the nearest neighbors. (Default: ‘auto’)

When it is ‘auto’, it will be set as ‘brute’(brute-force search approach). Unlike Scikit-learn, currently it supports only ‘brute’.

leaf_size: An unused parameter. (Default: 30)

metric: A string object parameter, specifying the distance metric to use for the tree. (Default: ‘euclidean’) Currently it only supports ‘euclidean’, ‘seuclidean’ and ‘cosine’ distance.

p: An unused parameter. (Default: 2)

metric_params: An unused parameter. (Default: None)

n_jobs: An unused parameter. (Default: None)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1(for DEBUG mode) or 2(for TRACE mode) for getting training time logs from frovedis server.

chunk_size: A positive float parameter, specifying the amount of data (in megabytes) to be processed in one time. (Default: 1.0)

batch_fraction: A positive double (float64) parameter used to calculate the batches of specific size. These batches are used to construct the distance matrix. It must be within the range of 0.0 to 1.0. (Default: None) When it is None (not specified explicitly), it will be set as `np.finfo(np.float64).max` value.

Purpose

It initializes a NearestNeighbors object with the given parameters.

The parameters: “leaf_size”, “p”, “metric_params” and “n_jobs” are simply kept in to make the interface uniform to the Scikit-learn NearestNeighbors module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

24.3.1.2 2. fit(X, y = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

Fit the model using X as training data

For example,

```
# loading sample data
samples = np.loadtxt("./input/knn_data.txt", dtype = np.float64)

# fitting input data on NearestNeighbors object
from frovedis.mllib.neighbors import NearestNeighbors
knn = NearestNeighbors(n_neighbors = 3, radius = 2.0, algorithm = 'brute',
                      metric = 'euclidean')
knn.fit(samples)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample data
samples = np.loadtxt("./input/knn_data.txt", dtype = np.float64)

# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# fitting input data on NearestNeighbors object
from frovedis.mllib.neighbors import NearestNeighbors
knn = NearestNeighbors(n_neighbors = 3, radius = 2.0, algorithm = 'brute',
                      metric = 'euclidean')
knn.fit(rmat)
```

Return Value

It simply returns “self” reference.

24.3.1.3 3. kneighbors(X = None, n_neighbors = None, return_distance = True)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_queries**, **n_features**), where ‘n_queries’ is the number of rows in the test data. (Default: None)

When it is None (not specified explicitly), it will be training data (X) used as input in fit().

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for ‘kneighbors’ queries. It must be within the range of 0 and n_queries. (Default: None)

When it is None (not specified explicitly), it will be ‘n_neighbors’ value used during NearestNeighbors object creation.

return_distance: A boolean parameter specifying whether or not to return the distances. (Default: True) If set to False, it will not return distances. Then, only indices are returned by this method.

Purpose

It finds the k-Neighbors of a point and returns the indices of neighbors and distances to the neighbors of each point.

For example,

```
distances, indices = knn.kneighbors(samples)
print('distances')
print(distances)
print('indices')
print(indices)
```

Output

```
distances
[[0.          1.          2.23606798]
 [0.          1.          1.41421356]
 [0.          1.41421356  2.23606798]
 [0.          1.          2.23606798]
 [0.          1.          1.41421356]
 [0.          1.41421356  2.23606798]]
indices
[[0 1 2]
 [1 0 2]
 [2 1 0]
 [3 4 5]
 [4 3 5]
 [5 4 3]]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the computation of indices and distances at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

distances, indices = knn.kneighbors(rmat)

# Here FrovedisRowmajorMatrix().debug_print() is used
print('distances')
distances.debug_print()

# Here FrovedisRowmajorMatrix().debug_print() is used
print('indices')
indices.debug_print()
```

Output

```
distances
matrix:
num_row = 6, num_col = 3
node 0
```

```

node = 0, local_num_row = 6, local_num_col = 3, val = 0 1 2.23607 0 1 1.41421 0 1.41421
2.23607 0 1 2.23607 0 1 1.41421 0 1.41421 2.23607
indices
matrix:
num_row = 6, num_col = 3
node 0
node = 0, local_num_row = 6, local_num_col = 3, val = 0 1 2 1 0 2 2 1 0 3 4 5 4 3 5 5 4 3

```

Return Value

1. When test data and training data used by fitted model are python native input:

- **distances**: A numpy array of float or double(float64) type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data. It is only returned by `kneighbors()` if `return_distance = True`.
- **indices**: A numpy array of int64 type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data.

2. When either test data or training data used by fitted model is frovedis-like input:

- **distances**: A FrovedisRowmajorMatrix of float or double(float64) type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data. It is only returned by `kneighbors()` if `return_distance = True`.
- **indices**: A FrovedisRowmajorMatrix of int64 type values. It has shape (**n_queries**, **n_neighbors**), where 'n_queries' is the number of rows in the test data.

24.3.1.4 4. `kneighbors_graph(X = None, n_neighbors = None, mode = 'connectivity')`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_queries**, **n_features**), where 'n_queries' is the number of rows in the test data. (Default: None)
When it is None (not specified explicitly), it will be training data (X) used as input in `fit()`.

n_neighbors: A positive integer parameter, specifying the number of neighbors to use by default for 'kneighbors' queries. It must be within the range of 0 and `n_queries`. (Default: None)

When it is None (not specified explicitly), it will be 'n_neighbors' value used during NearestNeighbors object creation.

mode: A string object parameter which can be either 'connectivity' or 'distance'. It specifies the type of returned matrix.

For 'connectivity', it will return the connectivity matrix with ones and zeros, whereas for 'distance', the edges are euclidean distance between points. Type of distance depends on the selected 'metric' value in NearestNeighbors class. (Default: 'connectivity')

Purpose

It computes the (weighted) graph of k-Neighbors for points in X.

For example, when `mode = 'connectivity'`

```

# Here 'mode = connectivity' by default
graph = knn.kneighbors_graph(samples)
print('kneighbors graph')
print(graph)

```

Output

```

kneighbors graph
(0, 0)      1.0
(0, 1)      1.0
(0, 2)      1.0
(1, 1)      1.0

```

```
(1, 0)      1.0
(1, 2)      1.0
(2, 2)      1.0
(2, 1)      1.0
(2, 0)      1.0
(3, 3)      1.0
(3, 4)      1.0
(3, 5)      1.0
(4, 4)      1.0
(4, 3)      1.0
(4, 5)      1.0
(5, 5)      1.0
(5, 4)      1.0
(5, 3)      1.0
```

For example, when `mode = 'distance'`

```
# Here 'mode = distance'
graph = knn.kneighbors_graph(samples, mode = 'distance')
print('kneighbors graph')
print(graph)
```

Output

```
kneighbors graph
(0, 0)      0.0
(0, 1)      1.0
(0, 2)      2.23606797749979
(1, 1)      0.0
(1, 0)      1.0
(1, 2)      1.4142135623730951
(2, 2)      0.0
(2, 1)      1.4142135623730951
(2, 0)      2.23606797749979
(3, 3)      0.0
(3, 4)      1.0
(3, 5)      2.23606797749979
(4, 4)      0.0
(4, 3)      1.0
(4, 5)      1.4142135623730951
(5, 5)      0.0
(5, 4)      1.4142135623730951
(5, 3)      2.23606797749979
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the graph construction at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCSRMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# Here 'mode = connectivity' by default
graph = knn.kneighbors_graph(rmat)
print('kneighbors graph')
```

```
# Here FrovedisCRSMatrix().debug_print() is used
graph.debug_print()
```

Output

```
graph
Active Elements:  18
matrix:
num_row = 6, num_col = 6
node 0
local_num_row = 6, local_num_col = 6
val : 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
idx : 0 1 2 1 0 2 2 1 0 3 4 5 4 3 5 4 3
off : 0 3 6 9 12 15 18
```

Return Value

- **When test data and training data used by fitted model are python native input :**

It returns a scipy sparse csr matrix of float or double (float64) type values. It has shape (**n_queries**, **n_samples_fit**), where 'n_queries' is the number of rows in the test data and 'n_samples_fit' is the number of samples in the fitted data.

- **When either test data or training data used by fitted model is frovedis-like input:**

It returns a FrovedisCRSMatrix of float or double(float64) type values. It has shape (**n_queries**, **n_samples_fit**), where 'n_queries' is the number of rows in the test data and 'n_samples_fit' is the number of samples in the fitted data.

24.3.1.5 5. radius_neighbors(X = None, radius = None, return_distance = True)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**). (Default: None)

When it is None (not specified explicitly), it will be training data (X) used as input in fit().

radius: A positive float parameter, specifying the limiting distance of neighbors to return. (Default: None)

When it is None (not specified explicitly), it will be 'radius' value used in NearestNeighbors object creation.

return_distance: A boolean parameter specifying whether or not to return the distances. (Default: True)

If set to False, it will not return distances. Then, only indices are returned by this method.

Purpose

It finds the neighbors within a given radius of a point or points and returns indices and distances to the neighbors of each point.

For example,

```
dist, ind = knn.radius_neighbors(samples)
print('radius neighbors')
for i in range(len(dist)):
    print(dist[i])
    print(ind[i])
```

Output

```
radius neighbors
[0. 1.]
[0 1]
[1.          0.          1.41421356]
[0 1 2]
[1.41421356 0.          ]
```

```
[1 2]
[0. 1.]
[3 4]
[1.          0.          1.41421356]
[3 4 5]
[1.41421356 0.          ]
[4 5]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the computation of indices and distances at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

rad_nbs = knn.radius_neighbors(rmat)
print('radius neighbors')

# Here FrovedisCRSMatrix().debug_print() is used
rad_nbs.debug_print()
```

Output

```
radius neighbors
Active Elements:  14
matrix:
num_row = 6, num_col = 6
node 0
local_num_row = 6, local_num_col = 6
val : 0 1 1 0 1.41421 1.41421 0 0 1 1 0 1.41421 1.41421 0
idx : 0 1 0 1 2 1 2 3 4 3 4 5 4 5
off : 0 2 5 7 9 12 14
```

Return Value

1. **When test data and training data used by fitted model are python native input :**
 - **distance:** A python list of float or double(float64) type values and has length `n_samples`. It is only returned by `radius_neighbors()` if `return_distance = True`.
 - **indices:** A python list of float or double(float64) type values and has length `n_samples`.
2. **When either test data or training data used by fitted model is frovedis-like input:**
It returns a `FrovedisCRSMatrix` of shape `(n_samples, n_samples_fit)`, where ‘`n_samples_fit`’ is the number of samples in the fitted data.

24.3.1.6 6. `radius_neighbors_graph(X = None, radius = None, mode = ‘connectivity’)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data of float or double (float64) type. It has shape `(n_samples, n_features)`. (Default: `None`)

When it is `None` (not specified explicitly), it will be training data (`X`) used as input in `fit()`.

radius: A positive float parameter, specifying the limiting distance of neighbors to return. (Default: `None`)
When it is `None` (not specified explicitly), it will be ‘radius’ value used in `NearestNeighbors` object creation.

mode: A string object parameter which can be either ‘connectivity’ or ‘distance’. It specifies the type of returned matrix.

For ‘connectivity’, it will return the connectivity matrix with ones and zeros, whereas for ‘distance’, the edges are euclidean distance between points. Type of distance depends on the selected ‘metric’ value in NearestNeighbors class. (Default: ‘connectivity’)

Purpose

It computes the (weighted) graph of Neighbors for points in X.

For example, when mode = ‘connectivity’

```
# Here 'mode = connectivity' by default
rad_graph = knn.radius_neighbors_graph(samples)
print('radius neighbors graph')
print(rad_graph)
```

Output

```
radius neighbors graph
(0, 0)      1.0
(0, 1)      1.0
(1, 0)      1.0
(1, 1)      1.0
(1, 2)      1.0
(2, 1)      1.0
(2, 2)      1.0
(3, 3)      1.0
(3, 4)      1.0
(4, 3)      1.0
(4, 4)      1.0
(4, 5)      1.0
(5, 4)      1.0
(5, 5)      1.0
```

For example, when mode = ‘distance’

```
# Here 'mode = distance'
rad_graph = knn.radius_neighbors_graph(samples, mode = 'distance')
print('radius neighbors graph')
print(rad_graph)
```

Output

```
radius neighbors graph
(0, 0)      0.0
(0, 1)      1.0
(1, 0)      1.0
(1, 1)      0.0
(1, 2)      1.4142135623730951
(2, 1)      1.4142135623730951
(2, 2)      0.0
(3, 3)      0.0
(3, 4)      1.0
(4, 3)      1.0
(4, 4)      0.0
(4, 5)      1.4142135623730951
(5, 4)      1.4142135623730951
(5, 5)      0.0
```

Like in fit(), frovedis-like input can be used to speed-up the computation of graph construction at server side.

For example,

```
# Since "samples" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(samples)

# Here 'mode = connectivity' by default
rad_graph = knn.radius_neighbors_graph(samples)
print('radius neighbors graph')

# Here FrovedisCRSMatrix().debug_print() is used
rad_graph.debug_print()
```

Output

```
radius neighbors graph
Active Elements: 14
matrix:
num_row = 6, num_col = 6
node 0
local_num_row = 6, local_num_col = 6
val : 1 1 1 1 1 1 1 1 1 1 1 1 1 1
idx : 0 1 0 1 2 1 2 3 4 3 4 5 4 5
off : 0 2 5 7 9 12 14
```

Return Value

- **When test data and training data used by fitted model are python native input :**

It returns a scipy sparse csr matrix of float or double(float64) type values. It has shape (**n_queries**, **n_samples_fit**), where 'n_queries' is the number of rows in the test data and 'n_samples_fit' is the number of samples in the fitted data.

- **When either test data or training data used by fitted model is frovedis-like input:**

It returns a FrovedisCRSMatrix of float or double(float64) type values. It has shape (**n_queries**, **n_samples_fit**), where 'n_queries' is the number of rows in the test data and 'n_samples_fit' is the number of samples in the fitted data.

24.3.1.7 7. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by NearestNeighbors. It is used to get parameters and their values of NearestNeighbors class.

For example,

```
print(knn.get_params())
```

Output

```
{'algorithm': 'brute', 'batch_fraction': 1.7976931348623157e+308, 'chunk_size': 1.0,
'leaf_size': 30, 'metric': 'euclidean', 'metric_params': None, 'n_jobs': None,
'n_neighbors': 3, 'p': 2, 'radius': 2.0, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

24.3.1.8 8. set_params(params)****Parameters**

****params:** All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by NearestNeighbors, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(knn.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
knn.set_params(n_neighbors = 4, radius = 1.0)
print("get parameters after setting:")
print(knn.get_params())
```

Output

```
get parameters before setting:
{'algorithm': 'brute', 'batch_fraction': 1.7976931348623157e+308, 'chunk_size': 1.0,
'leaf_size': 30, 'metric': 'euclidean', 'metric_params': None, 'n_jobs': None,
'n_neighbors': 3, 'p': 2, 'radius': 2.0, 'verbose': 0}
get parameters after setting:
{'algorithm': 'brute', 'batch_fraction': 1.7976931348623157e+308, 'chunk_size': 1.0,
'leaf_size': 30, 'metric': 'euclidean', 'metric_params': None, 'n_jobs': None,
'n_neighbors': 4, 'p': 2, 'radius': 1.0, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

24.3.1.9 9. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

Currently this method is not supported for NearestNeighbors. It is simply kept in NearestNeighbors module to maintain uniform interface for all estimators in frovedis.

Return Value

It simply raises an AttributeError.

24.3.1.10 10. load(fname)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

Purpose

Currently this method is not supported for NearestNeighbors. It is simply kept in NearestNeighbors module to maintain uniform interface for all estimators in frovedis.

Return Value

It simply raises an AttributeError.

24.3.1.11 11. debug_print()**Purpose**

Currently this method is not supported for NearestNeighbors. It is simply kept in NearestNeighbors module to maintain uniform interface for all estimators in frovedis.

Return Value

It simply raises an `AttributeError`.

24.3.1.12 12. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
knn.release()
```

This will remove the trained model, model-id present on server, along with releasing server side memory.

Return Value

It returns nothing.

24.3.1.13 13. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

24.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **KNeighbors Classifier in Frovedis**
- **KNeighbors Regressor in Frovedis**

Chapter 25

Principal Component Analysis

25.1 NAME

PCA - It's full form is Principal Component Analysis. It is an unsupervised learning algorithm that is used for dimensionality reduction in machine learning.

25.2 SYNOPSIS

```
frovedis.mllib.decomposition.pca(n_components=None, copy=True, whiten=False,  
                                svd_solver='auto', tol=0.0, iterated_power='auto',  
                                random_state=None)
```

25.2.1 Public Member Functions

```
fit(X, y = None)  
fit_transform(X)  
get_params(deep = True)  
inverse_transform(X)  
is_fitted()  
load(path, dtype)  
load_binary(path, dtype)  
release()  
save(path)  
save_binary(path)  
set_params(**params)  
transform(X)
```

25.3 DESCRIPTION

It is one of the popular tools that is used for exploratory data analysis and predictive modeling. It is a technique to draw strong patterns from the given dataset by reducing the variances.

PCA generally tries to find the lower-dimensional surface to project the high-dimensional data.

For dimensionality reduction, it uses Singular Value Decomposition of the data. The input data is centered but not scaled for each feature before applying the SVD.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn PCA interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for PCA on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When transform-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

Like Scikit-learn, PCA in frovedis only supports dense input data. However, when sparse input is provided in frovedis, internally, it will be converted into frovedis-like dense input before sending it to frovedis server.

25.3.1 Detailed Description

25.3.1.1 1. PCA()

Parameters

n_components: It accepts a positive integer value as parameter that specifies the number of components to keep. It must be in range `[1, min(n_samples_, n_features_) - 1]`. (Default: None)

When it is None (not specified explicitly), it will be set as `min(n_samples_, n_features_) - 1`.

copy: It is a boolean parameter that specifies whether input data passed is overwritten or not during fitting. (Default: True)

When it is False (specified explicitly), data passed to `fit()` is overwritten and running `fit(X).transform(X)` will not yield the expected results, so `fit_transform(X)` is to be used instead.

whiten: It is a boolean parameter. It removes some information from the transformed signal (the relative variance scales of the components) but can sometime improve the predictive accuracy of the downstream estimators. (Default: False)

When it is True (specified explicitly), the ***components_*** vectors are multiplied by the square root of ***n_samples*** and then divided by the singular values to ensure uncorrelated outputs with unit component-wise variances.

svd_solver: It accepts a string value as parameter that specifies which solver to use. **Currently it supports only 'arpack' implementation of svd.** (Default: 'auto')

When it is auto, then ***svd_solver*** = 'arpack'.

tol: This is an unused parameter. (Default: 0.0)

iterated_power: This is an unused parameter. (Default: 'auto')

random_state: This is an unused parameter. (Default: None)

Attributes

components_: It is a numpy ndarray of shape `(n_components, n_features)`. It specifies the principal axes in feature space, representing the directions of maximum variance in the data.

explained_variance_: It is a numpy ndarray of shape `(n_components,)`. It specifies the variance of the training samples transformed by a projection to each component.

explained_variance_ratio_: It is a numpy ndarray of shape `(n_components,)`. It specifies the percentage of variance explained by each of the selected components. If 'n_components' is not set, then all components are stored and the sum of the ratios is equal to 1.0.

mean_: It is a numpy ndarray of shape `(n_features,)`. It specifies the per-feature empirical mean, estimated from the training set. It is equal to `X.mean(axis=0)`.

`n_components`: It is a positive integer value that specifies the estimated number of components.
`n_features`: It is a positive integer value that specifies the number of features in the training data (X).
`n_samples`: It is a positive integer value that specifies the number of samples in the training data (X).
`noise_varaince`: It is a float (float32) value that specifies the estimated noise covariance. It is required to compute the estimated data covariance and score samples. It is equal to the **average of (min(`n_features`, `n_samples`) - `n_components`)** smallest eigenvalues of the covariance matrix of input matrix (X).
`singular_values`: It is a numpy ndarray of shape (**`n_components`**,) that specifies the singular values corresponding to each of the selected components.

Purpose

It initializes a PCA object with the given parameters.

The parameters: “tol”, “iterated_power” and “random_state” are simply kept in to make the interface uniform to the Scikit-learn PCA module. They are not used in frovedis implementation internally.

Return Value

It simply returns “self” reference.

25.3.1.2 2. fit(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**`n_samples`**, **`n_features`**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It will fit the model with input matrix (X).

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                 dtype = np.float64)

# fitting input matrix on PCA object
from frovedis.ml.lib.decomposition import PCA
pca = PCA().fit(mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                 dtype = np.float64)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
```

```
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# PCA with pre-constructed frovedis-like inputs
from frovedis.mllib.decomposition import PCA
pca = PCA().fit(rmat)
```

Return Value

It simply returns “self” reference.

25.3.1.3 3. fit_transform(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float or double (float64) type values. It can also be an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It will fit the model with input matrix (X) and apply the dimensionality reduction on input matrix (X).

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                 dtype = np.float64)

# fitting input matrix on PCA object and perform transform
from frovedis.mllib.decomposition import PCA
pca = PCA()
print(pca.fit_transform(mat))
```

Output

```
[[ -6.50037234  0.28016556]
 [ 1.0441383  -0.75923769]
 [ 5.45623404  0.47907213]]
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                 dtype = np.float64)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
```

```

rmat = FrovedisRowmajorMatrix(mat)

# Fitting PCA with pre-constructed frovedis-like inputs and perform transform
from frovedis.mllib.decomposition import PCA
pca = PCA()
print(pca.fit_transform(rmat))

```

Output

```

[[-6.50037234  0.28016556]
 [ 1.0441383  -0.75923769]
 [ 5.45623404  0.47907213]]

```

Return Value

For both frovedis-like input and python input, it returns a numpy ndarray of shape (**n_samples**, **n_components**) and double (float64) type values. It contains transformed values.

25.3.1.4 4. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by PCA. It is used to get parameters and their values of PCA class.

For example,

```
print(pca.get_params())
```

Output

```

{'copy': True, 'iterated_power': 'auto', 'n_components': 2, 'random_state': None,
'svd_solver': 'arpack', 'tol': 0.0, 'whiten': False}

```

Return Value

A dictionary of parameter names mapped to their values.

25.3.1.5 5. `inverse_transform(X, y = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It transforms data back to its original space.

In other words, it returns an input matrix (X_original) whose transform would be input matrix (X).

For example,

```

# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],

```

```

        [4.0, 0.0, 0.0, 6.0, 7.0]],
        dtype = np.float64)

# fitting input matrix on PCA object and perform transform
from frovedis.mllib.decomposition import PCA
pca = PCA().fit(mat)
X1 = pca.transform(mat)

# inverse_transform() demo to get original input data
print(pca.inverse_transform(X1))

```

Output

```

[[ 1.00000000e+00  0.00000000e+00  7.00000000e+00  5.77315973e-15
   1.24344979e-14]
 [ 2.00000000e+00  0.00000000e+00  3.00000000e+00  4.00000000e+00
   5.00000000e+00]
 [ 4.00000000e+00  0.00000000e+00 -7.99360578e-15  6.00000000e+00
   7.00000000e+00]]

```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```

# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                 dtype = np.float64)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCSRMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# PCA with pre-constructed frovedis-like inputs and perform transform
from frovedis.mllib.decomposition import PCA
pca = PCA().fit(rmat)
X1 = pca.transform(rmat)

# inverse_transform() demo to get original frovedis-like input data
X2 = pca.inverse_transform(X1)
X2.debug_print()

```

Output

```

matrix:
num_row = 3, num_col = 5
node 0
node = 0, local_num_row = 3, local_num_col = 5, val = 1 0 7 5.77316e-15 1.24345e-14
2 0 3 4 5 4 0 -7.99361e-15 6 7

```

Return Value

- When X is python native input:

It returns a numpy ndarray of shape (**n_samples**, **n_components**) and double (float64) type values. It contains original input data values.

- **When X is frovedis-like input:**

It returns a FrovedisRowmajorMatrix instance of shape (**n_samples**, **n_components**) and double (float64) type values. It contains original input data values.

25.3.1.6 6. `is_fitted()`

Purpose

It can be used to confirm if the model is already fitted or not. In case, `transform()` is used before training the model, then it can prompt the user to train the pca model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

25.3.1.7 7. `load(path, dtype)`

Parameters

path: A string object containing the name of the file having model information to be loaded.

dtype: It is the data-type of the loaded model with trainig data samples. Currently, expected input data-type is either float (float32) or double (float64).

Purpose

It loads the model from the specified file.

For example,

```
pca.load("./out/PCAModel", dtype = np.float64)
```

Return Value

It simply returns "self" reference.

25.3.1.8 8. `load_binary(path, dtype)`

Parameters

path: A string object containing the name of the binary file having model information to be loaded.

dtype: It is the data-type of the loaded model with trainig data samples. Currently, expected input data-type is double (float64).

Purpose

It loads the model from the specified file (having binary data).

For example,

```
pca.load("./out/PCA_BinaryModel", dtype = np.float64)
```

Return Value

It simply returns "self" reference.

25.3.1.9 9. `save(path)`

Parameters

path: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (components, mean, score, etc.) in the specified file. Otherwise, it throws an exception.

For example,

```
# To save the pca model
pca.save("./out/PCAModel")
```

This will save the pca model on the path '/out/PCAModel'. It would raise exception if the directory already exists with same name.

The 'PCAModel' directory has

PCAModel

```
|—components
|—mean
|—score
|—singular_values
|—variance
|—variance_ratio
```

The saved model contains the after-fitted attribute values.

Return Value

It returns nothing.

25.3.1.10 10. save__binary(path)**Parameters**

path: A string object containing the name of the binary file on which the target model is to be saved.

Purpose

On success, it writes the model information (components, mean, score, etc.) in binary format in the specified file. Otherwise, it throws an exception.

For example,

```
# To save the pca binary model
pca.save("./out/PCA_BinaryModel")
```

This will save the pca model on the path '/out/PCA_BinaryModel'. It would raise exception if the directory already exists with same name.

The 'PCA_BinaryModel' directory has

PCA_BinaryModel

```
|—components
|—mean
|—score
|—singular_values
|—variance
|—variance_ratio
```

The saved binary model contains the after-fitted attribute values.

Return Value

It returns nothing.

25.3.1.11 11. set_params(params)****Parameters**

****params:** All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by PCA, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(pca.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
pca.set_params(whiten=True)
print("get parameters after setting:")
print(pca.get_params())
```

Output

```
get parameters before setting:
{'copy': True, 'iterated_power': 'auto', 'n_components': 2, 'random_state': None,
'svd_solver': 'arpack', 'tol': 0.0, 'whiten': False}
get parameters after setting:
{'copy': True, 'iterated_power': 'auto', 'n_components': 2, 'random_state': None,
'svd_solver': 'arpack', 'tol': 0.0, 'whiten': True}
```

Return Value

It simply returns “self” reference.

25.3.1.12 12. transform(X, y = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It applies the dimensionality reduction on input matrix (X).

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                dtype = np.float64)

# fitting input matrix on PCA object and perform transform
from frovedis.ml.lib.decomposition import PCA
pca = PCA().fit(mat)
print(pca.transform(mat))
```

Output

```

[[-6.50037234  0.28016556]
 [ 1.0441383  -0.75923769]
 [ 5.45623404  0.47907213]]

```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```

# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[1.0, 0.0, 7.0, 0.0, 0.0],
                 [2.0, 0.0, 3.0, 4.0, 5.0],
                 [4.0, 0.0, 0.0, 6.0, 7.0]],
                 dtype = np.float64)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# Fitting PCA with pre-constructed frovedis-like inputs and perform transform
from frovedis.ml.lib.decomposition import PCA
pca = PCA().fit(rmat)
X1 = pca.transform(rmat)
X1.debug_print()

```

Output

```

matrix:
num_row = 3, num_col = 2
node 0
node = 0, local_num_row = 3, local_num_col = 2, val = -6.50037 0.280166 1.04414
-0.759238 5.45623 0.479072

```

Return Value

- **When X is python native input:**
It returns a numpy ndarray of shape (**n_samples**, **n_components**) and double (float64) type values. It contains the projection of input matrix (X) in the first principal components.
- **When X is frovedis-like input:**
It returns a FrovedisRowmajorMatrix instance of shape (**n_samples**, **n_components**) and double (float64) type values. It contains the projection of input matrix (X) in the first principal components.

25.4 SEE ALSO

- [Introduction to FrovedisRowmajorMatrix](#)
- [Introduction to FrovedisCRSMMatrix](#)
- [Latent Dirichlet Allocation in frovedis](#)

Chapter 26

RandomForestClassifier

26.1 NAME

RandomForestClassifier - A classification algorithm that contains multiple decision trees on various subsets of the given dataset. It contains the predictions of multiple decision trees and based on the majority votes of predictions, it predicts the final output.

26.2 SYNOPSIS

```
class frovedis.mllib.ensemble.forest.RandomForestClassifier(n_estimators=100, criterion='gini',
                                                            max_depth=None,
                                                            min_samples_split=2,
                                                            min_samples_leaf=1,
                                                            min_weight_fraction_leaf=0.0,
                                                            max_features='auto',
                                                            max_leaf_nodes=None,
                                                            min_impurity_decrease=0.0,
                                                            min_impurity_split=None,
                                                            bootstrap=True, oob_score=False,
                                                            n_jobs=None, random_state=None,
                                                            verbose=0, warm_start=False,
                                                            class_weight=None, ccp_alpha=0.0,
                                                            max_samples=None, max_bins=32)
```

26.2.1 Public Member Functions

```
fit(X, y)
predict(X)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
score(X, y, sample_weight = None)
save(fname)
debug_print()
```

```
release()
is_fitted()
```

26.3 DESCRIPTION

Random Forest Classifier is a supervised machine learning algorithm used for classification using decision trees. The classifier creates a set of decision trees from randomly selected subsets of the training set. It is basically a set of decision trees from a randomly selected subset of the training set and then it collects the votes from different decision trees to decide the final prediction. **Frovedis supports both binary and multinomial random forest classification algorithms.**

During training, the input **X** is the training data and **y** is the corresponding label values (Frovedis supports any values for labels, but internally it encodes the input binary labels to 0 and 1, and input multinomial labels to 0, 1, 2, ..., N-1 (where N is the no. of classes) before training at Frovedis server) which we want to predict.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn RandomForestClassifier interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for RandomForestClassifier on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

26.3.1 Detailed Description

26.3.1.1 1. RandomForestClassifier()

Parameters

n_estimators: A positive integer parameter that specifies the number of trees in the forest. (Default: 100)
criterion: A string object parameter that specifies the function to measure the quality of a split. Supported criteria are 'gini' and 'entropy'. (Default: 'gini')

- **'gini' impurity**: calculates the amount of probability of a specific feature that is classified incorrectly when selected randomly.

- **'entropy' (information gain)**: it is applied to quantify which feature provides maximal information about the classification based on the notion of entropy.

max_depth: A positive integer parameter that specifies the maximum depth of the tree. (Default: None)
 If it is None (not specified explicitly), then 'max_depth' is set to 4.

min_samples_split: An integer or float value that specifies the minimum number of samples required to split an internal node. (Default: 2)

min_samples_leaf: A positive integer or float value that specifies the minimum number of samples required to be at a leaf node. A split point at any depth will only be considered if it leaves at least 'min_samples_leaf' training samples in each of the left and right branches. (Default: 1)

- If it is an integer, then 'min_samples_leaf' should be greater than 0.

- If it is float, then it is set as `int(np.ceil(self.min_samples_split * self.n_samples_))`

min_weight_fraction_leaf: An unused parameter. (Default: 0.0)

max_features: A string object parameter that specifies the number of features to consider when looking for the best split:

- If it is an integer, then it will be set as $(\text{max_features} * 1.0) / \text{n_features_}$.
- If it is float, then it will be set as ‘max_features’ number of features at each split.
- If it is ‘auto’, then it will be set as $\text{sqrt}(\text{n_features_})$.
- If ‘sqrt’, then it will be set as $\text{sqrt}(\text{n_features_})$ (same as ‘auto’).
- If ‘log2’, then it will be set as $\text{log2}(\text{n_features_})$.
- If None, then it will be set as n_features_ . (Default: ‘auto’)

max_leaf_nodes: An unused parameter. (Default: None)

min_impurity_decrease: A positive double (float64) parameter. A node will be split if this split induces a decrease of the impurity greater than or equal to this value. (Default: 0.0)

min_impurity_split: An unused parameter. (Default: None)

bootstrap: An unused parameter. (Default: True)

oob_score: An unused parameter. (Default: False)

n_jobs: An unused parameter. (Default: None)

random_state: An integer parameter that controls the sampling of the features to consider when looking for the best split at each node (if $\text{max_features} < \text{n_features}$). (Default: None)

If it is None (not specified explicitly), then ‘random_state’ is set as -1.

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

warm_start: An unused parameter. (Default: False)

class_weight: An unused parameter. (Default: None)

ccp_alpha: An unused parameter. (Default: 0.0)

max_samples: An unused parameter. (Default: None)

max_bins: A positive integer parameter that specifies the maximum number of bins created by ordered splits. (Default: 32)

Attributes

classes_: It is a python ndarray (any type) of unique labels given to the classifier during training. It has shape $(\text{n_classes},)$.

n_features_: An integer value specifying the number of features when fitting the estimator.

Purpose

It initializes a RandomForestClassifier object with the given parameters.

The parameters: “min_weight_fraction_leaf”, “max_leaf_nodes”, “min_impurity_split”, “bootstrap”, “oob_score”, “n_jobs”, “warm_start”, “class_weight”, “ccp_alpha” and “max_samples” are simply kept in to make the interface uniform to the Scikit-learn RandomForestClassifier module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

26.3.1.2 2. fit(X, y)

Parameters

X: A numpy dense or any python array-like object or an instance of FrovedisColmajorMatrix for dense data. It has shape $(\text{n_samples}, \text{n_features})$. Currently, it supports only dense data as input.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape $(\text{n_samples},)$.

Purpose

It builds a forest of trees from the training data X and labels y.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]], dtype = np.float64)
lbl = np.array([100, 500, 100, 600], dtype = np.float64)

# fitting input matrix and label on RandomForestClassifier object
from frovedis.mllib.ensemble import RandomForestClassifier
rfc = RandomForestClassifier(n_estimators = 10, max_depth = 4, min_samples_split = 0.5
                             min_samples_leaf = 1.2, random_state = 324)

rfc.fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]], dtype = np.float64)
lbl = np.array([100, 500, 100, 600], dtype = np.float64)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# fitting input matrix and label on RandomForestClassifier object
from frovedis.mllib.ensemble import RandomForestClassifier
rfc = RandomForestClassifier(n_estimators = 10, max_depth = 4,
                             min_samples_split = 0.5, min_samples_leaf = 1.2,
                             random_state = 324)

rfc.fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

26.3.1.3 3. predict(X)

Parameters

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

Purpose

Predict class for X.

The predicted class of an input sample is a vote by the trees in the forest, weighted by their probability estimates. That is, the predicted class is the one with highest mean probability estimate across the trees.

For example,

```
# predicting on random forest classifier model
rfc.predict(mat)
```

Output

```
[100. 500. 100. 500.]
```

Like in `fit()`, `frovedis`-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on random forest classifier model
rfc.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[100. 500. 100. 500.]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape `(n_samples,)` containing the predicted classes.

26.3.1.4 4. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the `BaseEstimator` class inherited by `RandomForestClassifier`. It is used to get parameters and their values of `RandomForestClassifier` class.

For example,

```
print(rfc.get_params())
```

Output

```
{'bootstrap': True, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'gini',
 'max_bins': 32, 'max_depth': 4, 'max_features': 'auto', 'max_leaf_nodes': None,
 'max_samples': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
 'min_samples_leaf': 2, 'min_samples_split': 0.5, 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 10, 'n_jobs': None, 'oob_score': False, 'random_state': 324,
 'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

26.3.1.5 5. `set_params(**params)`

Parameters

*****params***: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by RandomForestClassifier, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(rfc.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
rfc.set_params(criterion = 'entropy', max_depth = 5)
print("get parameters after setting:")
print(rfc.get_params())
```

Output

```
get parameters before setting:
{'bootstrap': True, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'gini',
 'max_bins': 32, 'max_depth': 4, 'max_features': 'auto', 'max_leaf_nodes': None,
 'max_samples': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
 'min_samples_leaf': 2, 'min_samples_split': 0.5, 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 10, 'n_jobs': None, 'oob_score': False, 'random_state': 324,
 'verbose': 0, 'warm_start': False}
get parameters after setting:
{'bootstrap': True, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'entropy',
 'max_bins': 32, 'max_depth': 5, 'max_features': 'auto', 'max_leaf_nodes': None,
 'max_samples': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
 'min_samples_leaf': 2, 'min_samples_split': 0.5, 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 10, 'n_jobs': None, 'oob_score': False, 'random_state': 324,
 'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

26.3.1.6 6. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
rfc.load("./out/rf_classifier_model")
```

Return Value

It simply returns “self” reference.

26.3.1.7 7. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the random forest classifier model
dtc.save("./out/rf_classifier_model")
```

This will save the random forest classifier model on the path “/out/rf_classifier_model”. It would raise exception if the directory already exists with same name.

The ‘rf_classifier_model’ directory has

rf_classifier_model

```
|—label_map
|—metadata
|—model
```

‘label_map’ file contains information about labels mapped with their encoded value.

The ‘metadata’ file contains the number of classes, model kind and input datatype used for trained model.

The ‘model’ file contains the random forest model saved in binary format.

Return Value

It returns nothing.

26.3.1.8 8. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object containing the true labels for X. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of self.predict(X) wrt. y.

For example,

```
# calculate mean accuracy score on given test data and labels
rfc.score(mat,lbl)
```

Output

```
0.75
```

Return Value

It returns an accuracy score of float type.

26.3.1.9 9. debug_print()**Purpose**

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
rfc.debug_print()
```

Output

```
----- Classification Trees (Random Forest):: -----
# of trees: 10
---- [0] ----
  # of nodes: 3, height: 1
  <1> Split: feature[2] < 0.25, IG: 0.5
    \_ (2) Predict: 2 (100%)
    \_ (3) Predict: 0 (100%)
---- [1] ----
  # of nodes: 3, height: 1
  <1> Split: feature[6] < 0.25, IG: 0.375
    \_ (2) Predict: 1 (50%)
    \_ (3) Predict: 0 (100%)
---- [2] ----
  # of nodes: 1, height: 0
  (1) Predict: 0 (50%)
---- [3] ----
  # of nodes: 3, height: 1
  <1> Split: feature[6] < 0.25, IG: 0.125
    \_ (2) Predict: 0 (50%)
    \_ (3) Predict: 0 (100%)
---- [4] ----
  # of nodes: 3, height: 1
  <1> Split: feature[3] < 0.25, IG: 0.375
    \_ (2) Predict: 0 (100%)
    \_ (3) Predict: 1 (50%)
---- [5] ----
  # of nodes: 3, height: 1
  <1> Split: feature[1] < 0.25, IG: 0.5
    \_ (2) Predict: 2 (100%)
    \_ (3) Predict: 1 (100%)
---- [6] ----
  # of nodes: 1, height: 0
  (1) Predict: 0 (100%)
---- [7] ----
  # of nodes: 1, height: 0
  (1) Predict: 2 (75%)
---- [8] ----
  # of nodes: 1, height: 0
  (1) Predict: 1 (50%)
---- [9] ----
  # of nodes: 3, height: 1
  <1> Split: feature[5] < 0.25, IG: 0.375
    \_ (2) Predict: 0 (100%)
    \_ (3) Predict: 1 (50%)
```

This output will be visible on server side. It displays the random forest having maximum depth of 4 and total 10 decision trees.

No such output will be visible on client side.

Return Value

It returns nothing.

26.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
rfc.release()
```

This will reset the after-fit populated attributes (like `classes__`, `n_features__`) to `None`, along with releasing server side memory.

Return Value

It returns nothing.

26.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted, otherwise, it returns 'False'.

26.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisDvector**
- **Random Forest Regressor in Frovedis**
- **Decision Tree Classifier in Frovedis**

Chapter 27

RandomForestRegressor

27.1 NAME

RandomForestRegressor - A regression algorithm used for predicting the final output by taking the mean of all the predictions from multiple decision trees.

27.2 SYNOPSIS

```
class frovedis.mllib.ensemble.forest.RandomForestRegressor(n_estimators=100, criterion='mse',
                                                            max_depth=None,
                                                            min_samples_split=2,
                                                            min_samples_leaf=1,
                                                            min_weight_fraction_leaf=0.0,
                                                            max_features='auto',
                                                            max_leaf_nodes=None,
                                                            min_impurity_decrease=0.0,
                                                            min_impurity_split=None,
                                                            bootstrap=True, oob_score=False,
                                                            n_jobs=None, random_state=None,
                                                            verbose=0, warm_start=False,
                                                            ccp_alpha=0.0, max_samples=None,
                                                            max_bins=32)
```

27.2.1 Public Member Functions

```
fit(X, y)
predict(X)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
score(X, y, sample_weight = None)
debug_print()
release()
is_fitted()
```

27.3 DESCRIPTION

Random Forest Regressor is a supervised machine learning algorithm used for regression using decision trees. Every decision tree has high variance, but when we combine all of them together in parallel then the resultant variance is low as each decision tree gets perfectly trained on that particular sample data and hence the output doesn't depend on one decision tree but on multiple decision trees. The regressor creates a set of decision trees from randomly selected subsets of the training set. Then, it takes the average to improve the predictive accuracy of that dataset. So, instead of relying on one decision tree, it combines multiple decision trees in determining the final output.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn RandomForestRegressor interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for RandomForestRegressor on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

27.3.1 Detailed Description

27.3.1.1 1. RandomForestRegressor()

Parameters

n_estimators: A positive integer parameter that specifies the number of trees in the forest. (Default: 100)
criterion: A string object parameter that specifies the function to measure the quality of a split. (Default: 'mse')

Currently, supported criteria are 'mse' and 'mae'. The mean squared error (mse), which is equal to variance reduction as feature selection criterion. The mean absolute error (mae) uses reduction in Poisson deviance to find splits.

max_depth: A positive integer parameter that specifies the maximum depth of the tree. (Default: None) If it is None (not specified explicitly), then 'max_depth' is set to 4.

min_samples_split: An integer or float value that specifies the minimum number of samples required to split an internal node. (Default: 2)

min_samples_leaf: A positive integer or float value that specifies the minimum number of samples required to be at a leaf node. A split point at any depth will only be considered if it leaves at least 'min_samples_leaf' training samples in each of the left and right branches. (Default: 1)

- If it is an integer, then 'min_samples_leaf' should be greater than 0.

- If it is float, then it is set as `int(np.ceil(self.min_samples_split * self.n_samples_))`

min_weight_fraction_leaf: An unused parameter. (Default: 0.0)

max_features: A string object parameter that specifies the number of features to consider when looking for the best split:

- If it is an integer, then it will be set as `(max_features * 1.0) / n_features_`.

- If it is float, then it will be 'max_features' number of features at each split.

- If it is 'auto', then it will be set as `sqrt(n_features_)`.

- If 'sqrt', then it will be set as `sqrt(n_features_)` (same as 'auto').

- If 'log2', then it will be set as `log2(n_features_)`.

- If None, then it will be set as `n_features_`. (Default: 'auto')

max_leaf_nodes: An unused parameter. (Default: None)

min_impurity_decrease: A positive double (float64) parameter. A node will be split if this split induces a decrease of the impurity greater than or equal to this value. (Default: 0.0)

min_impurity_split: An unused parameter. (Default: None)

bootstrap: An unused parameter. (Default: True)

oob_score: An unused parameter. (Default: False)

n_jobs: An unused parameter. (Default: None)

random_state: An integer parameter that controls the sampling of the features to consider when looking for the best split at each node (if max_features < n_features). (Default: None)

If it is None (not specified explicitly), then 'random_state' is set as -1.

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

warm_start: An unused parameter. (Default: False)

ccp_alpha: An unused parameter. (Default: 0.0)

max_samples: An unused parameter. (Default: None)

max_bins: A positive integer parameter that specifies the maximum number of bins created by ordered splits. (Default: 32)

Attributes

n_features_: An integer value specifying the number of features when fitting the estimator.

Purpose

It initializes a RandomForestRegressor object with the given parameters.

The parameters: "min_weight_fraction_leaf", "max_leaf_nodes", "min_impurity_split", "bootstrap", "oob_score", "n_jobs", "warm_start", "ccp_alpha" and "max_samples" are simply kept in to make the interface uniform to the Scikit-learn RandomForestRegressor module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns "self" reference.

27.3.1.2 2. fit(X, y)

Parameters

X: A numpy dense or any python array-like object or an instance of FrovedisColmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**,).

Purpose

It builds a forest of trees from the training data X and labels y.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]], dtype = np.float64)
lbl = np.array([1.2, 0.3, 1.1, 1.9])

# fitting input matrix and label on RandomForestRegressor object
from frovedis.ml.lib.ensemble import RandomForestRegressor
rfr = RandomForestRegressor(n_estimators = 10, criterion = 'mae', max_depth = 5)
```

```
rfr.fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels dense data
import numpy as np
mat = np.array([[10, 0, 1, 0, 0, 1, 0],
                [0, 1, 0, 1, 0, 1, 0],
                [0, 1, 0, 0, 1, 0, 1],
                [1, 0, 0, 1, 0, 1, 0]], dtype = np.float64)
lbl = np.array([1.2, 0.3, 1.1, 1.9])

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# fitting input matrix and label on RandomForestRegressor object
from frovedis.mllib.ensemble import RandomForestRegressor
rfr = RandomForestRegressor(n_estimators = 10, criterion = 'mae', max_depth = 5)
rfr.fit(cmat, dlbl)
```

Return Value

It simply returns “self” reference.

27.3.1.3 3. predict(X)

Parameters

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

Purpose

Predict regression value for X.

The predicted regression target of an input sample is computed as the mean predicted regression targets of the trees in the forest.

For example,

```
# predicting on random forest regressor model
rfr.predict(mat)
```

Output

```
[1.29083333 0.8075      1.0475      1.48083333]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on random forest regressor model using pre-constructed input
rfr.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[1.29083333 0.8075      1.0475      1.48083333]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**,) containing the predicted values.

27.3.1.4 4. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by RandomForestRegressor. It is used to get parameters and their values of RandomForestRegressor class.

For example,

```
print(rfr.get_params())
```

Output

```
{'bootstrap': True, 'ccp_alpha': 0.0, 'criterion': 'mae', 'max_bins': 32,
 'max_depth': 5, 'max_features': 'auto', 'max_leaf_nodes': None,
 'max_samples': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
 'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 10, 'n_jobs': None, 'oob_score': False, 'random_state': -1,
 'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

27.3.1.5 5. `set_params(**params)`

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by RandomForestRegressor, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(rfr.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
rfr.set_params(max_depth = 4)
print("get parameters after setting:")
print(rfr.get_params())
```

Output

```
get parameters before setting:
{'bootstrap': True, 'ccp_alpha': 0.0, 'criterion': 'mae', 'max_bins': 32,
 'max_depth': 5, 'max_features': 'auto', 'max_leaf_nodes': None,
```

```
'max_samples': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'n_estimators': 10, 'n_jobs': None, 'oob_score': False, 'random_state': -1,
'verbose': 0, 'warm_start': False}
get parameters after setting:
{'bootstrap': True, 'ccp_alpha': 0.0, 'criterion': 'mae', 'max_bins': 32,
'max_depth': 4, 'max_features': 'auto', 'max_leaf_nodes': None,
'max_samples': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None,
'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0,
'n_estimators': 10, 'n_jobs': None, 'oob_score': False, 'random_state': -1,
'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

27.3.1.6 6. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

For example,

```
rfr.load("./out/rf_regressor_model")
```

Return Value

It simply returns “self” reference.

27.3.1.7 7. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the random forest regressor model
rfr.save("./out/rf_regressor_model")
```

This will save the random forest regressor model on the path “/out/rf_regressor_model”. It would raise exception if the directory already exists with same name.

The ‘rf_regressor_model’ directory has

```
rf_regressor_model
|—metadata
|—model
```

The ‘metadata’ file contains the model kind and input datatype used for trained model.

The ‘model’ file contains the random forest model saved in binary format.

Return Value

It returns nothing.

27.3.1.8 8. score(X, y, sample_weight = None)**Parameters**

X: A numpy dense or any python array-like object or an instance of FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**). Currently, it supports only dense data as input.

y: Any python array-like object containing the true values for X. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of self.predict(X) wrt. y.

The coefficient 'R2' is defined as $(1 - (u/v))$,

where 'u' is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and,

'v' is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
# calculate R2 score on given test data and labels
rfr.score(mat, lbl)
```

Output

0.65

Return Value

It returns an R2 score of float type.

27.3.1.9 9. debug_print()**Purpose**

It shows the target model information on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
rfr.debug_print()
```

Output

```
----- Regression Trees (Random Forest):: -----
# of trees: 10
---- [0] ----
  # of nodes: 3, height: 1
  <1> Split: feature[1] < 0.25, IG: 0.191667
    \_ (2) Predict: 1.43333
    \_ (3) Predict: 0.3
---- [1] ----
  # of nodes: 5, height: 2
  <1> Split: feature[2] < 0.25, IG: 0.0416667
    \_ <2> Split: feature[1] < 0.5, IG: 0.711111
      | \_ (4) Predict: 1.9
```

```

    | \_ (5) Predict: 0.3
    \_ (3) Predict: 1.2
---- [2] ----
    # of nodes: 1, height: 0
    (1) Predict: 1.375
---- [3] ----
    # of nodes: 5, height: 2
    <1> Split: feature[1] < 0.25, IG: 0.4
    \_ (2) Predict: 1.9
    \_ <3> Split: feature[3] < 0.5, IG: 0.4
        \_ (6) Predict: 1.1
        \_ (7) Predict: 0.3
---- [4] ----
    # of nodes: 3, height: 1
    <1> Split: feature[1] < 0.25, IG: 0.4
    \_ (2) Predict: 1.9
    \_ (3) Predict: 1.1
---- [5] ----
    # of nodes: 3, height: 1
    <1> Split: feature[4] < 0.25, IG: 0.4
    \_ (2) Predict: 0.3
    \_ (3) Predict: 1.1
---- [6] ----
    # of nodes: 5, height: 2
    <1> Split: feature[5] < 0.25, IG: 0.141667
    \_ (2) Predict: 1.1
    \_ <3> Split: feature[2] < 0.5, IG: 0.311111
        \_ (6) Predict: 1.9
        \_ (7) Predict: 1.2
---- [7] ----
    # of nodes: 7, height: 3
    <1> Split: feature[0] < 2.5, IG: 0.025
    \_ <2> Split: feature[6] < 0.5, IG: 1.11022e-16
        | \_ <4> Split: feature[1] < 0.5, IG: 0.8
            | \_ (8) Predict: 1.9
            | \_ (9) Predict: 0.3
        | \_ (5) Predict: 1.1
        \_ (3) Predict: 1.2
---- [8] ----
    # of nodes: 3, height: 1
    <1> Split: feature[2] < 0.25, IG: 0.2625
    \_ (2) Predict: 1.9
    \_ (3) Predict: 1.2
---- [9] ----
    # of nodes: 5, height: 2
    <1> Split: feature[3] < 0.25, IG: 0.279167
    \_ <2> Split: feature[2] < 0.5, IG: 0.0444444
        | \_ (4) Predict: 1.1
        | \_ (5) Predict: 1.2
        \_ (3) Predict: 0.3

```

This output will be visible on server side. It displays the random forest having maximum depth of 5 and total 10 decision trees.

No such output will be visible on client side.

Return Value

It returns nothing.

27.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
rfr.release()
```

This will reset the after-fit populated attributes (like `n_features__`) to `None`, along with releasing server side memory.

Return Value

It returns nothing.

27.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns ‘True’, if the model is already fitted, otherwise, it returns ‘False’.

27.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisDvector**
- **Random Forest Classifier in Frovedis**
- **Decision Tree Regressor in Frovedis**

Chapter 28

Ridge Regression

28.1 NAME

Ridge Regression - A regression algorithm used to predict the continuous output with L2 regularization.

28.2 SYNOPSIS

```
class frovedis.mllib.linear_model.Ridge(alpha=0.01, fit_intercept=True,
                                         normalize=False, copy_X=True,
                                         max_iter=None, tol=1e-3,
                                         solver='auto', random_state=None,
                                         lr_rate=1e-8, verbose=0,
                                         warm_start=False)
```

28.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

28.3 DESCRIPTION

Linear least squares is the most common formulation for regression problems. It is a linear method with the loss function given by the **squared loss**:

$$L(w; x, y) := 1/2(wTx - y)^2$$

During training, the input \mathbf{X} is the training data and \mathbf{y} is the corresponding label values which we want to predict. \mathbf{w} is the linear model (also known as weight) which uses a single weighted sum of features to make a prediction. The method is called linear since it can be expressed as a function of $\mathbf{w}^T\mathbf{x}$ and y . Ridge regression uses L2 regularization to address the overfit problem.

The gradient of the squared loss is: $(\mathbf{w}^T\mathbf{x}-\mathbf{y})\cdot\mathbf{x}$

The gradient of the regularizer is: \mathbf{w}

Frovedis provides implementation of ridge regression with two different optimizers:

- (1) stochastic gradient descent with minibatch
- (2) LBFGS optimizer

The simplest method to solve optimization problems of the form $\min \mathbf{f}(\mathbf{w})$ is gradient descent. Such first-order optimization methods well-suited for large-scale and distributed computation. Whereas, L-BFGS is an optimization algorithm in the family of quasi-Newton methods to solve the optimization problems of the similar form.

Like the original BFGS, L-BFGS (Limited Memory BFGS) uses an estimation to the inverse Hessian matrix to steer its search through feature space, but where BFGS stores a dense $n \times n$ approximation to the inverse Hessian (n being the number of features in the problem), L-BFGS stores only a few vectors that represent the approximation implicitly. L-BFGS often achieves rapider convergence compared with other first-order optimization.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn Ridge interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Ridge on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

28.3.1 Detailed Description

28.3.1.1 1. Ridge()

Parameters

alpha: A positive value of double (float64) type is called the regularization strength parameter. (Default: 0.01)

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the decision function. (Default: True)

normalize: An unused parameter (Default: False)

copy_X: An unused parameter (Default: True)

max_iter: An integer parameter specifying the maximum iteration count. (Default: None) When it is None(not specified explicitly), it will be set as 1000.

tol: Zero or a positive value of double (float64) type specifying the convergence tolerance value. (Default: 1e-3)

solver: A string object specifying the solver to use. It can be "sag" for frovedis side stochastic gradient descent or "lbfgs" for frovedis side LBFGS optimizer when optimizing the ridge regression model. Initially solver is "auto" by default. In such cases, it will select "sag" solver. Both "sag" and "lbfgs" can handle L2 penalty.

random_state: An unused parameter. (Default: None)

lr_rate: Zero or a positive value of double (float64) type containing the learning rate. (Default: 1e-8)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

Attributes

coef_: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)) of estimated coefficients for the ridge regression problem. It has shape (**n_features**,).

intercept_(**bias**): It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)). If fit_intercept is set to False, the intercept is set to zero. It has shape (**1**,).

n_iter_: A positive integer value used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a Ridge object with the given parameters.

The parameters: “normalize”, “copy_X” and “random_state” are simply kept in to to make the interface uniform to the Scikit-learn Ridge module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

28.3.1.2 2. fit(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisColmajorMatrix for dense data.

y: Any python array-like object or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**,).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a ridge regression model with L2 regularization with those data at frovedis server.

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

Return Value

It simply returns “self” reference.

28.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix as for dense data.

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

In case pre-constructed frovedis-like training data such as FrovedisColmajorMatrix (X) is provided during prediction, then “X.to_frovedis_rowmatrix()” will be used for prediction.

Return Value

It returns a numpy array of double (float64) type and has shape (**n_samples**,) containing the predicted outputs.

28.3.1.4 4. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data.

y: Any python array-like object containing the target values for X. It has shape (**n_samples**,).

sample_weight: Python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate the root mean square value on the given test data and labels i.e. R^2 (r-squared) of self.predict(X) wrt. y.

The coefficient ‘ R^2 ’ is defined as $(1 - (u/v))$,

where ‘u’ is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and

‘v’ is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R^2 score of 0.0.

Return Value

It returns an R^2 score of float type.

28.3.1.5 5. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by Ridge. It is used to get parameters and their values of Ridge class.

Return Value

A dictionary of parameter names mapped to their values.

28.3.1.6 6. set_params(**params)

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by Ridge, used to set parameter values.

Return Value

It simply returns “self” reference.

28.3.1.7 7. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file (having little-endian binary data).

Return Value

It simply returns “self” reference.

28.3.1.8 8. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

Suppose ‘RidgeModel’ directory is the model created, It will have

RidgeModel

```
|—— metadata
|—— model
```

The metadata file contains the number of classes, model kind, input datatype used for trained model. Here, the model file contains information about weights, intercept and threshold.

It would raise exception if the ‘RidgeModel’ directory already existed with same name.

Return Value

It returns nothing.

28.3.1.9 9. debug_print()**Purpose**

It shows the target model information like weight values, intercept on the server side user terminal. It is mainly used for debugging purpose.

No such output will be visible on client side.

Return Value

It returns nothing.

28.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server. With this, after-fit populated attributes are reset to None, along with releasing server side memory.

Return Value

It returns nothing.

28.3.1.11 11. `is_fitted()`**Purpose**

It can be used to confirm if the model is already fitted or not. In case, `predict()` is used before training the model, then it can prompt the user to train the ridge regression model first.

Return Value

It returns ‘True’, if the model is already fitted otherwise, it returns ‘False’.

28.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **Linear Regression in Frovedis**
- **Lasso Regression in Frovedis**

Chapter 29

SGDClassifier

29.1 NAME

SGDClassifier - A classification algorithm used to predict the labels with various loss functions. This estimator implements regularized linear models with stochastic gradient descent (SGD) learning.

29.2 SYNOPSIS

```
class frovedis.mllib.linear_model.SGDClassifier(loss='hinge', penalty='l2', alpha=0.0001,
                                              l1_ratio=0.15, fit_intercept=True,
                                              max_iter=1000, tol=1e-3,
                                              shuffle=True, verbose=0,
                                              epsilon=0.1, n_jobs=None,
                                              random_state=None,
                                              learning_rate="invscaling",
                                              eta0=1.0, power_t=0.5,
                                              early_stopping=False,
                                              validation_fraction=0.1,
                                              n_iter_no_change=5,
                                              class_weight=None,
                                              warm_start=False,
                                              average=False)
```

29.2.1 Public Member Functions

```
fit(X, y, coef_init = None, intercept_init = None, sample_weight = None)
predict(X)
predict_proba(X)
score(X, y, sample_weight = None)
load(fname, dtype = None)
save(fname)
get_params(deep = True)
set_params(**params)
debug_print()
```

```
release()
is_fitted()
```

29.3 DESCRIPTION

Stochastic Gradient Descent (SGD) is used for discriminative learning of linear classifiers under convex loss functions such as SVM and Logistic regression. It has been successfully applied to large-scale datasets because the update to the coefficients is performed for each training instance, rather than at the end of instances. **Frovedis supports both binary and multinomial Stochastic Gradient Descent (SGD) classifier algorithms.**

Stochastic Gradient Descent (SGD) classifier basically implements a plain SGD learning routine supporting various loss functions and penalties for classification. It implements regularized linear models with stochastic gradient descent (SGD) learning: the gradient of the loss is estimated each sample at a time and the model is updated along the way with a decreasing strength schedule. It is a linear method which uses the following loss functions:

- 1) **hinge**
- 2) **log**
- 3) **squared_loss**

It supports ZERO, L1 and L2 regularization to address the overfit problem.

During training, the input **X** is the training data and **y** are their corresponding label values (Frovedis supports any values as for labels, but internally it encodes the input binary labels to -1 and 1, before training at Frovedis server) which we want to predict.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn SGDClassifier (Stochastic Gradient Descent Classification) interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for SGDClassifier on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

29.3.1 Detailed Description

29.3.1.1 1. SGDClassifier()

Parameters

loss: A string object parameter containing the loss function type to use. Currently, frovedis supports 'hinge', 'log' and 'squared_loss' functions. (Default: 'hinge')

penalty: A string object parameter containing the regularizer type to use. Currently none, l1 and l2 are supported by Frovedis. (Default: 'l2')

If it is None (not specified explicitly), it will be set as 'ZERO' regularization type.

alpha: Zero or a positive double (float64) smoothing parameter. (Default: 0.0001)

l1_ratio: An unused parameter. (Default: 0.15)

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the

decision function. (Default: True)

max_iter: A positive integer parameter specifying maximum iteration count. (Default: 1000)

tol: A double (float64) parameter specifying the convergence tolerance value. It must be zero or a positive value. (Default: 1e-3)

shuffle: An unused parameter. (Default: True)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

epsilon: An unused parameter. (Default: 0.1)

n_jobs: An unused parameter. (Default: None)

random_state: An unused parameter. (Default: None)

learning_rate: A string object parameter containing the learning rate. (Default: 'invscaling')

Unlike sklearn, Frovedis only supports 'invscaling' learning rate. 'invscaling' gradually decreases the learning rate '**learning_rate**' at each time step '**t**' using an inverse scaling exponent of '**power_t**'.

learning_rate = $\text{eta0} / \text{pow}(\text{t}, \text{power_t})$

eta0: A double (float64) parameter specifying the initial learning rate for the 'invscaling' schedules. (Default: 1.0)

power_t: A double (float64) parameter specifying the exponent for inverse scaling learning rate. (Default: 0.5)

early_stopping: An unused parameter. (Default: False)

validation_fraction: An unused parameter. (Default: 0.1)

n_iter_no_change: An unused parameter. (Default: 5)

class_weight: An unused parameter. (Default: None)

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

average: An unused parameter. (Default: False)

Attributes

coef: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)). It is the weights assigned to the features.

Shape of this attribute depends on the **n_classes**.

- If '**classes**' is 2, then the shape is (**1**, **n_features**)

- If '**classes**' is more then 2, then the shape is (**n_classes**, **n_features**).

intercept: It is a python ndarray(float or double (float64) values depending on input matrix data type) and has shape (**1**,). It specifies the constants in decision function.

classes: It is a python ndarray(any type) of unique labels given to the classifier during training. It has shape (**n_classes**,). This attribute is not available for **squared_loss**.

n_iter: An integer value used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a SGDClassifier object with the given parameters.

The parameters: "l1_ratio", "shuffle", "epsilon", "n_jobs", "random_state", "early_stopping", "validation_fraction", "n_iter_no_change", "class_weight" and "average" are simply kept to make the interface uniform to Scikit-learn SGDClassifier module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns "self" reference.

29.3.1.2 2. fit(X, y, coef_init = None, intercept_init = None, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisColmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape

(**n_samples**,).

coef_init: An unused parameter that specifies the initial coefficients to warm-start the optimization. (Default: None)

intercept_init: An unused parameter that specifies the initial intercept to warm-start the optimization. (Default: None)

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a SGDClassifier model with specified regularization with those data at frovedis server.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# fitting input matrix and label on SGDClassifier object
from frovedis.mllib.linear_model import SGDClassifier
sgd_clf = SGDClassifier().fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_breast_cancer
mat, lbl = load_breast_cancer(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# and for scipy sparse data, FrovedisCRSMMatrix should be used.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# Linear SVC with pre-constructed frovedis-like inputs
from frovedis.mllib.linear_model import SGDClassifier
sgd_clf = SGDClassifier().fit(cmat,dlbl)
```

Return Value

It simply returns “self” reference.

29.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
sgd_clf.predict(mat)
```

Output:

```
[0 0 0 1 . . . . 0 0 0 1]
```

Like in `fit()`, frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on SGDClassifier using frovedis-like input
sgd_clf.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[0 0 0 1 . . . . 0 0 0 1]
```

Return Value

For **squared_loss** loss, it returns a numpy array of **double (float64)** type and for other loss functions it returns a numpy array of **int64** type containing the predicted outputs. It has shape **(n_samples,)**.

29.3.1.4 4. predict_proba(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data. It has shape **(n_samples, n_features)**.

Purpose

Perform classification on an array and return probability estimates for the test vector X.

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server. Unlike sklearn, it performs the classification on an array and returns the probability estimates for the test feature matrix (X).

This method is not available for “hinge” and “squared_loss” function.

For example,

```
# finds the probability sample for each class in the model
sgd_clf.predict_proba(mat)
```

Output

```
[[0. 1.]
 [0. 1.]
 [0. 1.]
 ...
 [0. 1.]
 [0. 1.]
 [0. 1.]]
```

Like in `fit()`, frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# finds the probability sample for each class in the model
sgd_clf.predict_proba(rmat)
```

Output

```
[[0. 1.]
 [0. 1.]
 [0. 1.]
 ...
 [0. 1.]
 [0. 1.]
 [0. 1.]]
```

Return Value

It returns a numpy array of float or double (float64) type and of shape (**n_samples**, **n_classes**) containing the predicted probability values.

29.3.1.5 5. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python array-like object containing the target labels. It has shape (**n_samples**,).

sample_weight: A python narray containing the intended weights for each input samples and it should be the shape of (**n_samples**,).

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample. (Default: None)

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of self.predict(X) wrt. y.

For ‘log’ and ‘hinge’ loss, ‘accuracy_score’ is calculated and for ‘squared_loss’, ‘r2_score’ is calculated.

For example,

```
# calculate mean accuracy score on given test data and labels
sgd_clf.score(mat, lbl)
```

Output

```
0.91
```

Return Value

For ‘log’ and ‘hinge’ loss, it returns ‘accuracy_score’ and for ‘squared_loss’, it returns ‘r2_score’ of double (float64) type.

29.3.1.6 6. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

For ‘squared_loss’ this method doesn’t load the saved file ‘label_map’.

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the SGDClassifier model
sgd_clf.load("./out/SCLFModel")
```

Return Value

It simply returns “self” instance.

29.3.1.7 7. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
sgd_clf.save("./out/SCLFModel")
```

The SCLFModel contains below directory structure:

SCLFModel

```
|——label_map
|——metadata
|——model
```

‘label_map’ contains information about labels mapped with their encoded value. **This information is not saved for ‘squared_loss’ loss function.**

‘metadata’ represents the detail about loss function, model_kind and datatype of training vector.

Here, the model file contains information about model_id, model_kind and datatype of training vector.

This will save the SGDClassifier model on the path ‘/out/SCLFModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

29.3.1.8 8. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by SGDClassifier. It is used to get parameters and their values of SGDClassifier class.

For example,

```
print(sgd_clf.get_params())
```

Output

```
{'alpha': 0.0001, 'average': False, 'class_weight': None,
'early_stopping': False, 'epsilon': 0.1, 'eta0': 1.0, 'fit_intercept': True,
'l1_ratio': 0.15, 'learning_rate': 'invscaling', 'loss': 'hinge', 'max_iter': 1000,
'n_iter_no_change': 5, 'n_jobs': None, 'penalty': 'l2', 'power_t': 0.5,
'random_state': None, 'shuffle': True, 'tol': 0.001, 'validation_fraction': 0.1,
'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

29.3.1.9 9. set_params(**params)

Parameters

****params**: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by SGDClassifier, used to set parameter values.

For example,

```
print("Get parameters before setting:")
print(sgd_clf.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
sgd_clf.set_params(penalty = 'l1', fit_intercept = False)
print("Get parameters before setting:")
print(sgd_clf.get_params())
```

Output

```
Get parameters before setting:
{'alpha': 0.0001, 'average': False, 'class_weight': None, 'early_stopping': False,
'epsilon': 0.1, 'eta0': 1.0, 'fit_intercept': True, 'l1_ratio': 0.15,
'learning_rate': 'invscaling', 'loss': 'hinge', 'max_iter': 1000,
'n_iter_no_change': 5, 'n_jobs': None, 'penalty': 'l2', 'power_t': 0.5,
'random_state': None, 'shuffle': True, 'tol': 0.001, 'validation_fraction': 0.1,
'verbose': 0, 'warm_start': False}
Get parameters before setting:
{'alpha': 0.0001, 'average': False, 'class_weight': None, 'early_stopping': False,
'epsilon': 0.1, 'eta0': 1.0, 'fit_intercept': False, 'l1_ratio': 0.15,
'learning_rate': 'invscaling', 'loss': 'hinge', 'max_iter': 1000,
'n_iter_no_change': 5, 'n_jobs': None, 'penalty': 'l1', 'power_t': 0.5,
'random_state': None, 'shuffle': True, 'tol': 0.001, 'validation_fraction': 0.1,
'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

29.3.1.10 10. debug_print()

Purpose

It shows the target model information(weight values, intercept, etc.) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
sgd_clf.debug_print()
```

Output:

```
----- Weight Vector:: -----
14072.2 22454.1 83330.6 45451.2 139.907 -12.3658 -191.893 -82.7364 265.627 109.118
46.0452 1510.72 -250.729 -38123.1 9.74564 -4.16035 -16.0092 0.837207 26.3526 3.59399
14821.1 29161.9 85356 -57710.4 183.668 -99.8164 -348.61 -70.554 386.649 113.928
Intercept:: 1845.32
Threshold:: 0
```

This output will be visible on server side. It displays the weights, intercept and threshold values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

29.3.1.11 11. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
sgd_clf.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

29.3.1.12 12. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, predict() is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

29.4 SEE ALSO

- Introduction to FrovedisRowmajorMatrix
- Introduction to FrovedisColmajorMatrix
- Introduction to FrovedisCRSMatrix
- Introduction to FrovedisDvector
- SGDRegressor in Frovedis

Chapter 30

SGDRegressor

30.1 NAME

SGDRegressor - A regression algorithm used to predict the labels with various loss functions. This estimator implements regularized linear models with stochastic gradient descent (SGD) learning.

30.2 SYNOPSIS

```
class frovedis.mllib.linear_model.SGDRegressor(loss='squared_loss', penalty='l2', alpha=0.0001,
                                              l1_ratio=0.15, fit_intercept=True,
                                              max_iter=1000, tol=1e-3,
                                              shuffle=True, verbose=0,
                                              epsilon=0.1, random_state=None,
                                              learning_rate="invscaling",
                                              eta0=0.001, power_t=0.25,
                                              early_stopping=False,
                                              validation_fraction=0.1,
                                              n_iter_no_change=5,
                                              warm_start=False,
                                              average=False)
```

30.2.1 Public Member Functions

```
fit(X, y, coef_init = None, intercept_init = None, sample_weight = None)
predict(X)
score(X, y, sample_weight = None)
load(fname, dtype = None)
save(fname)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

30.3 DESCRIPTION

Stochastic Gradient Descent (SGD) is used for discriminative learning of linear regressors under convex loss functions such as SVM and Logistic regression. It has been successfully applied to large-scale datasets because the update to the coefficients is performed for each training instance, rather than at the end of instances.

Stochastic Gradient Descent (SGD) regressor basically implements a plain SGD learning routine supporting various loss functions and penalties for regression.

It implements regularized linear models with stochastic gradient descent (SGD) learning: the gradient of the loss is estimated each sample at a time and the model is updated along the way with a decreasing strength schedule. It is a linear method which uses the following loss functions:

- 1) **squared_loss**
- 2) **epsilon_insensitive**
- 3) **squared_epsilon_insensitive**

It supports ZERO, L1 and L2 regularization to address the overfit problem.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn SGDRegressor (Stochastic Gradient Descent Regression) interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for SGDRegressor on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

30.3.1 Detailed Description

30.3.1.1 1. SGDRegressor()

Parameters

loss: A string object parameter containing the loss function type to use. Currently, frovedis supports 'squared_loss', 'epsilon_insensitive' and 'squared_epsilon_insensitive' functions. (Default: 'squared_loss')

penalty: A string object parameter containing the regularizer type to use. Currently none, l1 and l2 are supported by Frovedis. (Default: 'l2')

If it is None (not specified explicitly), it will be set as 'ZERO' regularization type.

alpha: Zero or a positive double (float64) smoothing parameter. (Default: 0.0001)

l1_ratio: An unused parameter. (Default: 0.15)

fit_intercept: A boolean parameter specifying whether a constant (intercept) should be added to the decision function. (Default: True)

max_iter: A positive integer parameter specifying maximum iteration count. (Default: 1000)

tol: A double (float64) parameter specifying the convergence tolerance value. It must be zero or a positive value. (Default: 1e-3)

shuffle: An unused parameter. (Default: True)

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

epsilon: A zero or positive double (float64) parameter used in the epsilon-insensitive loss function. (Default: 0.1)

random_state: An unused parameter. (Default: None)

learning_rate: A string object parameter containing the learning rate. (Default: ‘invscaling’)

Unlike sklearn, Frovedis only supports ‘invscaling’ learning rate. ‘invscaling’ gradually decreases the learning rate ‘**learning_rate**’ at each time step ‘**t**’ using an inverse scaling exponent of ‘**power_t**’.

learning_rate = $\text{eta0} / \text{pow}(\text{t}, \text{power_t})$

eta0: A double (float64) parameter specifying the initial learning rate for the ‘invscaling’ and ‘optimal’ schedules. (Default: 0.001)

power_t: An unused parameter which specifies the exponent for inverse scaling learning rate. Although, this parameter is unused in frovedis but it must be of double (float64) type. This is simply done to keep the behavior consistent with Scikit-learn. (Default: 0.25)

early_stopping: An unused parameter. (Default: False)

validation_fraction: An unused parameter. (Default: 0.1)

n_iter_no_change: An unused parameter. (Default: 5)

warm_start: A boolean parameter which when set to True, reuses the solution of the previous call to fit as initialization, otherwise, just erase the previous solution. (Default: False)

average: An unused parameter. (Default: False)

Attributes

coef: It is a python ndarray (containing float or double (float64) typed values depending on data-type of input matrix (X)). It is the weights assigned to the features. It is of shape (**n_features**).

intercept: It is a python ndarray (float or double (float64) values depending on input matrix data type) and has shape (**1**). It specifies the constants in decision function.

_n_iter: An integer value used to get the actual iteration point at which the problem is converged.

Purpose

It initializes a SGDRegressor object with the given parameters.

The parameters: “l1_ratio”, “shuffle”, “random_state”, “early_stopping”, “validation_fraction”, “n_iter_no_change” and “average” are simply kept to make the interface uniform to Scikit-learn SGDRegressor module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

30.3.1.2 2. fit(X, y, coef_init = None, intercept_init = None, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python ndarray or an instance of FrovedisCRSMMatrix for sparse data and FrovedisColmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python ndarray or an instance of FrovedisDvector containing the target values. It has shape (**n_samples**).

coef_init: An unused parameter that specifies the initial coefficients to warm-start the optimization. (Default: None)

intercept_init: An unused parameter that specifies the initial intercept to warm-start the optimization. (Default: None)

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It accepts the training feature matrix (X) and corresponding target values (y) as inputs from the user and trains a SGDRegressor model with specified regularization with those data at frovedis server.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_diabetes
```

```
mat, lbl = load_diabetes(return_X_y = True)

# fitting input matrix and label on SGDRegressor object
from frovedis.mllib.linear_model import SGDRegressor
sgd_reg = SGDRegressor(max_iter = 50000000, loss = "squared_epsilon_insensitive",
                       penalty = 'none', tol = 0.0000001).fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
from sklearn.datasets import load_diabetes
mat, lbl = load_diabetes(return_X_y = True)

# Since "mat" is numpy dense data, we have created FrovedisColmajorMatrix.
# and for scipy sparse data, FrovedisCRSMMatrix should be used.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)

# SGDRegressor with pre-constructed frovedis-like inputs
from frovedis.mllib.linear_model import SGDRegressor
sgd_reg = SGDRegressor(max_iter = 50000000, loss = "squared_epsilon_insensitive",
                       penalty = 'none', tol = 0.0000001).fit(cmat,dlbl)
```

Return Value

It simply returns “self” reference.

30.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
sgd_reg.predict(mat)
```

Output:

```
[187.30000503 79.39151311 168.59790769 155.75933185 129.99656706
 89.71373962 96.12127367 152.37303585 163.56868847 181.57865709
 .....
 .....
 123.18827921 64.13627628 185.64720615 136.37181437 139.34641795
 185.9899156 72.21133803]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "cmat" is FrovedisColmajorMatrix, we have created FrovedisRowmajorMatrix.
# predicting on SGDRegressor using frovedis-like input
sgd_reg.predict(cmat.to_frovedis_rowmatrix())
```

Output

```
[187.30000503  79.39151311 168.59790769 155.75933185 129.99656706
 89.71373962  96.12127367 152.37303585 163.56868847 181.57865709
 .....
 .....
123.18827921  64.13627628 185.64720615 136.37181437 139.34641795
185.9899156  72.21133803]
```

Return Value

It returns a numpy array of double (float64) type containing the predicted values. It is of shape (**n_samples**,).

30.3.1.4 4. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python ndarray or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python ndarray containing the target values. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

Calculate the root mean square value on the given test data and labels i.e. R2(r-squared) of self.predict(X) wrt. y.

The coefficient 'R2' is defined as $(1 - (u/v))$,

where 'u' is the residual sum of squares $((y_true - y_pred) ** 2).sum()$ and

'v' is the total sum of squares $((y_true - y_true.mean()) ** 2).sum()$.

The best possible score is 1.0 and it can be negative (because the model can be arbitrarily worse). A constant model that always predicts the expected value of y, disregarding the input features, would get a R2 score of 0.0.

For example,

```
# calculate R2 score on given test data and labels
sgd_reg.score(mat, lbl)
```

Output

```
0.52
```

Return Value

It returns an R2 score of double (float64) type.

30.3.1.5 5. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the SGDRegressor model
sgd_reg.load("./out/SGDRegressorModel")
```

Return Value

It simply returns “self” reference.

30.3.1.6 6. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
sgd_reg.save("./out/SGDRegressorModel")
```

The **SGDRegressorModel** contains below directory structure:

SGDRegressorModel

```
|— metadata
|— model
```

‘metadata’ represents the detail about loss function, model_kind and datatype of training vector.

The ‘model’ file contains the SGDRegressor model saved in binary format.

This will save the SGDRegressor model on the path ‘/out/SGDRegressorModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

30.3.1.7 7. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by SGDRegressor. It is used to get parameters and their values of SGDRegressor class.

For example,

```
print(sgd_reg.get_params())
```

Output

```
{'alpha': 0.0001, 'average': False, 'early_stopping': False, 'epsilon': 0.1,
'eta0': 0.001, 'fit_intercept': True, 'l1_ratio': 0.15, 'learning_rate': 'invscaling',
'loss': 'squared_loss', 'max_iter': 1000, 'n_iter_no_change': 5, 'penalty': 'l2',
'power_t': 0.25, 'random_state': None, 'shuffle': True, 'tol': 0.001,
'validation_fraction': 0.1, 'verbose': 0, 'warm_start': False}
```

Return Value

A dictionary of parameter names mapped to their values.

30.3.1.8 8. set_params(params)****Parameters**

****params:** All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by SGDRegressor, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(sgd_reg.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
sgd_reg.set_params(penalty = 'l1', fit_intercept = False)
print("get parameters before setting:")
print(sgd_reg.get_params())
```

Output

```
get parameters before setting:
{'alpha': 0.0001, 'average': False, 'early_stopping': False, 'epsilon': 0.1,
 'eta0': 0.001, 'fit_intercept': True, 'l1_ratio': 0.15, 'learning_rate': 'invscaling',
 'loss': 'squared_loss', 'max_iter': 1000, 'n_iter_no_change': 5, 'penalty': 'l2',
 'power_t': 0.25, 'random_state': None, 'shuffle': True, 'tol': 0.001,
 'validation_fraction': 0.1, 'verbose': 0, 'warm_start': False}
get parameters before setting:
{'alpha': 0.0001, 'average': False, 'early_stopping': False, 'epsilon': 0.1,
 'eta0': 0.001, 'fit_intercept': False, 'l1_ratio': 0.15, 'learning_rate': 'invscaling',
 'loss': 'squared_loss', 'max_iter': 1000, 'n_iter_no_change': 5, 'penalty': 'l1',
 'power_t': 0.25, 'random_state': None, 'shuffle': True, 'tol': 0.001,
 'validation_fraction': 0.1, 'verbose': 0, 'warm_start': False}
```

Return Value

It simply returns “self” reference.

30.3.1.9 9. debug_print()**Purpose**

It shows the target model information (weight values, intercept, etc.) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
sgd_reg.debug_print()
```

Output:

```
----- Weight Vector:: -----
-8.26881e+10 -1.23538e+11 -1.81876e+11 -9.355e+08 -8.27686e+09 -8.80985e+10 -1.0433e+12
-4.84075e+10 -1.69485e+11 -6.51606e+12 -2.66758e+11 -4.63121e+12 -2.0491e+11
Intercept:: -1.42647e+10
```

This output will be visible on server side. It displays the weights and intercept values on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

30.3.1.10 10. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
sgd_reg.release()
```

This will reset the after-fit populated attributes (like `coef__`, `intercept__`, `n_iter__`) to `None`, along with releasing server side memory.

Return Value

It returns nothing.

30.3.1.11 11. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, `predict()` is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

30.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisColmajorMatrix**
- **Introduction to FrovedisCRSMatrix**
- **Introduction to FrovedisDvector**
- **SGDRegressor in Frovedis**

Chapter 31

Spectral Clustering

31.1 NAME

Spectral Clustering - A clustering algorithm commonly used in EDA (exploratory data analysis). It uses the spectrum (eigenvalues) of the similarity matrix of the data to perform clustering.

31.2 SYNOPSIS

```
class frovedis.mllib.cluster.SpectralClustering(n_clusters=8, eigen_solver=None,
                                                n_components=None, random_state=None,
                                                n_init=10, gamma=1.0, affinity='rbf',
                                                n_neighbors=10, eigen_tol=0.0,
                                                assign_labels='kmeans', degree=3, coef0=1,
                                                kernel_params=None, n_jobs=None, verbose=0,
                                                max_iter=300, eps=1e-4, norm_laplacian=True,
                                                mode=3, drop_first=True)
```

31.2.1 Public Member Functions

```
fit(X, y = None)
fit_predict(X, y = None)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
load(fname, dtype = None)
save(fname)
debug_print()
release()
is_fitted()
```

31.3 DESCRIPTION

Clustering is an unsupervised learning problem where we aim to group subsets of entities with one another based on some notion of similarity.

In Spectral Clustering, the data points are treated as nodes of a graph. Thus, clustering is treated as a graph partitioning problem. The nodes are then mapped to a low-dimensional space that can be easily segregated to form clusters.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn Spectral Clustering interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Spectral Clustering on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

31.3.1 Detailed Description

31.3.1.1 1. SpectralClustering()

Parameters

n_clusters: A positive integer parameter specifying the number of clusters. The number of clusters should be greater than 0 and less than or equal to *n_samples*. (Default: 8)

eigen_solver: A string object parameter. It is the eigenvalue decomposition strategy to use. (Default: None)

When it is None (not specified explicitly), it will be set as 'arpack'. Only 'arpack' eigen solver is supported.

n_components: A positive integer parameter containing the number of components for clusters. It is used to compute the number of eigenvectors for spectral embedding. The number of components should be in between 1 to *n_features*. (Default: None)

When it is None (not specified explicitly), it will be equal to the number of clusters.

random_state: Zero or positive integer parameter. It is None by default.

When it is None (not specified explicitly), it will be set as 0. (unused)

n_init: A positive integer parameter that specifies the number of times the k-means algorithm will be run with different centroid seeds. (Default: 10)

gamma: The double (float64) parameter required for computing nearby relational meaningful eigenvalues. (Default: 1.0)

When it is None (specified explicitly), it will be set as 1.0.

Kernel coefficient for rbf is "[np.exp(-gamma * d(X,X) ** 2)]" kernel. Ignored for affinity='nearest_neighbors'.

affinity: A string object parameter which tells how to construct the affinity matrix. (Default: 'rbf')

When it is None (specified explicitly), it will be set as 'rbf'. Only 'rbf', 'nearest_neighbors' and 'precomputed' are supported.

- **'nearest_neighbors'**: construct the affinity matrix by computing a graph of nearest neighbors.

- **'rbf'**: construct the affinity matrix using a radial basis function (RBF) kernel.

- **'precomputed'**: interpret X as a precomputed affinity matrix, where larger values indicate greater similarity between instances.

n_neighbors: A positive integer parameter that specifies the number of neighbors to be used when constructing the affinity matrix using the nearest neighbors method. It must be in between 1 to *n_samples*. It is applicable only when affinity = 'nearest_neighbors'. (Default: 10)

eigen_tol: Stopping criterion for eigen decomposition of the Laplacian matrix when using 'arpack' eigen_solver. (unused)

assign_labels: A string object parameter that specifies the strategy to use to assign labels in the embedding

space. When it is None (specified explicitly), it will be set as 'kmeans'. Only 'kmeans' is supported. (Default: 'kmeans')

degree: Degree of the polynomial kernel. (unused)

coef0: Zero coefficient for polynomial and sigmoid kernels. (unused)

kernel_params: Parameters (keyword arguments) and values for kernel. (unused)

n_jobs: The number of parallel jobs to run. (unused)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

max_iter: A positive integer parameter containing the maximum number of iteration count for kmeans assignment. (Default: 300)

eps: Zero or a positive double parameter containing the tolerance value for kmeans. (Default: 1e-4)

norm_laplacian: A boolean parameter if set to True, then compute normalized Laplacian, else not. (Default: True)

mode: An integer parameter required to set the eigen computation method. It can be either 1 (for generic) or 3 (for shift-invert). It is applicable only for dense data. For more details refer ARPACK computation modes. (Default: 3)

drop_first: A boolean parameter if set to True, then drops the first eigenvector. The first eigenvector of a normalized Laplacian is full of constants, thus if it is set to True, then (n_components + 1) eigenvectors are computed and will drop the first vector. Otherwise, it will calculate 'n_components' number of eigenvectors. (Default: True)

Attributes

affinity_matrix_:

1. For python native dense input:

- When affinity = 'precomputed/rbf', it returns a numpy array
- When affinity = 'nearest_neighbors', it returns a scipy matrix

2. For frovedis-like dense input:

- When affinity = 'precomputed/rbf', returns a FrovedisRowmajorMatrix
- When affinity = 'nearest_neighbors', returns a FrovedisCRSMMatrix

3. For python native sparse input:

- When affinity = 'precomputed/nearest_neighbors', it returns a scipy matrix
- When affinity = 'rbf', it returns a numpy array

4. For frovedis-like sparse input:

- When affinity = 'precomputed/nearest_neighbors', it a returns FrovedisCRSMMatrix
- When affinity = 'rbf', it returns a FrovedisRowmajorMatrix

In all cases, the output is of float or double (float64) type and of shape (**n_samples**, **n_samples**).

labels_: A python ndarray of int64 type values and has shape (**n_clusters**,). It contains the predicted cluster labels for each point.

Purpose

It initializes a Spectral Clustering object with the given parameters.

The parameters: "eigen_tol", "degree", "coef0", "kernel_params" and "n_jobs", "random_state" are simply kept in to make the interface uniform to the Scikit-learn Spectral Clustering module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns "self" reference.

31.3.1.2 2. fit(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. If affinity="precomputed", it needs to be of

shape (**n_samples**, **n_samples**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, as in Scikit-learn as well.

Purpose

It clusters the given data points (X) into a predefined number of clusters.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/spectral_data.txt")

# fitting input matrix on SpectralClustering object
from frovedis.mllib.cluster import SpectralClustering
spec = SpectralClustering(n_clusters = 2).fit(mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/spectral_data.txt")

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCSRMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# Spectral Clustering with pre-constructed frovedis-like inputs
from frovedis.mllib.cluster import SpectralClustering
spec = SpectralClustering(n_clusters = 2).fit(rmat)
```

Return Value

It simply returns “self” reference.

31.3.1.3 3. fit_predict(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCSRMatrix for sparse data and FrovedisRowmajorMatrix for dense data. If affinity=“precomputed”, it needs to be of shape (**n_samples**, **n_samples**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, as in Scikit-learn as well.

Purpose

It fits the given data points (X) and returns the predicted labels based on cluster formed during the fit.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/spectral_data.txt")

# fitting input matrix on Spectral Clustering object
from frovedis.mllib.cluster import SpectralClustering
spec = SpectralClustering(n_clusters = 2)
print(spec.fit_predict(mat))
```

Output

```
[0 0 1 1 1]
```

It prints the predicted cluster labels after training is completed.

Like in `fit()`, we can also provide frovedis-like input in `fit_predict()` for faster computation.

For example,

```
# loading sample matrix data
mat = np.loadtxt("./input/sample_data.txt")

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# using pre-constructed input matrix
from frovedis.mllib.cluster import SpectralClustering
spec = SpectralClustering(n_clusters = 2)
print(spec.fit_predict(rmat))
```

Output

```
[0 0 1 1 1]
```

It prints the predicted cluster labels after training is completed.

Return Value

It returns a numpy array of int32 type values containing the cluster labels. It has a shape **(n_samples,)**.

31.3.1.4 4. `score(X, y, sample_weight = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of `FrovedisCRSMMatrix` for sparse data and `FrovedisRowmajorMatrix` for dense data. If `affinity="precomputed"`, it needs to be of shape **(n_samples, n_samples)**.

y: A python ndarray or an instance of `FrovedisVector` containing the true labels for X. It has shape **(n_samples,)**.

sample_weight: An unused parameter whose default value is `None`. It is simply ignored in frovedis implementation.

Purpose

It uses homogeneity score on given true labels and predicted labels i.e homogeneity score of `self.predict(X, y)` wrt. `y`.

For example,

```
spec.score(train_mat, [0, 0, 1, 1, 1])
```

Output

```
1.0
```

Return Value

It returns a homogeneity score of float type.

31.3.1.5 5. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by SpectralClustering. It is used to get parameters and their values of SpectralClustering class.

For example,

```
print(spec.get_params())
```

Output

```
{'affinity': 'rbf', 'assign_labels': 'kmeans', 'coef0': 1, 'degree': 3,
'drop_first': True, 'eigen_solver': 'arpack', 'eigen_tol': 0.0, 'eps': 0.0001,
'gamma': 1.0, 'kernel_params': None, 'max_iter': 300, 'mode': 3, 'n_clusters': 2,
'n_components': 2, 'n_init': 10, 'n_jobs': None, 'n_neighbors': 10,
'norm_laplacian': True, 'random_state': None, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

31.3.1.6 6. `set_params(**params)`

Parameters

*****params***: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by SpectralClustering, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(spec.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
spec.set_params(n_clusters = 3, affinity = 'precomputed')
print("get parameters after setting:")
print(spec.get_params())
```

Output

```
get parameters before setting:
{'affinity': 'rbf', 'assign_labels': 'kmeans', 'coef0': 1, 'degree': 3,
'drop_first': True, 'eigen_solver': 'arpack', 'eigen_tol': 0.0, 'eps': 0.0001,
'gamma': 1.0, 'kernel_params': None, 'max_iter': 300, 'mode': 3, 'n_clusters': 2,
'n_components': 2, 'n_init': 10, 'n_jobs': None, 'n_neighbors': 10,
'norm_laplacian': True, 'random_state': None, 'verbose': 0}
get parameters after setting:
{'affinity': 'precomputed', 'assign_labels': 'kmeans', 'coef0': 1, 'degree': 3,
'drop_first': True, 'eigen_solver': 'arpack', 'eigen_tol': 0.0, 'eps': 0.0001,
'gamma': 1.0, 'kernel_params': None, 'max_iter': 300, 'mode': 3, 'n_clusters': 3,
'n_components': 2, 'n_init': 10, 'n_jobs': None, 'n_neighbors': 10,
'norm_laplacian': True, 'random_state': None, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

31.3.1.7 7. load(fname, dtype = None)**Parameters**

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads a spectral clustering model stored previously from the specified file (having little-endian binary data).

For example,

```
spec.load("./out/MySpecClusteringModel", dtype = np.float64)
```

Return Value

It simply returns “self” reference.

31.3.1.8 8. save(fname)**Parameters**

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the spectral clustering model
spec.save("./out/MySpecClusteringModel")
```

This will save the spectral clustering model on the path “/out/MySpecClusteringModel”.

It would raise exception if the directory already exists with same name.

The ‘MySpecClusteringModel’ directory has

MySpecClusteringModel

```
|—metadata
|—model
|   |— aff_type
|   |— affinity
|   |— cluster_size
|   |— label
```

The metadata file contains the number of clusters, number of components, model kind, input datatype used for trained model.

Here, the **model** directory contains information about affinity type, labels, cluster size and **affinity** matrix (sparse or dense, depending upon python/frovedis input and affinity is ‘rbf’, ‘precomputed’ or ‘nearest_neighbors’).

Return Value

It returns nothing.

31.3.1.9 9. debug_print()**Purpose**

It shows the target model information (affinity matrix) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
spec.debug_print()
```

Output

```
affinity matrix:
num_row = 5, num_col = 5
node 0
node = 0, local_num_row = 5, local_num_col = 5, val = 1 0.970446 6.2893e-104 2.92712e-106
1.28299e-108 0.970446 1 1.27264e-101 6.2893e-104 2.92712e-106 6.2893e-104 1.27264e-101 1
0.970446 0.88692 2.92712e-106 6.2893e-104 0.970446 1 0.970446 1.28299e-108 2.92712e-106
0.88692 0.970446 1
labels:
0 0 1 1 1
ncluster: 2
```

This output will be visible on server side. It displays the affinity matrix.

No such output will be visible on client side.

Return Value

It returns nothing.

31.3.1.10 10. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
spec.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

31.3.1.11 11. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

31.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**

- **Introduction to FrovedisCRSMatrix**
- **Agglomerative Clustering in Frovedis**
- **DBSCAN in Frovedis**
- **KMeans in Frovedis**

Chapter 32

Spectral Embedding

32.1 NAME

Spectral Embedding - One of the accurate method for extraction of meaningful patterns in high dimensional data. It forms an affinity matrix given by the specified function and applies spectral decomposition to the corresponding graph laplacian. The resulting transformation is given by the value of the normalized eigenvectors for each data point.

32.2 SYNOPSIS

```
class frovedis.mllib.manifold.SpectralEmbedding(n_components=2, affinity='nearest_neighbors',
                                                gamma=1.0, random_state=None, eigen_solver=None,
                                                n_neighbors=None, n_jobs=None, verbose=0,
                                                norm_laplacian=True, mode=3, drop_first=True)
```

32.2.1 Public Member Functions

```
fit(X, y = None)
load(fname, dtype = None)
save(fname)
get_params(deep = True)
set_params(**params)
debug_print()
release()
is_fitted()
```

32.3 DESCRIPTION

Spectral embedding is particularly useful for reducing the dimensionality of data that is expected to lie on a low-dimensional manifold contained within a high-dimensional space. It yields a low-dimensional representation of the data that best preserves the structure of the original manifold in the sense that points that are close to each other on the original manifold will also be close after embedding. At the same time, the embedding emphasizes clusters in the original data.

This module provides a client-server implementation, where the client application is a normal python program. Frovedis is almost same as Scikit-learn manifold module providing Spectral Embedding support, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Spectral Embedding on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, the python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

32.3.1 Detailed Description

32.3.1.1 1. SpectralEmbedding()

Parameters

- n_components***: An integer parameter containing the number of component count. (Default: 2)
- affinity***: A string object parameter which specifies how to construct the affinity matrix. (Default: 'nearest_neighbors')
- **'nearest_neighbors'**: construct the affinity matrix by computing a graph of nearest neighbors.
- **'precomputed'**: interpret X as a precomputed affinity matrix, where larger values indicate greater similarity between instances.

Only 'nearest_neighbors' and 'precomputed' are supported.

gamma: The double (float64) parameter required for computing nearby relational meaningful eigenvalues. (Default: 1.0)

random_state: An unused parameter. (Default: None)

eigen_solver: An unused parameter. (Default: None)

n_neighbors: An unused parameter. (Default: None)

n_jobs: An unused parameter. (Default: None)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (for INFO mode and not specified explicitly). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

norm_laplacian: A boolean parameter if set to True, then computes the normalized Laplacian. (Default: True)

mode: An integer parameter required to set the eigen computation method. It can be either 1 (for generic) or 3 (for shift-invert). (Default: 3)

drop_first: A boolean parameter if set to True, then drops the first eigenvector. The first eigenvector of a normalized Laplacian is full of constants, thus if it is set to true, then (n_components + 1) eigenvectors are computed and will drop the first vector. Otherwise, it will calculate 'n_components' number of eigenvectors. (Default: True)

Attribute

affinity_matrix:

1. **For python native dense input:**
 - When affinity = 'precomputed', it returns a numpy array
 - When affinity = 'nearest_neighbors', it returns a numpy array
2. **For frovedis-like dense input:**

- When affinity = ‘precomputed’, returns a FrovedisRowmajorMatrix
- When affinity = ‘nearest_neighbors’, returns a FrovedisRowmajorMatrix
- 3. **For python native sparse input:**
 - When affinity = ‘precomputed’, it returns a scipy matrix
 - When affinity = ‘nearest_neighbors’, it returns a numpy array
- 4. **For frovedis-like sparse input:**
 - When affinity = ‘nearest_neighbors’, it returns a FrovedisRowmajorMatrix
 - When affinity = ‘precomputed’, it returns a FrovedisRowmajorMatrix

In all cases, the output is of float or double (float64) type and of shape (**n_samples, n_samples**).

embedding_:

1. **For python native dense input:**
 - When affinity = ‘precomputed’, it returns a numpy array
 - When affinity = ‘nearest_neighbors’, it returns a numpy array
2. **For frovedis-like dense input:**
 - When affinity = ‘precomputed’, returns a FrovedisRowmajorMatrix
 - When affinity = ‘nearest_neighbors’, returns a FrovedisRowmajorMatrix
3. **For python native sparse input:**
 - When affinity = ‘precomputed’, it returns a numpy array
 - When affinity = ‘nearest_neighbors’, it returns a numpy array
4. **For frovedis-like sparse input:**
 - When affinity = ‘nearest_neighbors’, it returns a FrovedisRowmajorMatrix
 - When affinity = ‘precomputed’, it returns a FrovedisRowmajorMatrix

In all cases, the output is of float or double (float64) type and of shape (**n_samples, n_components**).

Note: affinity = ‘precomputed’ should be used with square matrix input only, Otherwise, it throws an exception.

Purpose

It initializes a Spectral Embedding object with the given parameters.

The parameters: ‘random_state’, ‘eigen_solver’, ‘n_neighbors’ and ‘n_jobs’ are simply kept in to make the interface uniform to the Scikit-learn Spectral Embedding module. They are not used anywhere within frovedis implementation.

After getting the affinity matrix by computing distance co-relation, this is used to extract meaningful patterns in high dimensional data.

Return Value

It simply returns “self” reference.

32.3.1.2 2. fit(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples, n_features**).

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It extracts meaningful or co-related patterns obtained from normalized eigenvector computation.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("spectral_data.txt")

# fitting input matrix on Spectral Embedding object
from frovedis.mllib.manifold import SpectralEmbedding
sem = SpectralEmbedding(n_components = 2, drop_first = True).fit(train_mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading sample matrix data
train_mat = np.loadtxt("spectral_data.txt")

# Since "train_mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(train_mat)

# fitting input matrix on Spectral Embedding object
from frovedis.mllib.manifold import SpectralEmbedding
sem = SpectralEmbedding(n_components = 2, drop_first = True).fit(rmat)
```

Return Value

It simply returns “self” reference.

32.3.1.3 3. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the same model
sem.load("./out/MySemModel", dtype=np.float64)
```

Return Value

It simply returns “self” instance.

32.3.1.4 4. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information(metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
sem.save("./out/MySemModel")
```

The MySemModel contains below directory structure:

```
MySemModel
├── metadata
├── model
│   ├── aff_type
│   ├── affinity
│   └── embedding
```

‘metadata’ represents the detail about n_components, model_kind and datatype of training vector. Here, the **model** directory contains information about type of affinity matrix, affinity matrix and embedding matrix.

If the directory already exists with the same name then it will raise an exception.

Return Value

It returns nothing.

32.3.1.5 5. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by Spectral Embedding. It is used to get parameters and their values of Spectral Embedding class.

For example,

```
print(sem.get_params())
```

Output

```
{'affinity': 'nearest_neighbors', 'drop_first': True, 'eigen_solver': None, 'gamma': 1.0,
 'mode': 3, 'n_components': 2, 'n_jobs': None, 'n_neighbors': None, 'norm_laplacian': True,
 'random_state': None, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

32.3.1.6 6. set_params(**params)

Parameters

****params:** All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by Spectral Embedding, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(sem.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
sem.set_params(n_components = 3, drop_first = False)
print("get parameters after setting:")
print(sem.get_params())
```

Output

```
get parameters before setting:
{'affinity': 'nearest_neighbors', 'drop_first': True, 'eigen_solver': None,
'gamma': 1.0, 'mode': 3, 'n_components': 2, 'n_jobs': None, 'n_neighbors': None,
'norm_laplacian': True, 'random_state': None, 'verbose': 0}
get parameters after setting:
{'affinity': 'nearest_neighbors', 'drop_first': False, 'eigen_solver': None,
'gamma': 1.0, 'mode': 3, 'n_components': 3, 'n_jobs': None, 'n_neighbors': None,
'norm_laplacian': True, 'random_state': None, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

32.3.1.7 7. debug_print()

Purpose

It shows the target model information (affinity and embed matrix) on the server side user terminal. It is mainly used for debugging purpose.

For example,

```
sem.debug_print()
```

Output

```
affinity matrix:
num_row = 5, num_col = 5
node 0
node = 0, local_num_row = 5, local_num_col = 5, val = 1 0.970446 6.2893e-104 2.92712e-106
1.28299e-108 0.970446 1 1.27264e-101 6.2893e-104 2.92712e-106 6.2893e-104 1.27264e-101
1 0.970446 0.88692 2.92712e-106 6.2893e-104 0.970446 1 0.970446 1.28299e-108 2.92712e-106
0.88692 0.970446 1
embed matrix:
num_row = 5, num_col = 2
node 0
node = 0, local_num_row = 5, local_num_col = 2, val = -0.628988 -0.345834 -0.628988
-0.345834 -0.202594 0.368471 -0.202594 0.368471 -0.202594 0.368471
```

This output will be visible on server side. It displays the affinity matrix and embedding matrix on the trained model which is currently present on the server.

No such output will be visible on client side.

Return Value

It returns nothing.

32.3.1.8 8. release()**Purpose**

It can be used to release the in-memory model at frovedis server.

For example,

```
sem.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

32.3.1.9 9. is_fitted()**Purpose**

It can be used to confirm if the model is already fitted or not. In case, affinity_matrix is used before training the model, then it can prompt the user to train the clustering model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

32.4 SEE ALSO

- **Introduction to FrovedisRowmajorMatrix**
- **Introduction to FrovedisCRSMMatrix**
- **Spectral Clustering in Frovedis**

Chapter 33

StandardScaler

33.1 NAME

StandardScaler - It is a preprocessing technique which is used to perform scaling of distribution values so that mean of observed values is zero and the standard deviation is unit variance.

33.2 SYNOPSIS

```
frovedis.mllib.preprocessing.StandardScaler(copy = True, with_mean= True,  
                                             with_std=True, sam_std = False, verbose=0)
```

33.2.1 Public Member Functions

```
fit(X, y = None, sample_weight = None)  
fit_transform(X, y = None)  
get_params(deep = True)  
inverse_transform(X, copy = None)  
is_fitted()  
partial_fit(X, y = None, sample_weight = None)  
release()  
set_params(**params)  
transform(X, copy = None)
```

33.3 DESCRIPTION

This machine learning algorithm standardizes the features by removing the mean and scaling to unit variance.

The standard score of a sample x is calculated as:

$$z = (x - u) / s$$

where,

- u is the mean of the training samples or zero if `with_mean = False`.
- s is the standard deviation of the training samples or one if `with_std = False`.

Centering and scaling happens independently on each feature by computing the relevant statistics on the samples in the training set. Mean and standard deviation are then stored to be used later on during `transform()`.

Currently, this class in frovedis will first create a copy of input data and then will perform standard scaling. It does not support inplace scaling yet.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn StandardScaler interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for StandardScaler on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When transform-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

33.3.1 Detailed Description

33.3.1.1 1. StandardScaler()

Parameters

copy: This is an unused parameter. (Default: True)

with_mean: It is a boolean parameter that specifies whether to perform centering before scaling of data. (Default: True)

When it is True (not specified explicitly), it will center the data before scaling.

Note: Also, use 'with_mean = False' when attempting centering on sparse matrices. Otherwise it raises an exception.

When it is set as None (specified explicitly), it will be set as False.

with_std: It is a boolean parameter that specifies whether to scale the data to unit variance or not. (Default: True)

When it is True (not specified explicitly), it will scale the data to unit variance.

When it is set as None (specified explicitly), it will be set as False.

sam_std: It is a boolean parameter that specifies whether to enable unbiased or biased sample standard deviation. (Default: False)

- If it is False (not specified explicitly), it will compute biased standard deviation (where $1 / n_samples$ is used).
- If it is True (specified explicitly), it will compute unbiased standard deviation (where $1 / (n_samples - 1)$ is used).

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

Attributes

mean_: It is a numpy ndarray of shape (**n_features**,) and having double (float64) type values. It specifies the mean value for each feature in the training set.

var_: It is a numpy ndarray of shape (**n_features**,) and having double (float64) type values. It specifies the variance for each feature in the training set.

scale_: It is a numpy ndarray of shape (**n_features**,) and having double (float64) type values. It specifies the per feature relative scaling of the data to achieve zero mean and unit variance.

Purpose

It initializes a StandardScaler object with the given parameters.

The parameters: “copy” is simply kept in to make the interface uniform to the Scikit-learn StandardScaler module. They are not used in frovedis implementation internally.

Return Value

It simply returns “self” reference.

33.3.1.2 2. fit(X, y = None, sample_weight = None)**Parameters**

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float (float32) or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float (float32) or double (float64) type. It has shape (**n_samples**, **n_features**).
y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: None or any python array-like object (any shape). It is simply ignored in frovedis implementation.

Purpose

It computes the mean and standard deviation to be used for later scaling.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                  [0.2, 0.2, 0.2],
                  [9., 9., 9. ],
                  [9.1, 9.1, 9.1],
                  [9.2, 9.2, 9.2]])

# fitting input matrix on StandardScaler object
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler().fit(mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                  [0.2, 0.2, 0.2],
                  [9., 9., 9. ],
                  [9.1, 9.1, 9.1],
                  [9.2, 9.2, 9.2]])

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# StandardScaler with pre-constructed frovedis-like inputs
```

```
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler().fit(rmat)
```

Return Value

It simply returns “self” reference.

33.3.1.3 3. fit_transform(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float (float32) or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float (float32) or double (float64) type. It has shape (**n_samples**, **n_features**).
y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation.

Purpose

It will fit the model with input matrix (X) and then returns a transformed version of input matrix (X).

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                 [0.2, 0.2, 0.2],
                 [9., 9., 9. ],
                 [9.1, 9.1, 9.1],
                 [9.2, 9.2, 9.2]])

# fitting input matrix on StandardScaler object and perform transform
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler()
print(ss.fit_transform(mat))
```

Output

```
[[ -1.23598774 -1.23598774 -1.23598774]
 [ -1.21318353 -1.21318353 -1.21318353]
 [  0.79358622  0.79358622  0.79358622]
 [  0.81639042  0.81639042  0.81639042]
 [  0.83919462  0.83919462  0.83919462]]
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                 [0.2, 0.2, 0.2],
                 [9., 9., 9. ],
                 [9.1, 9.1, 9.1],
                 [9.2, 9.2, 9.2]])

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# Fitting StandardScaler with pre-constructed frovedis-like inputs and perform transform
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler()
print(ss.fit_transform(rmat))
```

Output

```
[[-1.23598774 -1.23598774 -1.23598774]
 [-1.21318353 -1.21318353 -1.21318353]
 [ 0.79358622  0.79358622  0.79358622]
 [ 0.81639042  0.81639042  0.81639042]
 [ 0.83919462  0.83919462  0.83919462]]
```

Return Value

- **When dense data is used as input:**
For both frovedis-like input and python input, it returns a numpy matrix of shape (**n_samples**, **n_features**) and double (float64) type values. It contains transformed values.
- **When sparse data is used as input:**
For both frovedis-like input and python input, it returns a scipy sparse matrix of shape (**n_samples**, **n_features**) and double (float64) type values. It contains transformed values.

33.3.1.4 4. `get_params(deep = True)`

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by StandardScaler. It is used to get parameters and their values of StandardScaler class.

For example,

```
print(ss.get_params())
```

Output

```
{'copy': True, 'sam_std': False, 'verbose': 0, 'with_mean': True, 'with_std': True}
```

Return Value

A dictionary of parameter names mapped to their values.

33.3.1.5 5. `inverse_transform(X, copy = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float (float32) or double (float64) type values. It can also be an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float (float32) or double (float64) type. It has shape (**n_samples**, **n_features**).
copy: This is an unused parameter.

Purpose

It scales back the data to the original representation.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                 [0.2, 0.2, 0.2],
                 [9., 9., 9. ],
                 [9.1, 9.1, 9.1],
                 [9.2, 9.2, 9.2]])

# fitting input matrix on StandardScaler object and perform transform
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler()
ss.fit(mat)
X1 = ss.transform(mat)

# inverse_transform() demo to get original input data
print(ss.inverse_transform(X1))
```

Output

```
[[0.1 0.1 0.1]
 [0.2 0.2 0.2]
 [9.  9.  9. ]
 [9.1 9.1 9.1]
 [9.2 9.2 9.2]]
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                 [0.2, 0.2, 0.2],
                 [9., 9., 9. ],
                 [9.1, 9.1, 9.1],
                 [9.2, 9.2, 9.2]])

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# StandardScaler with pre-constructed frovedis-like inputs and perform transform
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler()
ss.fit(rmat)
X1 = ss.transform(rmat)

# inverse_transform() demo to get original frovedis-like input data
print(ss.inverse_transform(X1))
```

Output

```
[[0.1 0.1 0.1]
 [0.2 0.2 0.2]]
```



```
[9.  9.  9. ]
[9.1 9.1 9.1]
[9.2 9.2 9.2]]
```

Return Value

- **When dense data is used as input:**
For both frovedis-like input and python input, it returns a numpy matrix of shape (**n_samples**, **n_features**) and double (float64) type values. It contains transformed values (original data).
- **When sparse data is used as input:**
For both frovedis-like input and python input, it returns a scipy sparse matrix of shape (**n_samples**, **n_features**) and double (float64) type values. It contains transformed values (original data).

33.3.1.6 6. `is_fitted()`

Purpose

It can be used to confirm if the model is already fitted or not. In case, `transform()` is used before training the model, then it can prompt the user to train the pca model first.

Return Value

It returns 'True', if the model is already fitted otherwise, it returns 'False'.

33.3.1.7 7. `partial_fit(X, y = None, sample_weight = None)`

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float (float32) or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float (float32) or double (float64) type. It has shape (**n_samples**, **n_features**).
y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

sample_weight: None or any python array-like object (any shape). It is simply ignored in frovedis implementation.

Purpose

It performs incremental computation of mean and standard deviation on each new batch of samples present in input matrix (X) and the finalised mean and standard deviation computed is later used for scaling.

All of input matrix (X) is processed as a single batch. This is intended for cases when fit is not feasible due to very large number of 'n_samples' or because input matrix (X) is read from a continuous stream.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                 [0.2, 0.2, 0.2],
                 [9., 9., 9. ],
                 [9.1, 9.1, 9.1],
                 [9.2, 9.2, 9.2]])

# partial_fit() demo with input matrix on StandardScaler object
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler().partial_fit(mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                 [0.2, 0.2, 0.2],
                 [9., 9., 9. ],
                 [9.1, 9.1, 9.1],
                 [9.2, 9.2, 9.2]])

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# StandardScaler with pre-constructed frovedis-like inputs
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler().partial_fit(rmat)
```

Return Value

It simply returns “self” reference.

33.3.1.8 8. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
ss.release()
```

This will reset the after-fit populated attributes to None, along with releasing server side memory.

Return Value

It returns nothing.

33.3.1.9 9. set_params(**params)

Parameters

****params**: All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by StandardScaler, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(ss.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
ss.set_params(with_mean = False, sam_std = False, verbose = 1)
print("get parameters after setting:")
print(ss.get_params())
```

Output

```
get parameters before setting:
{'copy': True, 'sam_std': False, 'verbose': 0, 'with_mean': True, 'with_std': True}
get parameters after setting:
{'copy': True, 'sam_std': False, 'verbose': 1, 'with_mean': False, 'with_std': True}
```

Return Value

It simply returns “self” reference.

33.3.1.10 10. transform(X, copy = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float (float32) or double (float64) type values. It can also be an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float (float32) or double (float64) type. It has shape (**n_samples**, **n_features**).
copy: This is an unused parameter.

Purpose

It performs standardization by centering and scaling.

It will always perform transform on a copy of input matrix (X).

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                  [0.2, 0.2, 0.2],
                  [9., 9., 9. ],
                  [9.1, 9.1, 9.1],
                  [9.2, 9.2, 9.2]])

# fitting input matrix on StandardScaler object and perform transform
from frovedis.ml.lib.preprocessing import StandardScaler
ss = StandardScaler()
ss.fit(mat)
print(ss.transform(mat))
```

Output

```
[[ -1.23598774 -1.23598774 -1.23598774]
 [ -1.21318353 -1.21318353 -1.21318353]
 [  0.79358622  0.79358622  0.79358622]
 [  0.81639042  0.81639042  0.81639042]
 [  0.83919462  0.83919462  0.83919462]]
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0.1, 0.1, 0.1],
                  [0.2, 0.2, 0.2],
                  [9., 9., 9. ],
                  [9.1, 9.1, 9.1],
                  [9.2, 9.2, 9.2]])
```

```

[9.2, 9.2, 9.2]])

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# Fitting StandardScaler with pre-constructed frovedis-like inputs and perform transform
from frovedis.mllib.preprocessing import StandardScaler
ss = StandardScaler()
ss.fit(rmat)
print(ss.transform(rmat))

```

Output

```

[[-1.23598774 -1.23598774 -1.23598774]
 [-1.21318353 -1.21318353 -1.21318353]
 [ 0.79358622  0.79358622  0.79358622]
 [ 0.81639042  0.81639042  0.81639042]
 [ 0.83919462  0.83919462  0.83919462]]

```

Return Value

- **When dense data is used as input:**
For both frovedis-like input and python input, it returns a numpy matrix of shape (**n_samples**, **n_features**) and double (float64) type values. It contains transformed values.
- **When sparse data is used as input:**
For both frovedis-like input and python input, it returns a scipy sparse matrix of shape (**n_samples**, **n_features**) and double (float64) type values. It contains transformed values.

33.4 SEE ALSO

- [Introduction to FrovedisRowmajorMatrix](#)
- [Introduction to FrovedisCRSMMatrix](#)

Chapter 34

SVC

34.1 NAME

SVC (Support Vector Classification) - A classification algorithm used to predict the binary output with different kernel functions.

34.2 SYNOPSIS

```
class frovedis.mllib.svm.SVC(C=1.0, kernel='rbf', degree=3, gamma='scale', coef0=0.0,
                             shrinking=True, probability=False, tol=0.001, cache_size=128,
                             class_weight=None, verbose=False, max_iter=-1,
                             decision_function_shape='ovr', break_ties=False,
                             random_state=None)
```

34.2.1 Public Member Functions

```
fit(X, y, sample_weight = None)
predict(X)
predict_proba(X) load(fname, dtype = None)
save(fname)
score(X, y, sample_weight = None)
get_params(deep = True)
set_params(**params)
release()
is_fitted()
```

34.3 DESCRIPTION

Support vector machines (SVMs) are a set of supervised learning methods used for classification and regression.

During training, the input X is the training data and y are their corresponding label values (Frovedis supports any values as for labels, but internally it encodes the input binary labels to -1 and 1, before training at Frovedis server) which we want to predict. **Frovedis supports only binary SVC classification algorithm.**

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn SVC (Support Vector Classification) interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for SVC on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When prediction-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

34.3.1 Detailed Description

34.3.1.1 1. SVC()

Parameters

C: A positive double (float64) parameter, also called as inverse of regularization strength. (Default: 1.0)

kernel: A string object parameter that specifies the kernel type to be used. Unlike sklearn, frovedis supports 'linear', 'poly', 'rbf' and 'sigmoid' kernel function. (Default: 'rbf')

degree: A positive integer parameter specifying the degree of the polynomial kernel. (Default: 3)

gamma: A string object parameter that specifies the kernel coefficient. (Default: 'scale')

For all python like input gamma = 'scale' will be used, otherwise 'auto' will be used.

Depending on the **kernel coefficient**, **gamma** will be calculated as:

- If **gamma** = 'scale' (default) is used then **gamma** = $1.0 / (n_features * variance)$,
- If **gamma** = 'auto' is used then **gamma** = $1.0 / n_features$

coef0: A double (float64) parameter which is independent term in kernel function. (Default: 0.0)

shrinking: An unused parameter. (Default: 'True')

probability: An unused parameter. (Default: 'False')

tol: A zero or positive double (float64) parameter specifying the convergence tolerance value. (Default: 0.001)

cache_size: An interger parameter that specifies the size of the kernel cache(in megabytes). It must be greater than 2. (Default: 128)

class_weight: An unused parameter. (Default: 'None')

verbose: A boolean parameter that specifies the log level to use. (Default: 'False')

Its value is False by default (for INFO mode). But it can be set to True (for DEBUG mode or TRACE mode) for getting training time logs from frovedis server.

max_iter: A positive integer parameter specifying the hard limit on iterations within solver, or -1 for no limit. (Default: -1)

decision_function_shape: An unused parameter. (Default: 'ovr')

break_ties: An unused parameter. (Default: 'False')

random_state: An unused parameter. (Default: 'None')

Attributes

coef_: It is a python ndarray(containing float or double (float64) typed values depending on data-type of input matrix (X)). It is the weights assigned to the features. It has shape **(1, n_features)**.

This attribute is supported only for 'linear' kernel, otherwise it will be None.

intercept_: It is a python ndarray(float or double (float64) values depending on input matrix data type) and has shape **(1,)**. It specifies the constants in decision function.

This attribute is supported only for 'linear' kernel, otherwise it will be None.

classes_: It is an int64 type python ndarray of unique labels given to the classifier during training. It has shape **(n_classes,)**.

support_: It is an int32 type python ndarray that specifies the support vectors indices. It has shape (**n_support_vectors_**), where **n_support_vectors_** = **len(support_vectors) / n_features**

support_vectors_: It is double (float64) type python ndarray specifies the support vectors. It has shape (**n_support_vectors_**, **n_features**).

Purpose

It initializes a SVC object with the given parameters.

The parameters: “shrinking”, “probability”, “class_weight”, “decision_function_shape”, “break_ties” and “random_state” are simply kept to make the interface uniform to Scikit-learn SVC module. They are not used anywhere within frovedis implementation.

Return Value

It simply returns “self” reference.

34.3.1.2 2. fit(X, y, sample_weight = None)

Parameters

X: A numpy dense or a scipy sparse matrix or any python ndarray. It has shape (**n_samples**, **n_features**). For Frovedis input data:

- if kernel = ‘linear’, then it can be an instance of FrovedisColmajorMatrix for dense data and FrovedisCRS-Matrix for sparse data.
- if kernel != ‘linear’, then it can be an instance of FrovedisRowmajorMatrix.

y: Any python array-like object or an instance of FrovedisDvector containing the target labels. It has shape (**n_samples**).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

It accepts the training feature matrix (X) and corresponding output labels (y) as inputs from the user and trains a SVC model at frovedis server.

Depending upon the input data, usage of ‘gamma’ will vary:

- For **python** dense data, **gamma** = ‘scale’ and **gamma** = ‘auto’ will be used.
- For **python** sparse input, **gamma** = ‘auto’ will be used.
- For all **frovedis** like input, **gamma** = ‘auto’ only will be used.

For example,

```
# loading a sample matrix and labels data
mat = np.array([[ -1.0, -1.0], [-2.0, -1.0], [ 1.0,  1.0], [ 2.0,  1.0]])
lbl = np.array([1.0, 1.0, 2.0, 2.0])

# fitting input matrix and label on SVC object
from frovedis.mllib.svc import SVC
svc = SVC(gamma='auto').fit(mat, lbl)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample matrix and labels data
mat = np.array([[ -1.0, -1.0], [-2.0, -1.0], [ 1.0,  1.0], [ 2.0,  1.0]])
lbl = np.array([1.0, 1.0, 2.0, 2.0])
```

```
# Since "mat" is numpy dense data and kernel = 'rbf' by default,
we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
rmat = FrovedisRowmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)
```

```
# SVC with pre-constructed frovedis-like inputs
from frovedis.mllib.svm import SVC
svc = SVC(gamma='auto').fit(rmat, dlbl)
```

User can also provide FrovedisColmajorMatrix or FrovedisCRSMMatrix as a training data but only with **kernel** = 'linear' and **gamma** = 'auto'.

```
# loading a sample matrix and labels data
mat = np.array([[ -1.0, -1.0], [-2.0, -1.0], [1.0, 1.0], [2.0, 1.0]])
lbl = np.array([1.0, 1.0, 2.0, 2.0])
```

```
# Since "mat" is numpy dense data and kernel = 'linear', we have created FrovedisColmajorMatrix
# For scipy sparse data, FrovedisCRSMMatrix should be used instead.
from frovedis.matrix.dense import FrovedisColmajorMatrix
from frovedis.matrix.dvector import FrovedisDvector
cmat = FrovedisColmajorMatrix(mat)
dlbl = FrovedisDvector(lbl)
```

```
# SVC with pre-constructed frovedis-like inputs
from frovedis.mllib.svm import SVC
svc = SVC(kernel = 'linear', gamma = 'auto').fit(cmat, dlbl)
```

Return Value

It simply returns "self" reference.

34.3.1.3 3. predict(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python ndarray or an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server.

For example,

```
svc.predict(mat)
```

Output:

```
[1. 1. 2. 2.]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# predicting on SVC using frovedis-like input
```



```
svc.predict(rmat)
```

Output

```
[1. 1. 2. 2.]
```

Return Value

It returns a numpy array of int64, float or double (float64) type containing the predicted outputs. It has shape (**n_samples**,).

34.3.1.4 4. predict_proba(X)

Parameters

X: A numpy dense or scipy sparse matrix or any python ndarray or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

Purpose

Perform classification on an array and return probability estimates for the test vector X.

It accepts the test feature matrix (X) in order to make prediction on the trained model at frovedis server. Unlike sklearn, it performs the classification on an array and returns the probability estimates for the test feature matrix (X).

This method is not available for kernel = 'linear'.

For example,

```
# finds the probability sample for each class in the SVC model
svc.predict_proba(mat)
```

Output

```
[[6.17597327e-04 9.99382403e-01]
 [1.98150882e-24 1.00000000e+00]
 [7.31058579e-01 2.68941421e-01]
 [9.38039081e-01 6.19609192e-02]]
```

Like in fit(), frovedis-like input can be used to speed-up the prediction making on the trained model at server side.

For example,

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)
```

```
# finds the probability sample for each class in the SVC model
svc.predict_proba(rmat)
```

Output

```
[[6.17597327e-04 9.99382403e-01]
 [1.98150882e-24 1.00000000e+00]
 [7.31058579e-01 2.68941421e-01]
 [9.38039081e-01 6.19609192e-02]]
```

Return Value

It returns a numpy ndarray of float or double (float64) type and of shape (**n_samples**, **n_classes**) containing the predicted probability values.

34.3.1.5 5. load(fname, dtype = None)

Parameters

fname: A string object containing the name of the file having model information to be loaded.

dtype: A data-type is inferred from the input data. Currently, expected input data-type is either float or double (float64). (Default: None)

Purpose

It loads the model from the specified file(having little-endian binary data).

For example,

```
# loading the SVC model
svc.load("./out/SVCModel")
```

Return Value

It simply returns “self” reference.

34.3.1.6 6. save(fname)

Parameters

fname: A string object containing the name of the file on which the target model is to be saved.

Purpose

On success, it writes the model information (label_map, metadata and model) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# saving the model
svc.save("./out/SVCModel")
```

The **SVCModel** contains below directory structure:

SVCModel

```
|—label_map
|—metadata
|—model
```

‘label_map’ contains information about labels mapped with their encoded value.

‘metadata’ represents the detail about number of classes, model_kind and datatype of training vector.

Here, the model file contains information about model in binary format.

This will save the SVC model on the path ‘/out/SVCModel’. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

34.3.1.7 7. score(X, y, sample_weight = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python ndarray or an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data. It has shape (**n_samples**, **n_features**).

y: Any python ndarray containing the target labels. It has shape (**n_samples**,).

sample_weight: A python ndarray containing the intended weights for each input samples and it should be the shape of (**n_samples**,). (Default: None)

When it is None (not specified explicitly), an uniform weight vector is assigned on each input sample.

Purpose

Calculate mean accuracy on the given test data and labels i.e. mean accuracy of `self.predict(X)` wrt. `y`.

For example,

```
# calculate mean accuracy score on given test data and labels
svc.score(mat, lbl)
```

Output

```
1.0
```

Return Value

It returns an accuracy score of double (float64) type.

34.3.1.8 8. get_params(deep = True)**Parameters**

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by SVC. It is used to get parameters and their values of SVC class.

For example,

```
print(svc.get_params())
```

Output

```
{'C': 1.0, 'break_ties': False, 'cache_size': 128, 'class_weight': None, 'coef0': 0.0,
'decision_function_shape': 'ovr', 'degree': 3, 'gamma': 0.5, 'kernel': 'rbf',
'max_iter': -1, 'probability': False, 'random_state': None, 'shrinking': True,
'tol': 0.001, 'verbose': False}
```

Return Value

A dictionary of parameter names mapped to their values.

34.3.1.9 9. set_params(params)****Parameters**

****params**: All the keyword arguments are passed this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by SVC, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(svc.get_params())
# User just needs to provide the arguments and internally it will create a
dictionary over the arguments given by user
svc.set_params( kernel = "linear", shrinking = False)
print("get parameters after setting:")
print(svc.get_params())
```

Output

```

get parameters before setting:
{'C': 1.0, 'break_ties': False, 'cache_size': 128, 'class_weight': None, 'coef0': 0.0,
 'decision_function_shape': 'ovr', 'degree': 3, 'gamma': 0.5, 'kernel': 'rbf',
 'max_iter': -1, 'probability': False, 'random_state': None, 'shrinking': True,
 'tol': 0.001, 'verbose': False}
get parameters after setting:
{'C': 1.0, 'break_ties': False, 'cache_size': 128, 'class_weight': None, 'coef0': 0.0,
 'decision_function_shape': 'ovr', 'degree': 3, 'gamma': 0.5, 'kernel': 'linear',
 'max_iter': -1, 'probability': False, 'random_state': None, 'shrinking': False,
 'tol': 0.001, 'verbose': False}

```

Return Value

It simply returns “self” reference.

34.3.1.10 10. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
svc.release()
```

This will reset the after-fit populated attributes (like `coef__`, `intercept__`, `classes__`, etc) to `None`, along with releasing server side memory.

Return Value

It returns nothing.

34.3.1.11 11. is_fitted()

Purpose

It can be used to confirm if the model is already fitted or not. In case, `predict()` is used before training the model, then it can prompt the user to train the model first.

Return Value

It returns ‘True’, if the model is already fitted otherwise, it returns ‘False’.

34.4 SEE ALSO

- Introduction to `FrovedisRowmajorMatrix`
- Introduction to `FrovedisColmajorMatrix`
- Introduction to `FrovedisCRSMatrix`
- Introduction to `FrovedisDvector`
- `LinearSVC` in `Frovedis`
- `LinearSVR` in `Frovedis`

Chapter 35

t-Distributed Stochastic Neighbor Embedding

35.1 NAME

TSNE - It's full form is T-distributed Stochastic Neighbor Embedding. It is an unsupervised algorithm primarily used for data exploration and visualizing high-dimensional data.

35.2 SYNOPSIS

```
frovedis.mllib.manifold.tsne(n_components=2, perplexity=30.0,
                             early_exaggeration=12.0, learning_rate=200.0,
                             n_iter=1000, n_iter_without_progress=300,
                             min_grad_norm=1e-7, metric="euclidean",
                             init="random", verbose=0, random_state=None,
                             method="exact", angle=0.5, n_jobs=None)
```

35.2.1 Public Member Functions

```
fit(X, y = None)
fit_transform(X, y = None)
get_params(deep = True)
set_params(**params)
```

35.3 DESCRIPTION

It is a nonlinear dimensionality reduction technique well-suited for embedding high-dimensional data for visualization in a low-dimensional space of two or three dimensions. Specifically, it models each high-dimensional object by a two or three dimensional point in such a way that similar objects are modeled by nearby points and dissimilar objects are modeled by distant points with high probability.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Scikit-learn TSNE interface, but it doesn't have any dependency with Scikit-learn. It can be used simply even if the system doesn't have Scikit-learn installed. Thus in this

implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for TSNE on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When transform-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

35.3.1 Detailed Description

35.3.1.1 1. TSNE()

Parameters

n_components: It is an integer parameter that specifies the dimension of the embedded space. (Default: 2) Currently, it supports '**n_components = 2**' only as parameter value.

perplexity: It must be a positive double (float64) parameter that specifies the number of nearest neighbors. Larger datasets usually require a larger perplexity. Consider selecting a value between 5 and 50. Different values can result in significantly different results. (Default: 30.0)

early_exaggeration: It must be a positive double (float64) parameter that controls how tight natural clusters in the original space are in the embedded space and how much space will be between them. For larger values, the space between natural clusters will be larger in the embedded space. (Default: 12.0)

learning_rate: It accepts the word 'auto' or a positive double (float64) as parameter value that controls the step size of the gradient updates. It is usually in the range [10.0, 1000.0]. **Currently, 'auto' as parameter value cannot be used.** (Default: 200.0)

- **If the 'learning_rate' is too high:** then the data may look like a 'ball' when plotted on a graph with any point approximately equidistant from its nearest neighbours.
- **If the 'learning_rate' is too low:** then most data points may look compressed in a **dense cloud** with few outliers when plotted on a graph.

n_iter: It must be a positive integer value that specifies the maximum number of iterations for the optimization. It must be atleast 250. (Default: 1000)

n_iter_without_progress: It is an integer parameter that specifies the maximum number of iterations without progress before we abort the optimization. (Default: 300)

min_grad_norm: It is a double (float64) parameter that specifies whether in case the gradient norm is below this threshold, then the optimization will be stopped. (Default: 1e-7)

metric: It is a string object parameter that specifies the metric to use when calculating distance between instances in a feature array. It supports '**euclidean**' or '**precomputed**' distances. (Default: 'euclidean')

init: It is a string object parameter that specifies the initialization of embedding. (Default: 'random')

Currently, only random initialization is supported for this method in frovedis.

verbose: An integer parameter specifying the log level to use. Its value is set as 0 by default (for INFO mode). But it can be set to 1 (for DEBUG mode) or 2 (for TRACE mode) for getting training time logs from frovedis server.

random_state: This is an unused parameter. (Default: None)

method: It is a string object parameter that specifies the t-SNE implementation method to use. (Default: 'exact')

- **'exact'**: it calculates the pair-wise distance between every pair of data points. **Currently, only exact implementation of tsne is supported in frovedis.**
- **'barnes_hut'**: it calculates the distance between each data point and its closest neighboring points only. **Currently, this implementation of tsne is not supported in frovedis.**

angle: This is an unused parameter. (Default: 0.5)

n_jobs: This is an unused parameter. (Default: None)

Attributes

n_iter: It is a positive integer value that specifies the number of iterations run.

kl_divergence: It is a double (float64) type value that specifies the Kullback-Leibler divergence after optimization.

embedding: It is a numpy ndarray of double (float64) type values or FrovedisRowmajorMatrix instance, having shape (**n_samples**, **n_components**), where **n_samples** is the number of samples in the input matrix (X). It stores the embedding vectors.

Purpose

It initializes a TSNE object with the given parameters.

The parameters: “random_state”, “angle”, “n_jobs” are simply kept in to make the interface uniform to the Scikit-learn TSNE module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

35.3.1.2 2. fit(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float or double (float64) type values. It can also be an instance of FrovedisCRSMMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**). If **metric = ‘precomputed’**, then input matrix (X) is assumed to be a **squared distance matrix**.

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It will fit input matrix (X) into an embedded space.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0, 0, 0, 0], [0, 1, 1, 1],
                 [1, 0, 1, 0], [1, 1, 1, 0],
                 [1, 1, 1, 1]], dtype = np.float64)
```

```
# fitting input matrix on TSNE object
from frovedis.ml.lib.manifold import TSNE
tsne = TSNE().fit(mat)
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0, 0, 0, 0], [0, 1, 1, 1],
                 [1, 0, 1, 0], [1, 1, 1, 0],
                 [1, 1, 1, 1]], dtype = np.float64)
```

```
# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
```

```
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# TSNE with pre-constructed frovedis-like inputs
from frovedis.mllib.manifold import TSNE
tsne = TSNE().fit(rmat)
```

Return Value

It simply returns “self” reference.

35.3.1.3 3. fit_transform(X, y = None)

Parameters

X: A numpy dense or scipy sparse matrix or any python array-like object of int, float or double (float64) type values. It can also be an instance of FrovedisCRSMatrix for sparse data and FrovedisRowmajorMatrix for dense data of float or double (float64) type. It has shape (**n_samples**, **n_features**). If **metric = ‘precomputed’**, then input matrix (X) is assumed to be a **squared distance matrix**.

y: None or any python array-like object (any shape). It is simply ignored in frovedis implementation, like in Scikit-learn.

Purpose

It will fit input matrix (X) into an embedded space and will return the transformed output.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0, 0, 0, 0], [0, 1, 1, 1],
                 [1, 0, 1, 0], [1, 1, 1, 0],
                 [1, 1, 1, 1]], dtype = np.float64)

# fitting input matrix on TSNE object and perform transform
from frovedis.mllib.manifold import TSNE
tsne = TSNE()
print(tsne.fit_transform(mat))
```

Output

```
[[ -15.15708437 -218.36584687]
 [  31.18096218   62.89769738]
 [ -79.72509032  -59.56844016]
 [ 129.59496966 -137.71666144]
 [ 195.56724491   14.28454464]]
```

When native python data is provided, it is converted to frovedis-like inputs and sent to frovedis server which consumes some data transfer time. Pre-constructed frovedis-like inputs can be used to speed up the training time, especially when same data would be used for multiple executions.

For example,

```
# loading a sample numpy dense data
import numpy as np
mat = np.matrix([[0, 0, 0, 0], [0, 1, 1, 1],
                 [1, 0, 1, 0], [1, 1, 1, 0],
                 [1, 1, 1, 1]], dtype = np.float64)

# Since "mat" is numpy dense data, we have created FrovedisRowmajorMatrix.
```



```
# For scipy sparse data, FrovedisCRSMatrix should be used instead.
from frovedis.matrix.dense import FrovedisRowmajorMatrix
rmat = FrovedisRowmajorMatrix(mat)

# TSNE with pre-constructed frovedis-like inputs and perform transform
from frovedis.mllib.manifold import TSNE
embedding = TSNE().fit_transform(rmat)
embedding.debug_print()
```

Output

```
matrix:
num_row = 5, num_col = 2
node 0
node = 0, local_num_row = 5, local_num_col = 2, val = -15.1571 -218.366 31.181 62.8977 -79.7251
-59.5684 129.595 -137.717 195.567 14.2845
```

Return Value

- **When X is python native input:**
It returns a python ndarray of shape (**n_samples**, **n_components**) and double (float64) type values. It contains the embedding of the training data in low-dimensional space.
- **When X is frovedis-like input:**
It returns a FrovedisRowmajorMatrix instance of shape (**n_samples**, **n_components**) and double (float64) type values, containing the embedding of the training data in low-dimensional space.

35.3.1.4 4. get_params(deep = True)

Parameters

deep: A boolean parameter, used to get parameters and their values for an estimator. If True, it will return the parameters for an estimator and contained subobjects that are estimators. (Default: True)

Purpose

This method belongs to the BaseEstimator class inherited by TSNE. It is used to get parameters and their values of TSNE class.

For example,

```
print(tsne.get_params())
```

Output

```
{'angle': 0.5, 'early_exaggeration': 12.0, 'init': 'random', 'learning_rate': 200.0,
'method': 'exact', 'metric': 'euclidean', 'min_grad_norm': 1e-07, 'n_components': 2,
'n_iter': 1000, 'n_iter_without_progress': 300, 'n_jobs': None, 'perplexity': 30.0,
'random_state': None, 'verbose': 0}
```

Return Value

A dictionary of parameter names mapped to their values.

35.3.1.5 5. set_params(**params)

Parameters

****params:** All the keyword arguments are passed to this function as dictionary. This dictionary contains parameters of an estimator with its given values to set.

Purpose

This method belongs to the BaseEstimator class inherited by TSNE, used to set parameter values.

For example,

```
print("get parameters before setting:")
print(tsne.get_params())
# User just needs to provide the arguments and internally it will create a
# dictionary over the arguments given by user
tsne.set_params(perplexity = 15, metric = 'precomputed')
print("get parameters after setting:")
print(tsne.get_params())
```

Output

```
get parameters before setting:
{'angle': 0.5, 'early_exaggeration': 12.0, 'init': 'random', 'learning_rate': 200.0,
'method': 'exact', 'metric': 'euclidean', 'min_grad_norm': 1e-07, 'n_components': 2,
'n_iter': 1000, 'n_iter_without_progress': 300, 'n_jobs': None, 'perplexity': 30.0,
'random_state': None, 'verbose': 0}
get parameters after setting:
{'angle': 0.5, 'early_exaggeration': 12.0, 'init': 'random', 'learning_rate': 200.0,
'method': 'exact', 'metric': 'precomputed', 'min_grad_norm': 1e-07, 'n_components': 2,
'n_iter': 1000, 'n_iter_without_progress': 300, 'n_jobs': None, 'perplexity': 15.0,
'random_state': None, 'verbose': 0}
```

Return Value

It simply returns “self” reference.

Note: In order to release the embedding vector from the server (generated by TSNE algorithm in frovedis), we can use `release()` of `FrovedisRowmajorMatrix` class.

For example,

```
tsne.embedding_.release()
```

This will remove the embedding vector from the server side memory.

35.4 SEE ALSO

- [Introduction to FrovedisRowmajorMatrix](#)
- [Introduction to FrovedisCRSMatrix](#)
- [Introduction to FrovedisDvector](#)
- [Spectral Embedding in frovedis](#)

Chapter 36

Word2Vec

36.1 NAME

Word2Vec - Word embedding is one of the most popular representation of document vocabulary. It is capable of capturing context of a word in a document, semantic and syntactic similarity, relation with other words, etc.

36.2 SYNOPSIS

```
class frovedis.mllib.feature.Word2Vec(sentences=None, corpusFile=None,
                                     outDirPath=None, hiddenSize=100,
                                     learningRate=0.025, n_iter=1,
                                     minCount=5, window=5, threshold=1e-3,
                                     negative=5, modelSyncPeriod=0.1, minSyncWords=1024,
                                     fullSyncTimes=0, messageSize=1024, numThreads=None)
```

36.2.1 Public Member Functions

```
build_vocab(corpusIterable = None, corpusFile = None, outDirPath = None, update = False)
build_vocab_and_dump(corpusIterable = None, corpusFile = None, outDirPath = None, update = False)
to_gensim_model()
train(corpusIterable = None, corpusFile = None)
fit(corpusIterable = None, corpusFile = None)
save(modelPath, binary = False)
transform(corpusIterable = None, corpusFile = None, func = None)
fit_transform(corpusIterable = None, corpusFile = None, func = None)
```

36.3 DESCRIPTION

Word2vec is a two-layer neural net that processes text by “vectorizing” words. Its input is a text corpus and its output is a set of vectors (feature vectors that represent words in that corpus). While Word2vec is not a deep neural network, it turns text into a numerical form that deep neural networks can understand.

Word2vec's applications extend beyond parsing sentences in the wild. It can be applied just as well to genes, code, likes, playlists, social media graphs and other verbal or symbolic series in which patterns may be discerned.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface is almost same as Gensim Word2Vec interface. It needs to be used when a system has Gensim installed. Thus in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Word2Vec on the frovedis server. Once the training is completed with the input data at the frovedis server, it returns an abstract model with a unique model ID to the client python program.

When transform-like request would be made on the trained model, python program will send the same request to the frovedis server. After the request is served at the frovedis server, the output would be sent back to the python client.

36.3.1 Detailed Description

36.3.1.1 1. Word2Vec()

Parameters

sentences: This parameter is an iterable of list of strings. If not provided, the 'corpusFile' parameter must be provided for creating the model. (Default: None)

corpusFile: A string object parameter which specifies the path to a corpus file. If not provided, the 'sentences' parameter must be provided for creating the model. (Default: None)

outDirPath: A string object parameter specifying the path of output directory. The newly built vocabulary generated from the input data file is dumped into the provided output file path. (Default: None)

hiddenSize: An integer parameter specifying the dimensionality of the word vectors. For fast computation, this value should be an even value and less than 512 during training. (Default: 100)

learningRate: An unused parameter specifying the initial learning rate. (Default: 0.025)

n_iter: An unused parameter specifying the number of iterations over the corpus. (Default: 1)

minCount: An integer parameter that ignores all words with total frequency lower than this value during vocabulary building. This value should be within 1 to largest total frequency of a word in vocabulary. (Default: 5)

window: An integer parameter specifying the maximum distance between the current and predicted word within a sentence. For fast computation, this value should be less than or equal to 8 during training. (Default: 5)

threshold: An unused parameter specifying the threshold for configuring which higher-frequency words are randomly downsampled. (Default: 1e-3)

negative: An integer parameter. This value specifies how many "noise words" should be drawn (usually between 5-20). The default value should be used for fast computation during training. (Default: 5)

modelSyncPeriod: An unused parameter specifying the model synchronous period. (Default: 0.1)

minSyncWords: An unused parameter specifying the minimum number of words to be synced at each model sync. (Default: 1024)

fullSyncTimes: An unused parameter specifying the full-model sync-up time during training. (Default: 0)

messageSize: An unused parameter specifying the message size in megabytes. (Default: 1024)

numThreads: An integer parameter specifying the number of worker threads to be used to train the model. Ideally, the number of threads should range between 1 and 8. (Default: None)

In case the environment variable 'VE_OMP_NUM_THREADS' is defined, then the value for 'numThreads' would be derived from VE_OMP_NUM_THREADS, otherwise, 'numThreads' would default to value 1.

Attributes

wv: The trained words are stored in KeyedVectors (mapping between keys such as words and embeddings as arrays) instance. It contains the mapping between words and embeddings.

For example,

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]

# Training the Word2Vec model using build_vocab() and train()
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(data)
wv_model.train(data)
print(wv_model.wv)
```

Output

```
{'cat': array([ 0.00400269,  0.0044194 , -0.00383026, ...,  0.0015239 ,  0.00305939,
                0.00019363]), 'say': array([ 0.00227966, -0.00495255,  0.00431488, ...,  0.00019882,
                -0.0016861 , -0.00112656])}
```

It displays words as keys and embeddings as arrays.

Purpose

It initializes a Word2Vec object with the given parameters.

The parameters: “learningRate”, “n_iter”, “threshold”, “modelSyncPeriod”, “minSyncWords”, “fullSyncTimes” and “messageSize” are simply kept in to make the interface uniform to the gensim Word2Vec module. They are not used anywhere within the frovedis implementation.

Return Value

It simply returns “self” reference.

36.3.1.2 2. build_vocab(corpusIterable = None, corpusFile = None, outDirPath = None, update = False)

Parameters

corpusIterable: Use the ‘sentences’ iterable which itself must be an iterable of list of list of tokens. If None, the ‘corpusFile’ must be provided for building the vocabulary. (Default: None)

corpusFile: A string object parameter specifying the path of a corpus file. If None, the ‘corpusIterable’ must be provided for building the vocabulary. (Default: None)

outDirPath: A string object parameter specifying the path of output directory ‘./out’. The newly built vocabulary generated from the input data file is dumped into provided output file path. (Default: None)

update: A boolean parameter if set to True, will add the new words present in sentences to the model’s vocabulary. (Default: False)

Purpose

It builds the vocabulary from input data file and dumped into provided output files. It also initializes the ‘wv’ attribute.

For example, building vocabulary from an iterable

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]

# Building vocabulary from input iterable data
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusIterable = data)
```

For example, building vocabulary from a text file

```
# Using a text file
textfile = "./input/text8-10k"
modelpath = "./out/text_model.txt"

# Building vocabulary from input text file
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusFile = textfile, outDirPath = modelpath)
```

Return Value

It simply returns “self” reference.

36.3.1.3 3. build_vocab_and_dump(corpusIterable = None, corpusFile = None, outDirPath = None, update = False)

Parameters

corpusIterable: Use the ‘sentences’ iterable which itself must be an iterable of list of list of tokens. If None, the ‘corpusFile’ must be provided for building the vocabulary. (Default: None)

corpusFile: A string object parameter specifying the path to a corpus file. If None, the ‘corpusIterable’ must be provided for building the vocabulary. (Default: None)

outDirPath: A string object parameter specifying the path to output directory ‘./out’. The newly build vocabulary generated from the input data file is dumped into provided output file path. (Default: None)

update: A boolean parameter if set to True, will add the new words present in sentences to the model’s vocabulary. (Default: False)

Purpose

It builds the vocabulary from input data file and dump into provided output files. It also initializes the ‘wv’ attribute. This method is an alias to build_vocab().

For example, building vocabulary from an iterable

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]

# Building vocabulary from input iterable data
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab_and_dump(corpusIterable = data)
```

For example, building vocabulary from a text file

```
# Using a text file
textfile = "./input/text8-10k"
modelpath = "./out/text_model.txt"

# Building vocabulary from input text file
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab_and_dump(corpusFile = textfile, outDirPath = modelpath)
```

Return Value

It simply returns “self” reference.

36.3.1.4 4. to_gensim_model()**Purpose**

It generates a gensim like 'wv' (KeyedVectors instance) attribute for Word2Vec model.

For example,

```
wv_model.to_gensim_model()
```

Note:- In order to use this method, gensim version installed by users needs to be 4.0.1 or above.

Return Value

It returns gensim like KeyedVectors instance.

36.3.1.5 5. train(corpusIterable = None, corpusFile = None)**Parameters**

corpusIterable: Use the 'sentences' iterable which itself must be an iterable of list of list of tokens. If None, the 'corpusFile' must be provided for building the vocabulary. (Default: None)

corpusFile: A string object parameter specifying the path to a corpus file. If None, the 'corpusIterable' must be provided for building the vocabulary. (Default: None)

Purpose

It trains the Word2Vec model on input vocabulary.

For example, training with an iterable data

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]
```

```
# Training the model
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusIterable = data)
wv_model.train(corpusIterable = data)
```

For example, training with a text file

```
# Using a text file
textfile = "./input/text8-10k"
modelpath = "./out/text_model.txt"
```

```
# Training the model
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusFile = textfile, outDirPath = modelpath)
wv_model.train(corpusFile = textfile)
```

Return Value

It simply returns "self" reference.

36.3.1.6 6. fit(corpusIterable = None, corpusFile = None)**Parameters**

corpusIterable: Use the 'sentences' iterable which itself must be an iterable of list of list of tokens. If None, the 'corpusFile' must be provided for building the vocabulary. (Default: None)

corpusFile: A string object parameter specifying the path to a corpus file. If None, the 'corpusIterable' must be provided for building the vocabulary. (Default: None)

Purpose

It trains the Word2Vec model on input vocabulary. This method is an alias to `train()`.

For example, training with an iterable data

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]

# Training the model
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusIterable = data)
wv_model.fit(corpusIterable = data)
```

For example, training with a text file

```
# Using a text file
textfile = "./input/text8-10k"
modelpath = "./out/text_model.txt"

# Training the model
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusFile = textfile, outDirPath = modelpath)
wv_model.fit(corpusFile = textfile)
```

Return Value

It simply returns “self” reference.

36.3.1.7 7. save(modelPath, binary = False)**Parameters**

modelPath: A string object parameter specifying the path of the output file in order to save the embeddings.

binary: A boolean parameter if set to True, will save the data in binary format, otherwise, it will be saved in plain text. (Default: False)

Purpose

It saves the word2vec model information to a file.

On success, it writes the model information (after-fit populated attribute like ‘wv’) in the specified file as little-endian binary data. Otherwise, it throws an exception.

For example,

```
# To save the word embeddings in a file
model = "./out/text_model.txt"
wv_model.save(model, binary = False)
```

This will save the word2vec model information on the path “/out/text_model.txt”.

It would raise exception if the ‘text_model.txt’ file already existed with same name.

The ‘text_model.txt’ file contains the count of all the words with total frequency greater than ‘minCount’, ‘hiddenSize’ and dictionary having words (as keys) and embeddings (as values).

Return Value

It returns nothing.

36.3.1.8 8. transform(corpusIterable = None, corpusFile = None, func = None)**Parameters**

corpusIterable: Use the ‘sentences’ iterable which itself must be an iterable of list of list of tokens. If None, the ‘corpusFile’ must be provided for building the vocabulary. (Default: None)

corpusFile: A string object parameter specifying the path to a corpus file. If None, the ‘corpusIterable’ must be provided for building the vocabulary. (Default: None)

func: A function to apply on document vector. (Default: None) If None, then np.mean() is used as ‘func’.

Purpose

It transforms the document text to word2vec embeddings.

For example, training with an iterable

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]

# Training the model
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusIterable = data)
wv_model.fit(corpusIterable = data)
embeddings = wv_model.transform(corpusIterable = data)
print(embeddings)
```

Output

```
[[ 0.00227966 -0.00495255  0.00431488 ...  0.00019882 -0.0016861
  -0.00112656]
 [ 0.00227966 -0.00495255  0.00431488 ...  0.00019882 -0.0016861
  -0.00112656]]
```

For example, training with a text file

```
# Using a text file
textfile = "./input/text8-10k"

# Training the model
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusFile = textfile)
wv_model.fit(corpusFile = textfile)
embeddings = wv_model.transform(corpusFile = textfile)
print(embeddings)
```

Output

```
[[-0.00013626 -0.00059639  0.00063703 ...  0.0004084  -0.00033636
  -0.00013291]]
```

Return Value

It returns document embeddings (numpy array) of shape (n_samples, hiddenSize).

36.3.1.9 9. fit_transform(corpusIterable = None, corpusFile = None, func = None)**Parameters**

corpusIterable: Use the ‘sentences’ iterable which itself must be an iterable of list of list of tokens. If None, the ‘corpusFile’ must be provided for building the vocabulary. (Default: None)

corpusFile: A string object parameter specifying the path to a corpus file. If None, the ‘corpusIterable’

must be provided for building the vocabulary. (Default: None)

func: A function to apply on document vector. (Default: None) If None, then `np.mean()` is used as 'func'.

Purpose

It trains the word2vec model on input document and transforms the document text to word2vec embeddings.

For example, training with an iterable

```
# Using an iterable data
data = [["cat", "say", "meow"], ["dog", "say", "woof"]]

# Training the model
from frovedis.mllib import Word2Vec
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusIterable = data)
embeddings = wv_model.fit_transform(corpusIterable = data)
print(embeddings)
```

Output

```
[[ 0.00227966 -0.00495255  0.00431488 ...  0.00019882 -0.0016861
  -0.00112656]
 [ 0.00227966 -0.00495255  0.00431488 ...  0.00019882 -0.0016861
  -0.00112656]]
```

For example, with a text file

```
# Using a text file
textfile = "./input/text8-10k"

# Training the model
wv_model = Word2Vec(minCount = 2)
wv_model.build_vocab(corpusFile = textfile)
embeddings = wv_model.fit_transform(corpusFile = textfile)
print(embeddings)
```

Output

```
[[-0.00013626 -0.00059639  0.00063703 ...  0.0004084  -0.00033636
  -0.00013291]]
```

Return Value

It returns document embeddings (numpy array) of shape (**n_samples**, **hiddenSize**).

Chapter 37

linalg

37.1 NAME

linalg - a frovedis module which provides user-friendly interfaces for commonly used linear algebra functions.

37.2 SYNOPSIS

```
import frovedis.linalg.linalg
```

37.3 Public Member Functions

```
1. dot(a, b, out=None)
2. eigsh(A, M=None, k=6, sigma=None, which='LM', v0=None, ncv=None, maxiter=None,
    tol=0., return_eigenvectors=True, Minv=None, OPinv=None, mode='normal')
3. inv(a)
4. matmul(x1, x2, out=None, casting='same_kind', order='K', dtype=None,
    subok=True, signature=None, extobj=None)
5. solve(a, b)
6. svd(a, full_matrices=False, compute_uv=True)
```

37.4 DESCRIPTION

The frovedis linear algebra functions rely on PBLAS and ScaLAPACK wrappers in frovedis to provide efficient low level implementations of standard linear algebra algorithms. These functions are used to compute matrices and vector product, matrix decomposition, solving linear equations, inverting matrices, etc.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis interface for all linear algebra functions is almost same as numpy linear algebra function interfaces, but it doesn't have any dependency on numpy. It can be used simply even if the system doesn't have numpy installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python matrix data to frovedis server side. Python data is converted into frovedis compatible data (blockcyclic matrix) internally and then python client can request frovedis server for any of the supported linalg functions on that matrix. Once the operation is completed, the frovedis server sends back the resultant matrix as equivalent python data.

37.4.1 Detailed Description

37.4.1.1 1. dot(a, b, out = None)

Parameters

a: It accepts scalar values and python array-like inputs (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

b: It accepts scalar values, and python array-like inputs (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

out: A numpy ndarray of int, float (float32) or double (float64) type values which is where the result is written. (Default: None)

When it is None (not specified explicitly), a freshly-allocated array having double (float64) type values is returned. Otherwise if provided, it must have a shape that matches the signature $(n,k),(k,m) \rightarrow (n,m)$ and same datatype as input matrices.

Purpose

It computes dot product of two arrays.

1. If both inputs ‘a’ and ‘b’ are scalar values,

For example,

```
# dot() demo with scalar values as input
from frovedis.linalg import dot
prod = dot(3,4)
print(prod)
```

Output

12

Here, result is also a scalar value.

2. If both ‘a’ and ‘b’ are 1D arrays,

For example,

```
import numpy as np
from frovedis.linalg import dot
vec1 = np.array([1., 9., 8.])
vec2 = np.array([4., 1., 7.])

# dot() demo with both inputs as 1D array
prod = dot(vec1,vec2)
print(prod)
```

Output

69.0

Here, dot product is the inner product of vectors.

3. If either ‘a’ or ‘b’ is a scalar value,

For example,

```
import numpy as np
from frovedis.linalg import dot
vec1 = np.array([[1, 9, 8], [7, 1, 5], [1, 2, 4]])
```

```
# dot() demo with 1D array and scalar value as input
prod = dot(vec1,4)
print(prod)
```

Output

```
[[ 4. 36. 32.]
 [28.  4. 20.]
 [ 4.  8. 16.]]
```

Here, result is multiplication of scalar over the vector.

4. If both a and b are 2D arrays,

For example,

```
import numpy as np
from frovedis.linalg import dot
mat1 = np.array([[1, 9, 8], [7, 1, 5], [1, 2, 4]])
mat2 = np.array([[4, 1, 7], [2, 2, 9], [1, 2, 3]])
```

```
# dot() demo with 2D arrays as input
prod = dot(mat1, mat2)
print(prod)
```

Output

```
[[ 30.  35. 112.]
 [ 35.  19.  73.]
 [ 12.  13.  37.]]
```

Here, result is simply matrix multiplication.

When both 'a' and 'b' inputs are matrices or instances of `FrovedisBlockCyclicMatrix`, then the column size of 'a' must be same as row size of 'b' before computing dot product as internally it performs matrix-matrix multiplication.

5. If a is an ND array and b is a 1-D array,

Currently, frovedis supports only $N \leq 2$, so that dot() between ND array and 1D array can be interpreted currently as:

- dot product between both 1D arrays.
- dot product between 2D array and 1D array.

For example,

```
import numpy as np
from frovedis.linalg import dot
mat1 = np.array([[1, 9, 8], [7, 1, 5], [1, 2, 4]])
mat2 = np.array([4, 2, 1])
```

```
# dot() demo with 'a' as 2D array, 'b' as 1D array as input
prod = dot(mat1, mat2)
print(prod)
```

Output

```
[30. 35. 12.]
```

In above example, result will be the sum product over the last axis of 'a' and 'b'.

Currently, dot product between inputs 'a' (an N-D array) and 'b' (an M-D array (where $M \geq 2$)) is not supported in frovedis.

For example,

```
import numpy as np
from frovedis.linalg import dot
mat1 = np.array([[1., 9., 8.], [7., 1., 5.], [1., 2., 4.]])
mat2 = np.array([[4., 1., 7.], [2., 2., 9.], [1., 2., 3.]])

# dot() demo with out != None
prod = dot(mat1, mat2, out = np.array([[1., 1., 1.], [1., 1., 1.], [1., 1., 1.])))
print(prod)
```

Output

```
[[ 30.  35. 112.]
 [ 35.  19.  73.]
 [ 12.  13.  37.]]
```

Here, the result will be written in the user provided ndarray and shape and datatype same as input matrix.

For example,

```
import numpy as np
from frovedis.linalg import dot
mat1 = np.array([[1, 9, 8], [7, 1, 5], [1, 2, 4]])
mat2 = np.array([[4, 1, 7], [2, 2, 9], [1, 2, 3]])

# creating FrovedisBlockcyclicMatrix instance
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
A_Bmat = FrovedisBlockcyclicMatrix(mat1, np.float64)
B_Bmat = FrovedisBlockcyclicMatrix(mat2, np.float64)

# dot() demo with FrovedisBlockcyclicMatrix instance
prod = dot(A_Bmat, B_Bmat)
prod.debug_print()
```

Output

```
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 30 35 12 35 19 13 112 73 37
```

This output will be only visible on server side terminal.

Return Value

It returns the dot product of 'a' and 'b'.

If 'a' and 'b' are both scalars or both 1-D arrays then a scalar is returned. Otherwise, an array is returned.

If out is given, then it is returned.

If either 'a' or 'b' is a FrovedisBlockCyclicMatrix instance, then it returns a FrovedisBlockCyclicMatrix instance.

37.4.1.2 2. eigsh(A, M = None, k = 6, sigma = None, which = 'LM', v0 = None, ncv = None, maxiter = None, tol = 0., return_eigenvectors = True, Minv = None, OPinv = None, mode = 'normal')

Parameters

A: A numpy dense or scipy sparse matrix or any python array-like object or an instance of FrovedisCRSMatrix

for sparse data and FrovedisRowmajorMatrix for dense data. The input matrix provided must be squared symmetric in nature.

M: This is an unused parameter. (Default: None)

k: A positive integer parameter that specifies the number of eigenvalues and eigenvectors desired. It must be in range 0 to N, where N is the number of rows in squared symmetric matrix. (Default: 6)

sigma: It accepts a float (float32) type parameter that is used to find eigenvalues near sigma using shift-invert mode. Currently, it is only supported for dense matrices. (Default: None)

When it is None (not specified explicitly), it will be set as largest possible value of float (float32) type.

which: A string object parameter that specifies which **k** eigenvectors and eigenvalues to find:

- ‘**LM**’ : Largest (in magnitude) eigenvalues. It is the default value.
- ‘**SM**’ : Smallest (in magnitude) eigenvalues.
- ‘**LA**’ : Largest (algebraic) eigenvalues.
- ‘**SA**’ : Smallest (algebraic) eigenvalues.
- ‘**BE**’ : Half (k/2) from each end of the spectrum.

When sigma != None, ‘which’ refers to the shifted eigenvalues.

v0: This is an unused parameter. (Default: None)

ncv: This is an unused parameter. (Default: None)

maxiter: A positive integer that specifies the maximum number of iterations allowed. The convergence may occur on or before the maximum iteration limit is reached. In case, the convergence does not occur before the iteration limit, then this method will not generate the resultant ‘**k**’ eigenvalues and eigenvectors. (Default: None)

When it is None (not specified explicitly), it will be set as 10 * N, where N is the number of rows in squared symmetric matrix.

tol: A float (float32) type parameter that specifies the tolerance values used to find relative accuracy for eigenvalues (stopping criterion). (Default: 0.)

return_eigenvectors: A boolean parameter that specifies whether to return eigenvectors in addition to eigenvalues. (Default: True)

When it is set to False, it will only return **k** eigenvalues.

Minv: This is an unused parameter. (Default: None)

OPinv: This is an unused parameter. (Default: None)

mode: A string object parameter that specifies the strategy to use for shift-invert mode. This parameter applies only when sigma != None. This parameter can be ‘**normal**’, ‘**cayley**’ or ‘**buckling**’ modes.

Currently, frovedis supports only normal mode. (Default: ‘normal’)

The choice of mode will affect which eigenvalues are selected by the ‘which’ parameter.

Purpose

It finds **k** eigenvalues and eigenvectors of the symmetric square matrix A.

The parameters: “M”, “vc0”, “ncv”, “Minv” and “OPinv” are simply kept in to to make the interface uniform to the scipy.sparse.linalg.eigsh() module. They are not used anywhere within the frovedis implementation.

For example,

```
# dense square symmetric matrix
import numpy as np
mat = np.asarray([[2.,-1.,0.,0.,-1.,0.], [-1.,3.,-1.,0.,-1.,0.], [0.,-1.,2.,-1.,0.,0.],
                  [0.,0.,-1.,3.,-1.,-1], [-1.,-1.,0.,-1.,3.,0], [0.,0.,0.,-1.,0.,1.]])

# eigsh() demo
from frovedis.linalg import eigsh
eigen_vals, eigen_vecs = eigsh(mat, k = 3)
print("frovedis computed eigen values:\n")
print(eigen_vals)
print("\nfrovedis computed eigen vectors:\n")
print(eigen_vecs)
```

Output

```
frovedis computed eigen values:
```

```
[3.          3.70462437  4.89121985]
```

```
frovedis computed eigen vectors:
```

```
[[-0.28867513  0.56702343  0.03232265]
 [-0.28867513 -0.65812747  0.46850031]
 [-0.28867513  0.20514371 -0.35640753]
 [ 0.57735027  0.30843449  0.56195221]
 [ 0.57735027 -0.30843449 -0.56195221]
 [-0.28867513 -0.11403968 -0.14441544]]
```

Since, input dense matrix has shape (3,3), so $0 < k \leq 3$.

For example,

```
# eigsh() demo when sigma != None
eigen_vals, eigen_vecs = eigsh(mat, k = 3, sigma = -1)
print("frovedis computed eigen values:\n")
print(eigen_vals)
print("\nfrovedis computed eigen vectors:\n")
print(eigen_vecs)
```

Output

```
frovedis computed eigen values:
```

```
[0.          0.72158639  1.68256939]
```

```
frovedis computed eigen vectors:
```

```
[ [ 0.40824829  0.41486979 -0.50529469]
 [ 0.40824829  0.30944167  0.04026854]
 [ 0.40824829  0.0692328   0.75901025]
 [ 0.40824829 -0.22093352  0.20066454]
 [ 0.40824829  0.22093352 -0.20066454]
 [ 0.40824829 -0.79354426 -0.29398409]]
```

Here, the resultant eigenvalues and eigenvectors are generated using shift invert mode.

For example,

```
# eigsh() demo where which = 'SM'
eigen_vals, eigen_vecs = eigsh(mat, k = 3, which = 'SM')
print("frovedis computed eigen values:\n")
print(eigen_vals)
print("\nfrovedis computed eigen vectors:\n")
print(eigen_vecs)
```

Output

```
frovedis computed eigen values:
```

```
[2.70826973e-16  7.21586391e-01  1.68256939e+00]
```

```
frovedis computed eigen vectors:
```



```
[[ 0.40824829  0.41486979  0.50529469]
 [ 0.40824829  0.30944167 -0.04026854]
 [ 0.40824829  0.0692328  -0.75901025]
 [ 0.40824829 -0.22093352 -0.20066454]
 [ 0.40824829  0.22093352  0.20066454]
 [ 0.40824829 -0.79354426  0.29398409]]
```

Here, the resultant eigenvalues and eigenvectors are smallest in magnitude for the given dense input matrix.

For example,

```
# eigsh() demo where maxiter = 10
eigen_vals, eigen_vecs = eigsh(mat, k = 3, maxiter = 10)
print("frovedis computed eigen values:\n")
print(eigen_vals)
print("\nfrovedis computed eigen vectors:\n")
print(eigen_vecs)
```

Output

frovedis computed eigen values:

```
[3.          3.70462437  4.89121985]
```

scipy computed eigen vectors:

```
[[-0.28867513  0.56702343  0.03232265]
 [-0.28867513 -0.65812747  0.46850031]
 [-0.28867513  0.20514371 -0.35640753]
 [ 0.57735027  0.30843449  0.56195221]
 [ 0.57735027 -0.30843449 -0.56195221]
 [-0.28867513 -0.11403968 -0.14441544]]
```

The eigenvalues and eigenvectors will be generated only when convergence occurs.

In above example, this will happen on or before maxiter = 10 is reached.

For example,

```
# eigsh() demo where return_eigenvectors = False
eigen_vals = eigsh(mat, k = 3, return_eigenvectors = False)
print("frovedis computed eigen values:\n")
print(eigen_vals)
```

Output

frovedis computed eigen values:

```
[3.          3.70462437  4.89121985]
```

Here, only eigenvalues are returned by this method.

Return Value

1. If return_eigenvectors = True:

- It returns eigenvectors in addition to eigenvalues as numpy ndarrays having double (float64) type values. Also, eigenvectors has shape **(k,1)** and eigenvalues has shape **(M,k)**, where **M** is the number of rows of input matrix 'A'.

2. If return_eigenvectors = False:

- It returns only eigenvalues as numpy ndarray having shape **(k,1)** and double (float64) type values.

37.4.1.3 3. inv(a)

Parameters

a: It accepts a python array-like input or numpy matrix (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It specifies the matrix to be inverted. It must be a square matrix of shape (\mathbf{M}, \mathbf{M}) , where \mathbf{M} is the number of rows and columns of input squared matrix.

Purpose

This method computes the inverse matrix of a given matrix.

Internally it uses `getrf()` and `getri()` methods present in `frovedis.linalg.scalapack` module.

For example,

```
import numpy as np
from frovedis.linalg import inv

# input array contains integer values
mat = np.array([[1, 2], [3, 4]])

# inv(0 demo with numpy array as input
matinv = inv(mat)
print(matinv)
```

Output

```
[[ -2.   1. ]
 [ 1.5 -0.5]]
```

It returns an ndarray as the inverse of given matrix having double (float64) type values.

For example,

```
import numpy as np
from frovedis.linalg import inv

# input array contains float (float320 type values
mat = np.array([[1., 2.], [3., 4.]], dtype = np.float32)

# inv() demo with numpy array as input
matinv = inv(mat)
print(matinv)
```

Output

```
[[ -2.0000002  1.0000001 ]
 [ 1.5000001  -0.50000006]]
```

It returns an ndarray as the inverse of given matrix having float (float32) type values.

For example,

```
import numpy as np
from frovedis.linalg import inv

# input array contains double (float64) type values
mat = np.array([[1., 2.], [3., 4.]])

# inv(0 demo with numpy array as input
matinv = inv(mat)
```

```
print(matinv)
```

Output

```
[[ -2.    1. ]
 [ 1.5 -0.5]]
```

It returns an ndarray as the inverse of given matrix having double (float64) type values.

For example,

```
import numpy as np
from frovedis.linalg import inv

# creating FrovedisBlockcyclicMatrix instance
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat = np.array([[1., 2.], [3., 4.]])
A_Bmat = FrovedisBlockcyclicMatrix(mat)

# inv() demo FrovedisBlockcyclicMatrix instance
matinv = inv(A_Bmat)
matinv.debug_print()
```

Output

```
matrix:
num_row = 2, num_col = 2
node 0
node = 0, local_num_row = 2, local_num_col = 2, type = 2, descriptor = 1 1 2 2 2 2 0 0 2 2 2
val = -2 1.5 1 -0.5
```

This output will be only visible on server side terminal.

Return Value

1. When numpy array-like input is used:

- It returns a numpy ndarray having float (float32) or double (float64) type values and has shape (M, M), where M is number of rows and columns of input ndarray.

2. When input is a numpy matrix:

- It returns a numpy matrix having float (float32) or double (float64) type values and has shape (M, M), where M is number of rows and columns of input matrix.

3. When input is a FrovedisBlockCyclicMatrix instance:

- It returns an instance of FrovedisBlockCyclicMatrix.

37.4.1.4 4. matmul(x1, x2, out = None, casting = 'same_kind', order = 'K', dtype = None, subok = True, signature = None, extobj = None)

Parameters

x1: It accepts a python array-like input or numpy maxtrix (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It must not be scalar values.

x2: It accepts scalar values, python array-like inputs or numpy maxtrix (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It must not be scalar values.

out: A numpy ndarray of int, float or double (float64) type values which is where the result is written. (Default: None)

When it is None (not specified explicitly), a freshly-allocated array having double (float64) type values is returned. Otherwise if provided, it must have a shape that matches the signature (n,k),(k,m)->(n,m) and same datatype as input matrices.

casting: This is an unused parameter. (Default: 'same_kind')

order: This is an unused parameter. (Default: 'K')

dtype: This is an unused parameter. (Default: None)

subok: This is an unused parameter. (Default: True)

signature: This is an unused parameter. (Default: None)

extobj: This is an unused parameter. (Default: None)

Purpose

It computes matrix product of two arrays.

The parameters: “casting”, “order”, “dtype”, “subok” and “signature” and “extobj” are simply kept in to to make the interface uniform to the `numpy.matmul()` module. They are not used anywhere within the `frovedis` implementation.

For example,

```
import numpy as np
from frovedis.linalg import matmul

# inputs are having integer type values
mat1 = np.array([[1, 9, 8], [7, 1, 5], [1, 2, 4]])
mat2 = np.array([[4, 1, 7], [2, 2, 9], [1, 2, 3]])

# matmul() demo with numpy array as input
prod = matmul(mat1, mat2)
print(prod)
```

Output

```
[[ 30.  35. 112.]
 [ 35.  19.  73.]
 [ 12.  13.  37.]]
```

Here, the result will be written in a newly generated array having double (float64) type values.

For inputs ‘mat1’ and ‘mat2’, the column size of ‘mat1’ must be same as row size of ‘mat2’ before computing matrix multiplication.

For example,

```
import numpy as np
from frovedis.linalg import matmul

# inputs are having double (float64) type values
mat1 = np.array([[1.5, 9, 8], [7, 1, 5.5], [1, 2.5, 4]])
mat2 = np.array([[4, 1, 7.5], [2.5, 2, 9], [1, 2, 3.5]])

# matmul() demo with numpy array as input
prod = matmul(mat1, mat2)
print(prod)
```

Output

```
[[ 36.5  35.5 120.25]
 [ 36.   20.   80.75]
 [14.25 14.   44.   ]]
```

Here, the result will be written in a newly generated array having double (float64) type values.

For inputs having float (float32) type values, the newly generated array will also have float (float32) type values.

For example,

```
import numpy as np
from frovedis.linalg import matmul

mat1 = np.array([[1., 9., 8.], [7., 1., 5.], [1., 2., 4.]])
mat2 = np.array([[4., 1., 7.], [2., 2., 9.], [1., 2., 3.]])

# matmul() demo with out != None
prod = matmul(mat1, mat2, out = np.array([[1.,1.,1.],[1.,1.,1.],[1.,1.,1.]])
print(prod)
```

Output

```
[[ 30.  35. 112.]
 [ 35.  19.  73.]
 [ 12.  13.  37.]]
```

Here, the result will be written in the user provided ndarray.

For example,

```
import numpy as np
from frovedis.linalg import matmul

mat1 = np.array([[1, 9, 8], [7, 1, 5], [1, 2, 4]])
mat2 = np.array([[4, 1, 7], [2, 2, 9], [1, 2, 3]])

# creating FrovedisBlockcyclicMatrix instance
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
A_Bmat = FrovedisBlockcyclicMatrix(mat1, np.float64)
B_Bmat = FrovedisBlockcyclicMatrix(mat2, np.float64)

# matmul() demo with FrovedisBlockcyclicMatrix instance
prod = matmul(A_Bmat, B_Bmat)
prod.debug_print()
```

Output

```
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 30 35 12 35 19 13 112 73 37
```

This output will be only visible on server side terminal.

Return Value

1. **If any python array like input (having dimensions ≤ 2) and out = None:**
- It returns a numpy ndarray as output and having float (float32) or double (float64) type values.
2. **If any python array like input (having dimensions ≤ 2) and out != None:**
- It returns a numpy ndarray as output. Datatype for output values will be same as datatype of input 'x1' such as int, float (float32) or double (float64) type.
3. **If numpy matrix (having dimensions ≤ 2) and out = None:**
- It returns a numpy matrix as output and having float (float32) or double (float64) type values.
4. **If numpy matrix (having dimensions ≤ 2) and out != None:**
- It returns a numpy matrix as output. Datatype for output values will be same as datatype of input 'x1' such as int, float (float32) or double (float64) type.
5. **If inputs are FrovedisBlockCyclicMatrix instances:**
- It returns a FrovedisBlockCyclicMatrix instance.

37.4.1.5 5. solve(a, b)

Parameters

a: It accepts a python array-like input or numpy maxtrix (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It specifies a coefficient matrix. It must be a square matrix of shape (**nRows**, **nCols**) , where **nRows** = **nCols**.

b: It accepts a python array-like inputs or numpy matrix (having dimensions ≤ 2) of int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It specifies the variables to be determined while solving the equation 'Ax = B'.

Purpose

This function solves a linear matrix equation $Ax = B$, or system of linear scalar equations.

Internally it uses **gesv()** method present in **frovedis.linalg.scalapack** module.

For example,

```
import numpy as np
from frovedis.linalg import solve

# inputs have integer type values
A = np.array([[3,4,3],[1,2,3],[4,2,1]])
B = np.array([[1,1,1], [3,0,2], [8,2,5]])

#solve() demo with numpy array input
X = solve(A,B)
print(X)
```

Output

```
[[ 3.57142857  0.57142857  2.14285714]
 [-4.57142857 -0.07142857 -2.64285714]
 [ 2.85714286 -0.14285714  1.71428571]]
```

Here, it returns an array with solution matrix having double (float64) type values.

For example,

```
import numpy as np
from frovedis.linalg import solve

# inputs have float (float32) type values
A = np.array([[3,4,3],[1,2,3],[4,2,1]], np.float32)
B = np.array([[1,1,1], [3,0,2], [8,2,5]], np.float32)

#solve() demo with numpy array input
X = solve(A,B)
print(X)
```

Output

```
[[ 3.5714288  0.57142854  2.142857  ]
 [-4.571429  -0.07142858 -2.6428573 ]
 [ 2.857143  -0.14285713  1.7142859 ]]
```

Here, it returns an array with solution matrix having float (float32) type values.

For example,

```
import numpy as np
from frovedis.linalg import solve

# inputs have integer type values
A = np.array([[3.,4.,3.],[1.,2.,3.],[4.,2.,1.]])
B = np.array([[1.,1.,1.], [3.,0.,2.], [8.,2.,5.]])

#solve() demo with numpy array input
X = solve(A,B)
print(X)
```

Output

```
[[ 3.57142857  0.57142857  2.14285714]
 [-4.57142857 -0.07142857 -2.64285714]
 [ 2.85714286 -0.14285714  1.71428571]]
```

Here, it returns an array with solution matrix having double (float64) type values.

For example,

```
import numpy as np
from frovedis.linalg import solve
A = np.array([[3,4,3],[1,2,3],[4,2,1]])
B = np.array([[1,1,1], [3,0,2], [8,2,5]])

#creating FrovedisBlockCyclicMatrix instance
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
A_Bmat = FrovedisBlockcyclicMatrix(A, np.float64)
B_Bmat = FrovedisBlockcyclicMatrix(B, np.float64)

#solve() demo with FrovedisBlockCyclicMatrix input
X = solve(A_Bmat,B_Bmat)
X.debug_print()
```

Output

```
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 3.57143 -4.57143 2.85714 0.571429 -0.0714286 -0.142857 2.14286 -2.64286 1.71429
```

It stores result as a FrovedisBlockCyclicMatrix instance.

Also, this output will be only visible on server side terminal.

Return Value

1. When python array-like input is used:

- It returns a numpy ndarray having float (float32) or double (float64) type values and shape (**nRows**, **nCols**), where **nRows** is number of rows and **nCols** is number of columns of resultant array.

2. When input is numpy matrix:

- It returns a numpy matrix having float (float32) or double (float64) type values and shape (**nRows**, **nCols**), where **nRows** is number of rows and **nCols** is number of columns of resultant matrix.

3. When input is a FrovedisBlockCyclicMatrix instance:

- It returns an instance of FrovedisBlockCyclicMatrix.

37.4.1.6 6. `svd(a, full_matrices = False, compute_uv = True)`**Parameters**

a: It accepts python array-like inputs or numpy matrix (having dimensions ≤ 2) containing int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

full_matrices: It is a boolean parameter. Currently, it can only be set as False. Also, the unitary 2D arrays ‘u’ and ‘vh’ shapes are (M, K) and (K, N), respectively, where $K = \min(M, N)$, M is the number of rows and N is the number of columns of input matrix. (Default: False)

compute_uv: It is a boolean parameter that specifies whether or not to compute unitary arrays ‘u’ and ‘vh’ in addition to 1D array of singular values ‘s’. (Default: True)

Purpose

It computes the factor of an array by Singular Value Decomposition.

In a nutshell, this method performs factorization of matrix into three matrices ‘u’, ‘vh’ and ‘s’.

Here,

- **u** and **vh** are orthogonal matrices.
- **s** is a diagonal matrix of singular values.

This method internally **gesvd()** method present in **frovedis.linalg.scalapack** module.

For example,

```
import numpy as np
from frovedis.linalg import svd

# input is having integer values
mat1 = np.array([[3,4,3],[1,2,3],[4,2,1]])

# svd() demo with numpy 2D array
u, s, vh = svd(mat1)
print(u)
print(s)
print(vh)
```

Output

```
[[-0.73553325 -0.18392937 -0.65204358]
 [-0.42657919 -0.62196982  0.65664582]
 [-0.52632788  0.76113306  0.37901904]]

[7.87764972  2.54031671  0.69958986]

[[-0.60151068  0.73643349  0.30959751]
 [-0.61540527 -0.18005275 -0.76737042]
 [-0.5093734  -0.65210944  0.5615087  ]]
```

Here it returns ‘u’, ‘s’ and ‘vh’ as arrays having double (float64) dtype values.

For example,

```
import numpy as np
from frovedis.linalg import svd

# input is having float (float32) type values
mat1 = np.array([[3.,4.,3.],[1.,2.,3.],[4.,2.,1.]], dtype = np.float32)

# svd() demo with numpy 2D array
```



```
u, s, vh = svd(mat1)
print(u)
print(s)
print(vh)
```

Output

```
[[-0.7355336 -0.18392955 -0.65204364]
 [-0.4265793 -0.62197      0.656646  ]
 [-0.5263281  0.7611333  0.3790189 ]]
[7.8776484 2.540317  0.69959  ]
[[-0.60151064  0.73643357  0.30959752]
 [-0.6154053  -0.1800527  -0.76737046]
 [-0.5093734  -0.6521094   0.5615088  ]]
```

Here it returns ‘u’, ‘s’ and ‘vh’ as arrays having float (float32) dtype values.

For example,

```
import numpy as np
from frovedis.linalg import svd

# input is having double (float64) type values
mat1 = np.array([[3.,4.,3.],[1.,2.,3.],[4.,2.,1.]])

# svd() demo with numpy 2D array
u, s, vh = svd(mat1)
print(u)
print(s)
print(vh)
```

Output

```
[[-0.73553325 -0.18392937 -0.65204358]
 [-0.42657919 -0.62196982  0.65664582]
 [-0.52632788  0.76113306  0.37901904]]

[7.87764972 2.54031671 0.69958986]

[[-0.60151068  0.73643349  0.30959751]
 [-0.61540527 -0.18005275 -0.76737042]
 [-0.5093734  -0.65210944  0.5615087  ]]
```

Here it returns ‘u’, ‘s’ and ‘vh’ as arrays having double (float64) dtype values.

For example,

```
import numpy as np
from frovedis.linalg import svd
mat1 = np.array([[3,4,3],[1,2,3],[4,2,1]])

# svd() demo with compute_uv = False
s = svd(mat1, compute_uv = False)
print(s)
```

Output

```
[7.87764972 2.54031671 0.69958986]
```

Here, it returns only ‘s’ as an array having double (float64) type values.

For example,

```
import numpy as np
from frovedis.linalg import svd
mat1 = np.array([[3,4,3],[1,2,3],[4,2,1]])

# create FrovedisBlockcyclicMatrix instance
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat2 = FrovedisBlockcyclicMatrix(mat1, np.float64)

# svd() demo with FrovedisBlockcyclicMatrix object as input
u, s, vh = svd(mat2)
u.debug_print()
s.debug_print()
vh.debug_print()
```

Output

```
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -0.735533 -0.426579 -0.526328 -0.183929 -0.62197 0.761133 -0.652044 0.656646 0.379019
```

```
vector:
[7.87764972 2.54031671 0.69958986]
```

```
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -0.601511 -0.615405 -0.509373 0.736433 -0.180053 -0.652109 0.309598 -0.76737 0.561509
```

This output will be only visible on server side terminal.

For example,

```
import numpy as np
from frovedis.linalg import svd
mat1 = np.array([[3,4,3],[1,2,3],[4,2,1]])

# create FrovedisBlockcyclicMatrix instance
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat2 = FrovedisBlockcyclicMatrix(mat1, np.float64)

# svd() demo with FrovedisBlockcyclicMatrix object as input and compute_cv = False
s = svd(mat2, compute_uv = False)
s.debug_print()
```

Output

```
vector:
[7.87764972 2.54031671 0.69958986]
```

This output will be only visible on server side terminal.

Return Value

1. When input is numpy array:

- If `compute_cv = True`, then it returns a tuple (**u**, **s**, **vh**), where **u**, **s** and **vh** are numpy ndarrays having

float (float32) or double (float64) type values.

- If `compute_cv = False`, then it returns only array of singular values **s** of float (float32) or double (float64) type.

2. **When input is numpy matrix:**

- If `compute_cv = True`, then it returns a tuple (**u**, **s**, **vh**), where **u**, **vh** are matrices and **s** is a numpy ndarray having float (float32) or double (float64) type values.

- If `compute_cv = False`, then it returns only array of singular values **s** of float (float32) or double (float64) type.

3. **When input is a FrovedisBlockcyclicMatrix instance:**

- If `compute_cv = True`, then it returns a tuple (**u**, **s**, **vh**), where **u**, **vh** are FrovedisBlockcyclicMatrix instances and **s** is FrovedisVector instance.

- If `compute_cv = False`, then it returns **s** is an instance of FrovedisVector.

37.5 SEE ALSO

- Scalapack Functions

Chapter 38

scalapack

38.1 NAME

scalapack - it contains wrapper functions created for low level functions from ScaLAPACK library (present in frovedis).

38.2 SYNOPSIS

```
import frovedis.linalg.scalapack
```

38.3 Public Member Functions

1. dgels(a, b, trans='N', lwork=0, overwrite_a=0, overwrite_b=0)
2. gels(a, b, trans='N', lwork=0, overwrite_a=0, overwrite_b=0, dtype=np.float64)
3. sgels(a, b, trans='N', lwork=0, overwrite_a=0, overwrite_b=0)
4. dgesv(a, b, overwrite_a=0, overwrite_b=0)
5. gesv(a, b, overwrite_a=0, overwrite_b=0, dtype=np.float64)
6. sgesv(a, b, overwrite_a=0, overwrite_b=0)
7. dgesvd(a, compute_uv=1, full_matrices=0, lwork=0, overwrite_a=0)
8. gesvd(a, compute_uv=1, full_matrices=0, lwork=0, overwrite_a=0, dtype=np.float64)
9. sgesvd(a, compute_uv=1, full_matrices=0, lwork=0, overwrite_a=0)
10. dgetrf(a, overwrite_a=0)
11. getrf(a, overwrite_a=0, dtype=np.float64)
12. sgetrf(a, overwrite_a=0)
13. dgetri(lu, piv, lwork=0, overwrite_lu=0)
14. getri(lu, piv, lwork=0, overwrite_lu=0, dtype=np.float64)
15. sgetri(lu, piv, lwork=0, overwrite_lu=0)
16. dgetrs(lu, piv, b, trans=0, overwrite_b=0)
17. getrs(lu, piv, b, trans=0, overwrite_b=0, dtype=np.float64)
18. sgetrs(lu, piv, b, trans=0, overwrite_b=0)

38.4 DESCRIPTION

In Frovedis, the linalg module contains wrapper for ScaLAPACK (Scalable LAPACK: the LAPACK routines distributed in nature) routines, whereas scipy.linalg contains wrapper for LAPACK routines. Here, the method names, arguments, purpose etc. in frovedis.linalg wrapper routines are similar to scipy.linalg wrapper routines.

In order to use dgetrf() wrapper present in linalg module of both frovedis and scipy:

In frovedis:

```
from frovedis.linalg.scalapack import dgetrf
```

In scipy:

```
from scipy.linalg.lapack import dgetrf
```

This python module implements a client-server application, where the python client can send the python matrix data to frovedis server side in order to create blockcyclic matrix at frovedis server and then perform the supported ScaLAPACK operation requested by the python client on that matrix. After request is completed, frovedis server sends back the resultant matrix and it can then create equivalent python data.

We have supported ‘overwrite’ option in frovedis routine wrappers.

However, unlike scipy.linalg, if ‘overwrite’ option is enabled and input is a valid ndarray or an instance of FrovedisBlockcyclicMatrix, we would always overwrite (copy-back) to the same (irrespective of its dtype).

This will slow down the overall computation. This should be enabled only when it’s wanted to check the intermediate results (like LU factor etc.)

38.4.1 Detailed Description

38.4.1.1 1. dgels(a, b, trans = ‘N’, lwork = 0, overwrite_a = 0, overwrite_b = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values.

b: It accepts a python array-like input or right hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values. It should have number of rows $\geq \max(M, N)$ and at least 1 column.

trans: It accepts a string object parameter like ‘N’ or ‘T’ which specifies if transpose of ‘a’ is needed to be computed before solving linear equation. If set to ‘T’, then transpose is computed. (Default: ‘N’)

lwork: This is an unused parameter. (Default: 0)

overwrite_a: It accepts an integer parameter, if set to 0, then ‘a’ will remain unchanged. Otherwise, ‘a’ would be overwritten with QR or LQ factor. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then ‘b’ will remain unchanged. Otherwise, ‘b’ would be overwritten with the solution matrix. (Default: 0)

Purpose

This method solves overdetermined or underdetermined real linear systems involving a left hand side matrix ‘a’ or its transpose, using a QR or LQ factorization of ‘a’. It is assumed that the matrix ‘a’ has full rank. It returns the LQ or QR factor, solution matrix with double (float64) precision.

The parameter: “lwork” is simply kept in to to make the interface uniform to the scipy.linalg.lapack.dgels() module. They are not used anywhere within the frovedis implementation.

This method internally uses Scalapack.gels() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
```

```
# dgels() demo
from frovedis.linalg.scalapack import dgels
rf = dgels(mat1,mat2)
print(rf)
```

Output

```
(array([[-1.41421356,  3.53553391, -1.41421356],
        [-0.41421356, -4.63680925,  4.74464202],
        [ 0.          ,  0.36709178, -7.77742709]]), array([[-7.54901961,  2.68627451, -1.          ],
        [-1.50980392,  1.1372549 ,  0.          ],
        [-0.7254902 ,  0.15686275,  0.          ]]), 0)
```

For example,

```
# dgels() demo and using transpose of a
from frovedis.linalg.scalapack import dgels
rf = dgels(mat1,mat2,trans='T')
print(rf)
```

Output

```
(array([[-1.41421356,  3.53553391, -1.41421356],
        [-0.41421356, -4.63680925,  4.74464202],
        [ 0.          ,  0.36709178, -7.77742709]]), array([[-7.82352941,  3.29411765, -0.70588235],
        [ 1.17647059,  0.29411765,  0.29411765],
        [-1.96078431,  0.50980392, -0.15686275]]), 0)
```

For example,

```
# dgels() demo and overwriting of a with QR or LQ factor
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import dgels
rf = dgels(mat1,mat2,overwrite_a=1)
print('overwritten matrix with LQ or QF factor: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

```
overwritten matrix with LQ or QF factor:
[[-1.41421356  3.53553391 -1.41421356]
 [-0.41421356 -4.63680925  4.74464202]
 [ 0.          0.36709178 -7.77742709]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, then LQ or QR factor would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# dgels() demo and overwriting of b with solution matrix
print('original matrix mat2: ')
print(mat2)
from frovedis.linalg.scalapack import dgels
rf = dgels(mat1,mat2,overwrite_b=1)
print('overwritten matrix with solution matrix: ')
print(mat2)
```

Output

```
original matrix mat2:
[[-9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]

overwritten matrix with solution matrix:
[[-7.54901961  2.68627451 -1.         ]
 [-1.50980392  1.1372549   0.         ]
 [-0.7254902   0.15686275  0.         ]]
```

Here, if the input ‘mat2’ is double (float64) type and overwrite is enabled, then solution matrix would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# dgels() demo and a and b as an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgels
rf = dgels(bcm1,bcm2)

# Unpacking the tuple
rf[0].debug_print()
rf[1].debug_print()
print(rf[2])
```

Output

```
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -1.41421 -0.414214 0 3.53553 -4.63681 0.367092 -1.41421 4.74464 -7.77743
matrix:
num_row = 3, num_col = 3
node 0
```



```
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0
```

Return Value

1. If ‘a’ and ‘b’ are python inputs such as numpy matrices:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It is a numpy matrix having double (float64) type values (by default) and containing the QR or LQ factor of input matrix ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **x**: It is also a numpy matrix having double (float64) type values (by default) and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If ‘a’ and ‘b’ are instances of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It returns instance of FrovedisBlockcyclicMatrix containing the QR or LQ factor of input matrix ‘a’.
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

Here, the original inputs ‘a’ and ‘b’ are not overwritten even when overwrite_a/overwrite_b is enabled.

38.4.1.2 2. gels(a, b, trans = ‘N’, lwork = 0, overwrite_a = 0, overwrite_b = 0, dtype = np.float64)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

b: It accepts a python array-like input or right hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It should have number of rows $\geq \max(M, N)$ and at least 1 column.

trans: It accepts a string object parameter like ‘N’ or ‘T’ which specifies if transpose of ‘a’ is needed to be computed before solving linear equation. If set to ‘T’, then transpose is computed. (Default: ‘N’)

lwork: This is an unused parameter. (Default: 0)

overwrite_a: It accepts an integer parameter, if set to 0, then ‘a’ will remain unchanged. Otherwise, ‘a’ would be overwritten with QR or LQ factor. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then ‘b’ will remain unchanged. Otherwise, ‘b’ would be overwritten with the solution matrix. (Default: 0)

dtype: It specifies the datatype to be used for setting the precision level (single for float32 / double for float64) for the values returned by this method. (Default: np.float64)

Currently, it supports float (float32) or double (float64) datatypes.

Purpose

This method solves overdetermined or underdetermined real linear systems involving a left hand side matrix ‘a’ or its transpose, using a QR or LQ factorization of ‘a’. It is assumed that the matrix ‘a’ has full rank. It returns the LQ or QR factor, solution matrix for the system of linear equations with single (float32) or double (float64) precision depending on the ‘dtype’ parameter provided by user.

The parameter: “lwork” is simply kept in to to make the interface uniform with other modules in frovedis. They are not used anywhere within the frovedis implementation.

This method is present only in frovedis.

This method internally uses Scalapack.gels() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
```

```
mat2 = np.array([[ -9., 3., -1.], [0., 3., 1.], [2., 2., 0.]])
```

```
# gels() demo
from frovedis.linalg.scalapack import gels
rf = gels(mat1, mat2)
print(rf)
```

Output

```
(array([[ -1.41421356,  3.53553391, -1.41421356],
        [ -0.41421356, -4.63680925,  4.74464202],
        [ 0.          ,  0.36709178, -7.77742709]]), array([[ -7.54901961,  2.68627451, -1.          ],
        [ -1.50980392,  1.1372549 ,  0.          ],
        [ -0.7254902 ,  0.15686275,  0.          ]]), 0)
```

For example,

```
# gels() demo and using transpose of a
from frovedis.linalg.scalapack import gels
rf = gels(mat1, mat2, trans='T')
print(rf)
```

Output

```
(array([[ -1.41421356,  3.53553391, -1.41421356],
        [ -0.41421356, -4.63680925,  4.74464202],
        [ 0.          ,  0.36709178, -7.77742709]]), array([[ -7.82352941,  3.29411765, -0.70588235],
        [ 1.17647059,  0.29411765,  0.29411765],
        [ -1.96078431,  0.50980392, -0.15686275]]), 0)
```

For example,

```
# gels() demo and overwriting of a with QR or LQ factor
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import gels
rf = gels(mat1, mat2, overwrite_a=1)
print('overwritten matrix with LQ or QF factor: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

```
overwritten matrix with LQ or QF factor:
[[ -1.41421356  3.53553391 -1.41421356]
 [ -0.41421356 -4.63680925  4.74464202]
 [ 0.          0.36709178 -7.77742709]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, then LQ or QR factor would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# gels() demo and overwriting of b with solution matrix
```

```

print('original matrix mat2: ')
print(mat2)
from frovedis.linalg.scalapack import gels
rf = gels(mat1,mat2,overwrite_b=1)
print('overwritten matrix with solution matrix: ')
print(mat2)

```

Output

```

original matrix mat2:
[[-9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]

overwritten matrix with solution matrix:
[[-7.54901961  2.68627451 -1.         ]
 [-1.50980392  1.1372549   0.         ]
 [-0.7254902   0.15686275  0.         ]]

```

Here, if the input ‘mat2’ is double (float64) type and overwrite is enabled, then solution matrix would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```

# gels() demo and specifying the dtypes of output matrices LQ or QR factor and solution matrix
from frovedis.linalg.scalapack import gels
rf = gels(mat1,mat2,dtype=np.float32)
print(rf)

```

Output

```

(array([[ -1.4142135 ,  3.535534 , -1.4142137 ],
        [-0.41421354, -4.6368093 ,  4.744642  ],
        [ 0.         ,  0.36709177, -7.777427  ]], dtype=float32),
 array([[ -7.5490184 ,  2.686274 , -1.0000001 ],
        [-1.5098035 ,  1.1372546 ,  0.         ],
        [-0.7254901 ,  0.15686269,  0.         ]], dtype=float32), 0)

```

For example,

```

from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# gels() demo and a and b as an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import gels
rf = gels(bcm1,bcm2)

# Unpacking the tuple
rf[0].debug_print()
rf[1].debug_print()
print(rf[2])

```

Output

```

0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -1.41421 -0.414214 0 3.53553 -4.63681 0.367092 -1.41421 4.74464 -7.77743
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0

```

Note:- ‘dtype’ for the wrapper function and FrovedisBlockcyclicMatrix instance must be same during computation. Otherwise, it will raise an excpetion.

Return Value

1. If ‘a’ and ‘b’ are python inputs such as numpy matrices and dtype = np.float64:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It is a numpy matrix having double (float64) type values (by default) and containing the QR or LQ factor of input matrix ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **x**: It is also a numpy matrix having double (float64) type values (by default) and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If ‘a’ and ‘b’ are python inputs such as numpy matrices and dtype = np.float32:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It is a numpy matrix having float (float32) type values (by default) and containing the QR or LQ factor of input matrix ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **x**: It is also a numpy matrix having float (float32) type values (by default) and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

3. If ‘a’ and ‘b’ are instances of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It returns instance of FrovedisBlockcyclicMatrix containing the QR or LQ factor of input matrix ‘a’.
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.3 3. sgels(a, b, trans = ‘N’, lwork = 0, overwrite_a = 0, overwrite_b = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values.

b: It accepts a python array-like input or right hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values. It should have number of rows $\geq \max(M, N)$ and at least 1 column.

trans: It accepts a string object parameter like ‘N’ or ‘T’ which specifies if transpose of ‘a’ is needed to be computed before solving linear equation. If set to ‘T’, then transpose is computed. (Default: ‘N’)

lwork: This is an unused parameter. (Default: 0)

overwrite_a: It accepts an integer parameter, if set to 0, then ‘a’ will remain unchanged. Otherwise, ‘a’ would be overwritten with QR or LQ factor. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then ‘b’ will remain unchanged. Otherwise, ‘b’ would be overwritten with the solution matrix. (Default: 0)

Purpose

This method solves overdetermined or underdetermined linear systems involving a left hand side matrix ‘a’ or its transpose, using a QR or LQ factorization of ‘a’. It is assumed that the matrix ‘a’ has full rank. It returns the LQ or QR factor, solution matrix for the system of linear equations with single (float32) precision.

The parameter: “lwork” is simply kept in to to make the interface uniform to the `scipy.linalg.lapack.sgels()` module. They are not used anywhere within the `frovedis` implementation.

This method internally uses `Scalapack.gels()` (**present in `frovedis.matrix` module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
```

```
# sgels() demo
from frovedis.linalg.scalapack import sgels
rf = sgels(mat1,mat2)
print(rf)
```

Output

```
(array([[-1.4142135 ,  3.535534 , -1.4142137 ],
        [-0.41421354, -4.6368093 ,  4.744642  ],
        [ 0.          ,  0.36709177, -7.777427  ]], dtype=float32),
 array([[-7.5490184 ,  2.686274 , -1.0000001 ],
        [-1.5098035 ,  1.1372546 ,  0.          ],
        [-0.7254901 ,  0.15686269,  0.          ]], dtype=float32), 0)
```

For example,

```
# sgels() demo and using transpose of a
from frovedis.linalg.scalapack import sgels
rf = sgels(mat1,mat2,trans='T')
print(rf)
```

Output

```
(array([[-1.4142135 ,  3.535534 , -1.4142137 ],
        [-0.41421354, -4.6368093 ,  4.744642  ],
        [ 0.          ,  0.36709177, -7.777427  ]], dtype=float32),
 array([[-7.8235283 ,  3.2941177 , -0.7058824 ],
        [ 1.1764708 ,  0.29411745,  0.29411763],
        [-1.9607842 ,  0.509804 , -0.15686275]], dtype=float32), 0)
```

For example,

```
# sgels() demo and overwriting of a with QR or LQ factor
print('original matrix mat1: ', mat1)
from frovedis.linalg.scalapack import sgels
rf = sgels(mat1,mat2,overwrite_a=1)
print('overwritten matrix with LQ or QF factor: ', mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
overwritten matrix with LQ or QF factor:
[[-1.41421354  3.53553391 -1.41421366]
```

```
[-0.41421354 -4.63680935  4.74464178]
[ 0.          0.36709177 -7.7774272  ]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, then LQ or QR factor would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# sgels() demo and overwriting of b with solution matrix
print('original matrix mat2: ')
print(mat2)
from frovedis.linalg.scalapack import sgels
rf = sgels(mat1,mat2,overwrite_b=1)
print('overwritten matrix with solution matrix: ')
print(mat2)
```

Output

```
original matrix mat2:
[[-9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]

overwritten matrix with solution matrix:
[[-7.54901838  2.68627405 -1.00000012]
 [-1.50980353  1.1372546  0.          ]
 [-0.72549009  0.15686269  0.          ]]
```

Here, if the input ‘mat2’ is double (float64) type and overwrite is enabled, then solution matrix would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# sgels() demo and a and b as an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import sgels
rf = sgels(bcm1,bcm2)

# Unpacking the tuple
rf[0].debug_print()
rf[1].debug_print()
print(rf[2])
```

Output

```
0
matrix:
num_row = 3, num_col = 3
node 0
```

```

node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -1.41421 -0.414214 0 3.53553 -4.63681 0.367092 -1.41421 4.74464 -7.77743
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0

```

Return Value

1. If 'a' and 'b' are python inputs such as numpy matrices:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It is a numpy matrix having float (float32) type values (by default) and containing the QR or LQ factor of input matrix 'a'. In case 'overwrite_a' is enabled, then dtype for the matrix will depend on input 'a' dtype.
- **x**: It is also a numpy matrix having float (float32) (by default) type values and containing the solution matrix. In case 'overwrite_b' is enabled then dtype for the matrix will depend on input 'b' dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If 'a' and 'b' are instances of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lqr**, **x**, **stat**) where,
- **lqr**: It returns an instance of FrovedisBlockcyclicMatrix containing the QR or LQ factor of input matrix 'a'.
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.4 4. dgesv(a, b, overwrite_a = 0, overwrite_b = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values.

b: It accepts a python array-like input or right hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values. It should have number of rows \geq the number of rows in 'a' and at least 1 column in it.

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise, 'a' would be overwritten with LU factor. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then 'b' will remain unchanged. Otherwise, 'b' would be overwritten with the solution matrix. (Default: 0)

Purpose

It solves a system of linear equations, $AX = B$ with a left hand side square matrix, 'a' by computing its LU factors internally. It returns the LU factor, solution matrix for the system of linear equations with double (float64) precision.

This method internally uses Scalapack.dgesv() (**present in frovedis.matrix module**).

For example,

```

import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])

# dgesv() demo
from frovedis.linalg.scalapack import dgesv
rf = dgesv(mat1,mat2)
print(rf)

```

Output

```
(array([[ 1. ,  0. ,  2. ],
       [-1. ,  5. ,  2. ],
       [ 0. ,  0.6, -10.2]]), <frovedis.matrix.results.GetrfResult object at 0x7f30f7e25208>,
array([[-7.54901961,  2.68627451, -1.          ],
       [-1.50980392,  1.1372549 ,  0.          ],
       [-0.7254902 ,  0.15686275,  0.          ]]), 0)
```

This resultant tuple of `dgesv()` wrapper function contains `GetrfResult` instance in `frovedis`. Whereas in `scipy`, it contains an array.

Both here are responsible to hold pivot array information.

For example,

```
# dgesv() demo and overwriting of a with LU factor
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import dgesv
rf = dgesv(mat1,mat2,overwrite_a=1)
print('overwritten matrix with LU factor: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

```
overwritten matrix with LU factor:
[[ 1.    0.    2. ]
 [-1.    5.    2. ]
 [ 0.    0.6 -10.2]]
```

Here, if the input ‘`mat1`’ is double (`float64`) type and overwrite is enabled, then LU factor would also be double (`float64`) type.

Same applies for other types (`int`, `float (float32)`) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# dgesv() demo and overwriting of b with solution matrix
print('original matrix mat2: ')
print(mat2)
from frovedis.linalg.scalapack import dgesv
rf = dgesv(mat1,mat2,overwrite_b=1)
print('overwritten matrix with solution matrix: ')
print(mat2)
```

Output

```
original matrix mat2:
[[-9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]
```

```
overwritten matrix with solution matrix:
[[-7.54901961  2.68627451 -1.          ]
 [-1.50980392  1.1372549   0.          ]
```



```
[-0.7254902  0.15686275  0.      ]]
```

Here, if the input 'mat2' is double (float64) type and overwrite is enabled, then solution matrix would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[-9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# dgesv() demo and a and b as an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgesv
rf = dgesv(bcm1,bcm2)

# Unpacking the tuple
rf[0].debug_print()
print(rf[1])
rf[2].debug_print()
print(rf[3])
```

Output

```
<frovedis.matrix.results.GetrfResult object at 0x7f7ce5798cf8>
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 1 -1 0 0 5 0.6 2 2 -10.2
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0
```

Return Value

1. If 'a' and 'b' are python inputs such as numpy matrices:

- It returns a tuple (**lu**, **piv**, **x**, **rs_stat**) where,
- **lu**: It is a numpy matrix having double (float64) type values (by default) and containing the LU factor of input 'a'. In case 'overwrite_a' is enabled, then dtype for the matrix will depend on input 'a' dtype.
- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It is also a numpy matrix having double (float64) type values (by default) and containing the solution matrix. In case 'overwrite_b' is enabled, then dtype for the matrix will depend on input 'b' dtype.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If 'a' and 'b' are instances of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lu**, **piv**, **x**, **rs_stat**) where,
- **lu**: It returns an instance of FrovedisBlockcyclicMatrix containing the LU factor of input 'a'.
- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.5 5. gesv(a, b, overwrite_a = 0, overwrite_b = 0, dtype = np.float64)**Parameters**

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

b: It accepts a python array-like input or right hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It should have number of rows \geq the number of rows in 'a' and at least 1 column in it.

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise, 'a' would be overwritten with LU factor. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then 'b' will remain unchanged. Otherwise, 'b' would be overwritten with the solution matrix. (Default: 0)

dtype: It specifies the datatype to be used for setting the precision level (single for float32 / double for float64) for the values returned by this method. (Default: np.float64)

Currently, it supports float (float32) or double (float64) datatypes.

Purpose

It solves a system of linear equations, $AX = B$ with a left hand side square matrix, 'a' by computing it's LU factors internally. It returns the LU factor computed using getrf() and solution matrix computed using getrf() for the system of linear equations with float (float32) or double (float64) precision depending on the 'dtype' parameter provided by user.

This method is present only in frovedis.

This method internally uses Scalapack.gesv() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[-9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])

# gesv() demo
from frovedis.linalg.scalapack import gesv
rf = gesv(mat1,mat2)
print(rf)
```

Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]]), <frovedis.matrix.results.GetrfResult object at 0x7f30f7e25208>,
array([[-7.54901961,  2.68627451, -1.          ],
        [-1.50980392,  1.1372549 ,  0.          ],
        [-0.7254902 ,  0.15686275,  0.          ]]), 0)
```

For example,

```
# dgesv() demo and overwriting of a with LU factor
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import dgesv
rf = dgesv(mat1,mat2,overwrite_a=1)
print('overwritten matrix with LU factor: ')
print(mat1)
```

Output

original matrix mat1:

```
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

overwritten matrix with LU factor:

```
[[ 1.    0.    2. ]
 [-1.    5.    2. ]
 [ 0.    0.6 -10.2]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, then LU factor would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# dgesv() demo and overwriting of b with solution matrix
print('original matrix mat2: ')
print(mat2)
from frovedis.linalg.scalapack import dgesv
rf = dgesv(mat1,mat2,overwrite_b=1)
print('overwritten matrix with solution matrix: ')
print(mat2)
```

Output

original matrix mat2:

```
[[ -9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]
```

overwritten matrix with solution matrix:

```
[[ -7.54901961  2.68627451 -1.          ]
 [-1.50980392  1.1372549   0.          ]
 [-0.7254902   0.15686275  0.          ]]
```

Here, if the input ‘mat2’ is double (float64) type and overwrite is enabled, then solution matrix would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# gesv() demo and specifying the dtypes of output matrices LU factor and solution matrix
from frovedis.linalg.scalapack import gesv
rf = gesv(mat1,mat2, dtype=np.float32)
print(rf)
```

Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]], dtype=float32),
 <frovedis.matrix.results.GetrfResult object at 0x7f03d450ad30>,
 array([[ -7.54902 ,  2.6862745 , -1.          ],
        [-1.509804 ,  1.137255 ,  0.          ],
        [-0.7254902 ,  0.15686277,  0.          ]], dtype=float32), 0)
```

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# dgesv() demo and a and b as an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgesv
rf = dgesv(bcm1,bcm2)

# Unpacking the tuple
rf[0].debug_print()
print(rf[1])
rf[2].debug_print()
print(rf[3])
```

Output

```
<frovedis.matrix.results.GetrfResult object at 0x7f7ce5798cf8>
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 1 -1 0 0 5 0.6 2 2 -10.2
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0
```

Note:- ‘dtype’ for the wrapper function and FrovedisBlockcyclicMatrix instance must be same during computation. Otherwise, it will raise an exception.

Return Value

1. If ‘a’ and ‘b’ are python inputs such as numpy matrices and dtype = np.float32:

- It returns a tuple (lu, piv, x, rs_stat) where,
- **lu**: It is a numpy matrix having float (float32) type values (by default) and containing the LU factor of input ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It is also a numpy matrix having float (float32) type values (by default) and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If ‘a’ and ‘b’ are python inputs such as numpy matrices and dtype = np.float64:

- It returns a tuple (lu, piv, x, rs_stat) where,
- **lu**: It is a numpy matrix having double (float64) type values (by default) and containing the LU factor of input ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It is also a numpy matrix having double (float64) type values (by default) and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

3. If ‘a’ and ‘b’ are instances of FrovedisBlockcyclicMatrix:

- It returns a tuple (lu, piv, x, rs_stat) where,
- **lu**: It returns an instance of FrovedisBlockcyclicMatrix containing the LU factor of input ‘a’.

- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.6 6. sgesv(a, b, overwrite_a = 0, overwrite_b = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values.

b: It accepts a python array-like input or right hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values. It should have number of rows \geq the number of rows in 'a' and at least 1 column in it.

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise, 'a' would be overwritten with LU factor. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then 'b' will remain unchanged. Otherwise, 'b' would be overwritten with the solution matrix. (Default: 0)

Purpose

It solves a system of linear equations, $AX = B$ with a left hand side square matrix, 'a' by computing it's LU factors internally. It returns the LU factor, solution matrix for the system of linear equations with single (float32) precision.

This method internally uses Scalapack.sgesv() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]],dtype = np.float32)
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]],dtype = np.float32)
```

```
# sgesv() demo
from frovedis.linalg.scalapack import sgesv
rf = sgesv(mat1,mat2)
print(rf)
```

Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]], dtype=float32),
 <frovedis.matrix.results.GetrfResult object at 0x7f49657f3cf8>,
 array([[ -7.54902   ,  2.6862745 , -1.          ],
        [-1.509804   ,  1.137255   ,  0.          ],
        [-0.7254902   ,  0.15686277,  0.          ]], dtype=float32), 0)
```

This resultant tuple of sgesv() wrapper function contains GetrfResult instance in frovedis. Whereas in scipy, it contains an array.

Both here are responsible to hold pivot array information.

For example,

```
# sgesv() demo and overwriting of a with LU factor
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import sgesv
rf = sgesv(mat1,mat2,overwrite_a=1)
print('overwritten matrix with LU factor: ')
```

```
print(mat1)
```

Output

original matrix mat1:

```
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

overwritten matrix with LU factor:

```
[[ 1.    0.    2. ]
 [-1.    5.    2. ]
 [ 0.    0.6 -10.2]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, then LU factor would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# sgesv() demo and overwriting of b with solution matrix
print('original matrix mat2: ')
print(mat2)
from frovedis.linalg.scalapack import sgesv
rf = sgesv(mat1,mat2,overwrite_b=1)
print('overwritten matrix with solution matrix: ')
print(mat2)
```

Output

original matrix mat2:

```
[[ -9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]
```

overwritten matrix with solution matrix:

```
[[ -7.54901981  2.68627453 -1.         ]
 [-1.50980401  1.13725495  0.         ]
 [-0.72549021  0.15686277  0.         ]]
```

Here, if the input ‘mat2’ is double (float64) type and overwrite is enabled, then solution matrix would also be double (float64) type.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype=np.float32)
bcm2 = FrovedisBlockcyclicMatrix(mat2,dtype=np.float32)

# sgesv() demo and a and b as an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import sgesv
rf = sgesv(bcm1,bcm2)
```

```
# Unpacking the tuple
rf[0].debug_print()
print(rf[1])
rf[2].debug_print()
print(rf[3])
```

Output

```
<frovedis.matrix.results.GetrfResult object at 0x7f67ddeccd30>
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 1 -1 0 0 5 0.6 2 2 -10.2
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0
```

Return Value

1. If 'a' and 'b' are python inputs such as numpy matrices:

- It returns a tuple (**lu**, **piv**, **x**, **rs_stat**) where,
- **lu**: It is a numpy matrix having float (float32) type values (by default) and containing the LU factor of input 'a'. In case 'overwrite_a' is enabled, then dtype for the matrix will depend on input 'a' dtype.
- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It is also a numpy matrix having float (float32) type values (by default) and containing the solution matrix. In case 'overwrite_b' is enabled, then dtype for the matrix will depend on input 'b' dtype.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If 'a' and 'b' are instances of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lu**, **piv**, **x**, **rs_stat**) where,
- **lu**: It returns an instance of FrovedisBlockcyclicMatrix containing the LU factor of input 'a'.
- **piv**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **rs_stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.7 7. dgesvd(a, compute_uv = 1, full_matrices = 0, lwork = 0, overwrite_a = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values.

compute_uv: It accepts an integer parameter that specifies whether left singular matrix and right singular matrix will be computed. If it is set as '0', a numpy matrix of shape (1,1) and datatype same as 'a' will be assigned to the left and right singular matrix. (Default: 1)

full_matrices: It accepts an integer parameter. Currently, it can only be set as 0. Also, the left singular matrix and right singular matrix shapes are (M, K) and (K, N), respectively, where $K = \min(M, N)$, M is the number of rows and N is the number of columns of input matrix. (Default: 0)

lwork: This is an unused parameter. (Default: 0)

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise 'a' would be consumed internally and its contents will be destroyed. (Default: 0)

Purpose

It computes the singular value decomposition (SVD) of matrix 'a' with double (float64) precision.

The parameter: “lwork” is simply kept in to make the interface uniform to the `scipy.linalg.lapack.dgesvd()` module. They are not used anywhere within the `frovedis` implementation.

This method internally uses `Scalapack.gesvd()` (**present in `frovedis.matrix` module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# dgesvd() demo
from frovedis.linalg.scalapack import dgesvd
rf = dgesvd(mat1)
print(rf)
```

Output

```
(array([[ 0.19046167, -0.12914878, -0.97316234],
        [-0.20606538, -0.97448299,  0.08899413],
        [-0.95982364,  0.18358509, -0.21221475]]), array([9.83830146, 4.80016509, 1.0799257 ]),
 array([[ 0.04030442, -0.39740577,  0.91675744],
        [ 0.17610524, -0.9003148 , -0.39802035],
        [-0.98354588, -0.17748777, -0.03369857]]), 0)
```

For example,

```
# dgesvd() demo and compute_uv = 0
from frovedis.linalg.scalapack import dgesvd
rf = dgesvd(mat1,compute_uv=0)
print(rf)
```

Output

```
(array([[0.]]), array([9.83830146, 4.80016509, 1.0799257 ]), array([[0.]]), 0)
```

For example,

```
# dgesvd() demo and overwriting of a
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import dgesvd
rf = dgesvd(mat1,overwrite_a=1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]

overwritten matrix:
[[-1.41421356 -3.80788655 -0.1925824 ]
 [-0.41421356  6.71975985  5.52667534]
 [ 0.          0.64659915 -5.36662718]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)

# dgesvd() demo and a is an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgesvd
rf = dgesvd(bcm1)

# Unpacking the tuple
rf[0].debug_print()
rf[1].debug_print()
rf[2].debug_print()
```

Output

```
vector:
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.190462 -0.206065 -0.959824 -0.129149 -0.974483 0.183585 -0.973162 0.0889941 -0.212215
[9.83830146 4.80016509 1.0799257 ]
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.0403044 -0.397406 0.916757 0.176105 -0.900315 -0.39802 -0.983546 -0.177488 -0.0336986
```

Return Value

1. If 'a' is a python input such as numpy matrix:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It is a numpy matrix having double (float64) type values and containing the left singular matrix.
- **s**: It is a numpy matrix having double (float64) type values and containing the singular matrix.
- **vt**: It is a numpy matrix having double (float64) type values and containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If 'a' is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It returns an instance of FrovedisBlockcyclicMatrix containing the left singular matrix.
- **s**: It is a numpy matrix having double (float64) type values and containing the singular matrix.
- **vt**: It returns an instance of FrovedisBlockcyclicMatrix containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.8 8. gesvd(a, compute_uv = 1, full_matrices = 0, lwork = 0, overwrite_a = 0, dtype = np.float64)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

compute_uv: It accepts an integer parameter that specifies whether left singular matrix and right singular matrix will be computed. If it is set as '0', a numpy matrix of shape (1,1) and datatype same as 'a' will be

assigned to the left and right singular matrix. (Default: 1)

full_matrices: It accepts an integer parameter. Currently, it can only be set as 0. Also, the left singular matrix and right singular matrix shapes are (\mathbf{M}, \mathbf{K}) and (\mathbf{K}, \mathbf{N}) , respectively, where $\mathbf{K} = \min(\mathbf{M}, \mathbf{N})$, \mathbf{M} is the number of rows and \mathbf{N} is the number of columns of input matrix. (Default: 0)

lwork: This is an unused parameter. (Default: 0)

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise 'a' would be consumed internally and its contents will be destroyed. (Default: 0)

dtype: It specifies the datatype to be used for setting the precision level (single for float32 / double for float64) for the values returned by this method. (Default: np.float64)

Currently, it supports float (float32) or double (float64) datatypes.

Purpose

It computes the singular value decomposition (SVD) of matrix 'a' with single (float32) or double (float64) precision depending on the 'dtype' parameter provided by user.

The parameter: "lwork" is simply kept in to to make the interface uniform to the scipy.linalg.lapack.dgesvd() module. They are not used anywhere within the frovedis implementation.

This method is present only in frovedis.

This method internally uses Scalapack.gesvd() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# gesvd() demo
from frovedis.linalg.scalapack import gesvd
rf = gesvd(mat1)
print(rf)
```

Output

```
(array([[ 0.19046167, -0.12914878, -0.97316234],
        [-0.20606538, -0.97448299,  0.08899413],
        [-0.95982364,  0.18358509, -0.21221475]]),
 array([9.83830146, 4.80016509, 1.0799257 ]),
 array([[ 0.04030442, -0.39740577,  0.91675744],
        [ 0.17610524, -0.9003148 , -0.39802035],
        [-0.98354588, -0.17748777, -0.03369857]]), 0)
```

For example,

```
# gesvd() demo and compute_uv = 0
from frovedis.linalg.scalapack import gesvd
rf = gesvd(mat1,compute_uv=0)
print(rf)
```

Output

```
(array([[0.]]), array([9.83830146, 4.80016509, 1.0799257 ]), array([[0.]]), 0)
```

For example,

```
# gesvd() demo and overwriting of a
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import gesvd
rf = gesvd(mat1,overwrite_a=1)
print('overwritten matrix: ')
```

```
print(mat1)
```

Output

original matrix mat1:

```
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

overwritten matrix:

```
[[ -1.41421356 -3.80788655 -0.1925824 ]
 [ -0.41421356  6.71975985  5.52667534]
 [  0.          0.64659915 -5.36662718]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# gesvd() demo and specifying the dtypes of output matrices
from frovedis.linalg.scalapack import gesvd
rf = gesvd(mat1,dtype=np.float32)
print(rf)
```

Output

```
(array([[ 0.19046175, -0.12914878, -0.9731621 ],
        [-0.20606534, -0.97448295,  0.08899409],
        [-0.9598237 ,  0.1835851 , -0.2122148 ]], dtype=float32),
 array([9.838303 , 4.800165 , 1.0799255], dtype=float32),
 array([[ 0.04030442, -0.3974058 ,  0.91675735],
        [ 0.17610519, -0.90031475, -0.39802024],
        [-0.98354584, -0.17748773, -0.03369856]], dtype=float32), 0)
```

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)

# gesvd() demo and a is an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import gesvd
rf = gesvd(bcm1)
```

Unpacking the tuple

```
rf[0].debug_print()
rf[1].debug_print()
rf[2].debug_print()
```

Output

vector:

matrix:

num_row = 3, num_col = 3

node 0

node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3

val = 0.190462 -0.206065 -0.959824 -0.129149 -0.974483 0.183585 -0.973162 0.0889941 -0.212215

```
[9.83830146 4.80016509 1.0799257 ]
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.0403044 -0.397406 0.916757 0.176105 -0.900315 -0.39802 -0.983546 -0.177488 -0.0336986
```

Note:- ‘dtype’ for the wrapper function and FrovedisBlockcyclicMatrix instance must be same during computation. Otherwise, it will raise an excpetion.

Return Value

1. If ‘a’ is a python input such as numpy matrix and dtype = np.float32:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It is a numpy matrix having float (float32) type values and containing the left singular matrix.
- **s**: It is a numpy matrix having float (float32) type values and containing the singular matrix.
- **vt**: It is a numpy matrix having float (float32) type values and containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If ‘a’ is a python input such as numpy matrix and dtype = np.float64:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It is a numpy matrix having double (float64) type values and containing the left singular matrix.
- **s**: It is a numpy matrix having double (float64) type values and containing the singular matrix.
- **vt**: It is a numpy matrix having double (float64) type values and containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

3. If ‘a’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It returns an instance of FrovedisBlockcyclicMatrix containing the left singular matrix.
- **s**: It is a numpy matrix having double (float64) type values and containing the singular matrix.
- **vt**: It returns an instance of FrovedisBlockcyclicMatrix containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.9 9. sgesvd(a, compute_uv = 1, full_matrices = 0, lwork = 0, overwrite_a = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values.

compute_uv: It accepts an integer parameter that specifies whether left singular matrix and right singular matrix will be computed. If it is set as ‘0’, a numpy matrix of shape (1,1) and datatype same as ‘a’ will be assigned to the left and right singular matrix. (Default: 1)

full_matrices: It accepts an integer parameter. Currently, it can only be set as 0. Also, the left singular matrix and right singular matrix shapes are (M, K) and (K, N), respectively, where **K** = min(M, N), **M** is the number of rows and **N** is the number of columns of input matrix. (Default: 0)

lwork: This is an unused parameter. (Default: 0)

overwrite_a: It accepts an integer parameter, if set to 0, then ‘a’ will remain unchanged. Otherwise ‘a’ would be consumed internally and its contents will be destroyed. (Default: 0)

Purpose

It computes the singular value decomposition (SVD) of matrix ‘a’ with single (float32) precision.

The parameter: “lwork” is simply kept in to to make the interface uniform to the scipy.linalg.lapack.sgesvd() module. They are not used anywhere within the frovedis implementation.

This method internally uses Scalapack.gesvd() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
```

```
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
```

```
# sgesvd() demo
from frovedis.linalg.scalapack import sgesvd
rf = sgesvd(mat1)
print(rf)
```

Output

```
(array([[ 0.19046175, -0.12914878, -0.9731621 ],
        [-0.20606534, -0.97448295,  0.08899409],
        [-0.9598237 ,  0.1835851 , -0.2122148 ]], dtype=float32),
 array([9.838303 , 4.800165 , 1.0799255], dtype=float32),
 array([[ 0.04030442, -0.3974058 ,  0.91675735],
        [ 0.17610519, -0.90031475, -0.39802024],
        [-0.98354584, -0.17748773, -0.03369856]], dtype=float32), 0)
```

For example,

```
# sgesvd() demo and compute_uv = 0
from frovedis.linalg.scalapack import sgesvd
rf = sgesvd(mat1,compute_uv=0)
print(rf)
```

Output

```
(array([[0.]], dtype=float32), array([9.838302 , 4.800165 , 1.0799258], dtype=float32),
 array([[0.]], dtype=float32), 0)
```

For example,

```
# sgesvd() demo and overwriting of a
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import sgesvd
rf = sgesvd(mat1,overwrite_a=1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]

overwritten matrix:
[[-1.41421354 -3.80788684 -0.19258241]
 [-0.41421354  6.71975994  5.52667427]
 [ 0.          0.64659911 -5.36662769]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
```

```

bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)

# sgesvd() demo and a is an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import sgesvd
rf = sgesvd(bcm1)

# Unpacking the tuple
rf[0].debug_print()
rf[1].debug_print()
rf[2].debug_print()

Output

vector:
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.190462 -0.206065 -0.959824 -0.129149 -0.974483 0.183585 -0.973162 0.0889941 -0.212215
[9.838303  4.800165  1.0799255]
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.0403044 -0.397406 0.916757 0.176105 -0.900315 -0.39802 -0.983546 -0.177488 -0.0336986

```

Return Value

1. If 'a' is a python input such as numpy matrices:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It is a numpy matrix having float (float32) type values and containing the left singular matrix.
- **s**: It is a numpy matrix having float (float32) type values and containing the singular matrix.
- **vt**: It is a numpy matrix having float (float32) type values and containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If 'a' is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**u**, **s**, **vt**, **stat**) where,
- **u**: It returns an instance of FrovedisBlockcyclicMatrix containing the left singular matrix.
- **s**: It is a numpy matrix having float (float32) type values and containing the singular matrix.
- **vt**: It returns an instance of FrovedisBlockcyclicMatrix containing the right singular matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

38.4.1.10 10. dgetrf(a, overwrite_a = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise 'a' will be overwritten with 'L' and 'U' factors. (Default: 0)

Purpose

It computes LU factorization of matrix 'a' with double (float64) precision.

It computes an LU factorization of matrix 'a', using partial pivoting with row interchanges.

This method internally uses Scalapack.getrf() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
```

```
# dgetrf() demo
from frovedis.linalg.scalapack import dgetrf
rf = dgetrf(mat1)
print(rf)
```

Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]]),
      <frovedis.matrix.results.GetrfResult object at 0x7f80614aacf8>, 0)
```

This resultant tuple of `dgetrf()` wrapper function contains `GetrfResult` instance in `frovedis`. Whereas in `scipy`, it contains an array.

Both here are responsible to hold pivot array information.

For example,

```
# dgetrf() demo and overwriting of a
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import dgetrf
rf = dgetrf(mat1,overwrite_a=1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

```
overwritten matrix:
[[ 1.  0.  2.]
 [-1.  5.  2.]
 [ 0.  0.6 -10.2]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float64)
```

```
# dgetrf() demo and bcm1 is an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgetrf
rf = dgetrf(bcm1)
```

```
# Unpacking the tuple
rf[0].debug_print()
```

```
print(rf[1])
print(rf[2])
```

Output

```
<frovedis.matrix.results.GetrfResult object at 0x7f785d244978>
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 1 -1 0 0 5 0.6 2 2 -10.2
```

Return Value

1. If 'a' is a python inputs such as numpy matrix:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It is a numpy matrix having double (float64) type values (by default) and containing the LU factor of input matrix 'a'. In case 'overwrite_a' is enabled, then dtype for the matrix will depend on input 'a' dtype.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

2. If 'a' is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It returns an instance of FrovedisBlockcyclicMatrix containing the LU factor of input matrix 'a'.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

38.4.1.11 11. getrf(a, overwrite_a = 0, dtype = np.float64)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

overwrite_a: It accepts an integer parameter, if set to 0, then 'a' will remain unchanged. Otherwise 'a' will be overwritten with 'LU'. (Default: 0)

dtype: It specifies the datatype to be used for setting the precision level (single for float32 / double for float64) for the values returned by this method. (Default: np.float64)

Currently, it supports float (float32) or double (float64) datatypes.

Purpose

It computes the LU factorization of matrix 'a' with single (float32) or double (float64) precision depending on 'dtype' parameter provided by user.

It computes an LU factorization of matrix 'a', using partial pivoting with row interchanges.

This method is present only in frovedis.

This method internally uses Scalapack.getrf() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# getrf() demo
from frovedis.linalg.scalapack import getrf
rf = getrf(mat1)
print(rf)
```


Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]]),
      <frovedis.matrix.results.GetrfResult object at 0x7f80614aacf8>, 0)
```

For example,

```
# getrf() demo and overwriting of a
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import getrf
rf = getrf(mat1,overwrite_a=1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
```

```
overwritten matrix:
[[ 1.    0.    2. ]
 [-1.    5.    2. ]
 [ 0.    0.6 -10.2]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
# getrf() demo and specifying the dtype of output matrix LU factor
from frovedis.linalg.scalapack import getrf
rf = getrf(mat1,dtype=np.float32)
print(rf)
```

Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]], dtype=float32),
      <frovedis.matrix.results.GetrfResult object at 0x7f7c0bdbbd30>, 0)
```

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)

# getrf() demo and bcm1 is an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import getrf
rf = getrf(bcm1,dtype = np.float32)

# Unpacking the tuple
```

```
rf[0].debug_print()
print(rf[1])
print(rf[2])
```

Output

```
<frovedis.matrix.results.GetrfResult object at 0x7f56091379b0>
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 1 -1 0 0 5 0.6 2 2 -10.2
```

Note:- ‘dtype’ for the wrapper function and FrovedisBlockcyclicMatrix instance must be same during computation. Otherwise, it will raise an exception.

Return Value

1. If ‘a’ is a python input such as numpy matrix and dtype = np.float32:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It is a numpy matrix having float (float32) type values (by default) and containing the LU factor of input matrix ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

1. If ‘a’ is a python input such as numpy matrix and dtype = np.float64:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It is a numpy matrix having double (float64) type values (by default) and containing the LU factor of input matrix ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

3. If ‘a’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It returns an instance of FrovedisBlockcyclicMatrix containing the LU factor of input matrix ‘a’.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

38.4.1.12 12. sgetrf(a, overwrite_a = 0)

Parameters

a: It accepts a python array-like input or left hand side numpy matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values.

overwrite_a: It accepts an integer parameter, if set to 0, then ‘a’ will remain unchanged. Otherwise ‘a’ will be overwritten with ‘LU’. (Default: 0)

Purpose

It computes the LU factorization of matrix ‘a’ with single (float32) precision.

This method internally uses Scalapack.getrf() (**present in frovedis.matrix module**).

It computes an LU factorization of matrix ‘a’, using partial pivoting with row interchanges.

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# sgetrf() demo
```

```
from frovedis.linalg.scalapack import sgetrf
rf = sgetrf(mat1)
print(rf)
```

Output

```
(array([[ 1. ,  0. ,  2. ],
        [-1. ,  5. ,  2. ],
        [ 0. ,  0.6, -10.2]], dtype=float32),
 <frovedis.matrix.results.GetrfResult object at 0x7f1498750cf8>, 0)
```

This resultant tuple of `sgetrf()` wrapper function contains `GetrfResult` instance in `frovedis`. Whereas in `scipy`, it contains an array.

Both here are responsible to hold pivot array information.

For example,

```
# sgetrf() demo and overwriting of a
print('original matrix mat1: ')
print(mat1)
from frovedis.linalg.scalapack import sgetrf
rf = sgetrf(mat1,overwrite_a=1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
overwritten matrix:
[[ 1.          0.          2.          ]
 [-1.         5.          2.          ]
 [ 0.          0.60000002 -10.19999981]]
```

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)

# sgetrf() demo and bcm1 is an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import getrf
rf = getrf(bcm1,dtype = np.float32)

# Unpacking the tuple
rf[0].debug_print()
print(rf[1])
print(rf[2])
```

Output

```
<frovedis.matrix.results.GetrfResult object at 0x7f6199f28908>
```

```

0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 1 -1 0 0 5 0.6 2 2 -10.2

```

Return Value

1. If ‘a’ is a python input such as numpy matrix:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It is a numpy matrix having float (float32) type values (by default) and containing the LU factor of input matrix ‘a’. In case ‘overwrite_a’ is enabled, then dtype for the matrix will depend on input ‘a’ dtype.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

2. If ‘a’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**lu**, **res**, **stat**) where,
- **lu**: It returns an instance of FrovedisBlockcyclicMatrix containing the LU factor of input matrix ‘a’.
- **res**: It returns an instance of GetrfResult containing server side pointer of pivot array.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrf.

38.4.1.13 13. dgetri(lu, piv, lwork = 0, overwrite_lu = 0)

Parameters

lu: It accepts a python array-like input or numpy matrix having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values computed using dgetrf().

piv: It accepts an instance of GetrfResult containing server side pointer of pivot array computed using dgetrf().

lwork: This is an unused parameter. (Default: 0)

overwrite_lu: It accepts an integer parameter, if set to 0, then ‘lu’ will remain unchanged. Otherwise ‘lu’ will be overwritten with inverse matrix. (Default: 0)

Purpose

It computes the inverse matrix with double (float64) precision.

The parameter: “lwork” is simply kept in to to make the interface uniform to the scipy.linalg.lapack.dgetri() module. They are not used anywhere within the frovedis implementation.

This method internally uses Scalapack.getri() (**present in frovedis.matrix module**).

For example,

```

import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# dgetrf() demo
from frovedis.linalg.scalapack import dgetrf,dgetri

# compute lu and piv using dgetrf()
rf = dgetrf(mat1)

# dgetri() demo
ri = dgetri(rf[0],rf[1])
print(ri)

```

Output

```
(array([[ 0.88235294, -0.11764706,  0.19607843],
       [ 0.17647059,  0.17647059,  0.03921569],
       [ 0.05882353,  0.05882353, -0.09803922]]), 0)
```

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# dgetrf() demo
from frovedis.linalg.scalapack import dgetrf,dgetri

# compute lu and piv using dgetrf()
rf = dgetrf(mat1,overwrite_a = 1)

# dgetri() demo and overwriting of lu
print('original matrix mat1: ')
print(mat1)
ri = dgetri(rf[0],rf[1], overwrite_lu = 1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
overwritten matrix:
[[ 0.88235294 -0.11764706  0.19607843]
 [ 0.17647059  0.17647059  0.03921569]
 [ 0.05882353  0.05882353 -0.09803922]]
```

The input matrix is only processed by `dgetrf()` first to generate LU factor. The LU factor and ipiv information is then used with `dgetri()`.

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with inverse matrix having double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# dgetri() demo and bcm1 and bcm2 are an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgetrf,dgetri
rf = dgetrf(bcm1)
ri = dgetri(rf[0],rf[1],bcm2)

# Unpacking the tuple
ri[0].debug_print()
print(ri[1])
```

Output

```
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.882353 0.176471 0.0588235 -0.117647 0.176471 0.0588235 0.196078 0.0392157 -0.0980392
```

Return Value

1. If 'lu' is python input such as numpy matrix:

- It returns a tuple (**inv_a**, **stat**) where,
- **inv_a**: It is a numpy matrix having double (float64) type values (by default) and containing the inverse matrix. In case 'overwrite_lu' is enabled, then dtype for the matrix will depend on intermediate input 'LU' factor dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

2. If 'lu' is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**inv_a**, **stat**) where,
- **inv_a**: It returns an instance of FrovedisBlockcyclicMatrix containing the inverse matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

38.4.1.14 14. getri(lu, piv, lwork = 0, overwrite_lu = 0, dtype = np.float64)

Parameters

lu: It accepts a python array-like input or numpy matrix having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values.

piv: It accepts an instance of GetrfResult containing server side pointer of pivot array.

lwork: This is an unused parameter. (Default: 0)

overwrite_lu: It accepts an integer parameter, if set to 0, then 'lu' will remain unchanged. Otherwise 'lu' will be overwritten with inverse matrix. (Default: 0)

dtype: It specifies the datatype to be used for setting the precision level (single for float32 / double for float64) for the values returned by this method. (Default: np.float64)

Currently, it supports float (float32) or double (float64) datatypes.

Purpose

It computes the inverse matrix with single (float32) or double (float64) precision depending on 'dtype' parameter provided by user.

The parameter: "lwork" is simply kept in to to make the interface uniform to the scipy.linalg.lapack.dgetri() module. They are not used anywhere within the frovedis implementation.

This method is present only in frovedis.

This method internally uses Scalapack.getri() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# getrf() demo
from frovedis.linalg.scalapack import getrf,getri

# compute lu and piv using getrf()
rf = getrf(mat1)
```

```
# getri() demo
ri = getri(rf[0],rf[1])
print(ri)
```

Output

```
(array([[ 0.88235294, -0.11764706,  0.19607843],
        [ 0.17647059,  0.17647059,  0.03921569],
        [ 0.05882353,  0.05882353, -0.09803922]]), 0)
```

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
```

```
# getrf() demo
from frovedis.linalg.scalapack import getrf,getri
```

```
# compute lu and piv using getrf()
rf = getrf(mat1, overwrite_a = 1)
```

```
# getri() demo and overwriting of lu
print('original matrix mat1: ')
print(mat1)
ri = getri(rf[0],rf[1], overwrite_lu = 1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]
overwritten matrix:
[[ 0.88235294 -0.11764706  0.19607843]
 [ 0.17647059  0.17647059  0.03921569]
 [ 0.05882353  0.05882353 -0.09803922]]
```

The input matrix is only processed by `getrf()` first to generate LU factor. The LU factor and ipiv information is then used with `getri()`.

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with inverse matrix having double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
```

```
# getrf() demo
from frovedis.linalg.scalapack import getrf,getri
```

```
# compute lu and piv using getrf()
rf = getrf(mat1)
```

```
# getri() demo and specifying the dtype of output LU matrix
from frovedis.linalg.scalapack import getri
rf = getri(rf[0],rf[1],dtype=np.float32)
print(rf)
```

Output

```
(array([[ 0.88235295, -0.11764707,  0.19607843],
        [ 0.1764706 ,  0.1764706 ,  0.03921569],
        [ 0.05882353,  0.05882353, -0.09803922]]), dtype=float32), 0)
```

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)
bcm2 = FrovedisBlockcyclicMatrix(mat2,dtype = np.float32)

# sgetri() demo and bcm1 and bcm2 are an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import getrf,getri
rf = getrf(bcm1,dtype = np.float32)
ri = getri(rf[0],rf[1],bcm2, dtype = np.float32)

# Unpacking the tuple
ri[0].debug_print()
print(ri[1])
```

Output

```
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.882353 0.176471 0.0588235 -0.117647 0.176471 0.0588235 0.196078 0.0392157 -0.0980392
```

Note:- ‘dtype’ for the wrapper function and FrovedisBlockcyclicMatrix instance must be same during computation. Otherwise, it will raise an exception.

Return Value

1. If ‘lu’ is python input such as numpy matrix and dtype = np.float32:

- It returns a tuple (inv_a, stat) where,
- **inv_a**: It is a numpy matrix having float (float32) type values (by default) and containing the inverse matrix. In case ‘overwrite_lu’ is enabled, then dtype for the matrix will depend on intermediate input ‘LU’ factor dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

2. If ‘lu’ is python input such as numpy matrix and dtype = np.float64:

- It returns a tuple (inv_a, stat) where,
- **inv_a**: It is a numpy matrix having double (float64) type values (by default) and containing the inverse matrix. In case ‘overwrite_lu’ is enabled, then dtype for the matrix will depend on intermediate input ‘LU’ factor dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

3. If ‘lu’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (inv_a, stat) where,
- **inv_a**: It returns an instance of FrovedisBlockcyclicMatrix containing the inverse matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

38.4.1.15 15. sgetri(lu, piv, lwork = 0, overwrite_lu = 0)**Parameters**

lu: It accepts a python array-like input or numpy matrix having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values.

piv: It accepts an instance of GetrfResult containing server side pointer of pivot array.

lwork: This is an unused parameter. (Default: 0)

overwrite_lu: It accepts an integer parameter, if set to 0, then 'lu' will remain unchanged. Otherwise 'lu' will be overwritten with inverse matrix. (Default: 0)

Purpose

It computes the inverse matrix with single (float32) precision.

The parameter: "lwork" is simply kept in to to make the interface uniform to the scipy.linalg.lapack.dgetri() module. They are not used anywhere within the frovedis implementation.

This method internally uses Scalapack.getri() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# sgetrf() demo
from frovedis.linalg.scalapack import sgetrf,sgetri

# compute lu and piv using sgetrf()
rf = sgetrf(mat1)

# sgetri() demo
ri = sgetri(rf[0],rf[1])
print(ri)
```

Output

```
(array([[ 0.88235295, -0.11764707,  0.19607843],
        [ 0.1764706 ,  0.1764706 ,  0.03921569],
        [ 0.05882353,  0.05882353, -0.09803922]], dtype=float32), 0)
```

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# sgetrf() demo
from frovedis.linalg.scalapack import sgetrf,sgetri

# compute lu and piv using sgetrf()
rf = sgetrf(mat1, overwrite_a = 1)

# sgetri() demo and overwriting of lu
print('original matrix mat1: ')
print(mat1)
ri = sgetri(rf[0],rf[1], overwrite_lu = 1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat1:
[[ 1.  0.  2.]
 [-1.  5.  0.]
 [ 0.  3. -9.]]

overwritten matrix:
[[ 0.88235295 -0.11764707  0.19607843]
 [ 0.17647059  0.17647059  0.03921569]
 [ 0.05882353  0.05882353 -0.09803922]]
```

The input matrix is only processed by `sgetrf()` first to generate LU factor. The LU factor and ipiv information is then used with `sgetri()`.

Here, if the input ‘mat1’ is double (float64) type and overwrite is enabled, it’s overwritten with inverse matrix having double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)
bcm2 = FrovedisBlockcyclicMatrix(mat2,dtype = np.float32)

# sgetri() demo and bcm1 and bcm2 are an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import sgetrf,sgetri
rf = sgetrf(bcm1)
ri = sgetri(rf[0],rf[1],bcm2)

# Unpacking the tuple
ri[0].debug_print()
print(ri[1])
```

Output

```
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = 0.882353 0.176471 0.0588235 -0.117647 0.176471 0.0588235 0.196078 0.0392157 -0.0980392
```

Return Value

1. If ‘lu’ is python input such as numpy matrix:

- It returns a tuple (**inv_a**, **stat**) where,
- **inv_a**: It is a numpy matrix having float (float32) type values (by default) and containing the inverse matrix. In case ‘overwrite_lu’ is enabled, then dtype for the matrix will depend on intermediate input ‘LU’ factor dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

2. If ‘lu’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**inv_a**, **stat**) where,
- **inv_a**: It returns an instance of FrovedisBlockcyclicMatrix containing the inverse matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

38.4.1.16 16. dgetrs(lu, piv, b, trans = 0, overwrite_b = 0)**Parameters**

lu: It accepts a python array-like input or numpy matrix having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values computed using dgetrf.

piv: It accepts an instance of GetrfResult containing server side pointer of pivot array computed using dgetrf().

b: It accepts a python array-like input or right hand side matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having double (float64) type values. It should have number of rows \geq the number of rows in 'a' and at least 1 column in it.

trans: It accepts an integer parameter indicating if transpose of 'lu' needs to be computed before solving linear equation. If it is not 0, then the transpose is computed. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then 'b' will remain unchanged. Otherwise 'b' will be overwritten with the solution matrix. (Default: 0)

Purpose

It solves a system of linear equations, $AX = B$ with matrix 'a' using the LU factorization computed by getrf(). Thus before calling this function, it is required to obtain the factored matrix 'lu' (along with piv information) by calling getrf().

It computes the solution matrix for the system of linear equations with double (float64) precision.

This method internally uses Scalapack.getrs() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])

# dgetrf() demo
from frovedis.linalg.scalapack import dgetrf,dgetrs

# compute lu and piv using dgetrf()
rf = dgetrf(mat1)

# dgetrs() demo
ri = dgetrs(rf[0],rf[1],mat2)
print(ri)
```

Output

```
(array([[-7.54901961,  2.68627451, -1.          ],
        [-1.50980392,  1.1372549 ,  0.          ],
        [-0.7254902 ,  0.15686275,  0.          ]]), 0)
```

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])

# dgetrf() demo
from frovedis.linalg.scalapack import dgetrf,dgetrs

# compute lu and piv using dgetrf()
rf = dgetrf(mat1, overwrite_a = 1)
```

```
# dgetrs() demo and overwriting of b
print('original matrix mat2: ')
print(mat2)
ri = dgetrs(rf[0],rf[1],mat2,overwrite_b = 1)
print('overwritten matrix: ')
print(mat1)
```

Output

```
original matrix mat2:
[[-9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]
overwritten matrix:
[[-7.54901961  2.68627451 -1.          ]
 [-1.50980392  1.1372549  0.          ]
 [-0.7254902   0.15686275  0.          ]]
```

Here, if the inputs ‘mat2’ is double (float64) type and overwrite is enabled in dgetrs(), it’s overwritten with solution having double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```
from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1)
bcm2 = FrovedisBlockcyclicMatrix(mat2)

# dgetrs() demo and bcm1 and bcm2 are an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import dgetrf,dgetrs
rf = dgetrf(bcm1)
ri = dgetrs(rf[0],rf[1],bcm2)

# Unpacking the tuple
ri[0].debug_print()
print(ri[1])
```

Output

```
0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0
```

Return Value

1. If ‘b’ is python input such as numpy matrix:

- It returns a tuple (x, stat) where,
- x: It is a numpy matrix having double (float64) type values (by default) and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- stat: It returns an integer containing status (info) of native scalapack dgetrs.

2. If ‘b’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (**x**, **stat**) where,
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

38.4.1.17 17. getrs(lu, piv, b, trans = 0, overwrite_b = 0, dtype = np.float64)

Parameters

lu: It accepts a python array-like input or numpy matrix having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values computed using dgetrf.

piv: It accepts an instance of GetrfResult containing server side pointer of pivot array computed using dgetrf().

b: It accepts a python array-like input or right hand side matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) or double (float64) type values. It should have number of rows \geq the number of rows in 'a' and at least 1 column in it.

trans: It accepts an integer parameter indicating if transpose of 'lu' needs to be computed before solving linear equation. If it is not 0, then the transpose is computed. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then 'b' will remain unchanged. Otherwise 'b' will be overwritten with the solution matrix. (Default: 0)

dtype: It specifies the datatype to be used for setting the precision level (single for float32 / double for float64) for the values returned by this method. (Default: np.float64)

Currently, it supports float (float32) or double (float64) datatypes.

Purpose

It solves a system of linear equations, $AX = B$ with matrix 'a' using the LU factorization computed by getrf(). Thus before calling this function, it is required to obtain the factored matrix 'lu' by calling getrf().

It computes the solution matrix for the system of linear equations with float or double (float64) precision depending on 'dtype' parameter provided by user.

This method is present only in frovedis.

This method internally uses Scalapack.getrs() (**present in frovedis.matrix module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
```

```
# getrf() demo
from frovedis.linalg.scalapack import getrf,getrs
```

```
# compute lu and piv using getrf()
rf = getrf(mat1)
```

```
# getrs() demo
ri = getrs(rf[0],rf[1],mat2)
print(ri)
```

Output

```
(array([[ -7.54901961,  2.68627451, -1.          ],
        [-1.50980392,  1.1372549 ,  0.          ],
        [-0.7254902 ,  0.15686275,  0.          ]]), 0)
```

For example,

```

import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[-9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])

# getrf() demo
from frovedis.linalg.scalapack import getrf,getrs

# compute lu and piv using getrf()
rf = getrf(mat1, overwrite_a = 1)

# getrs() demo and overwriting of b
print('original matrix mat2: ')
print(mat2)
ri = getrs(rf[0],rf[1], mat2, overwrite_b = 1)
print('overwritten matrix: ')
print(mat2)

```

Output

```

original matrix mat2:
[[-9.  3. -1.]
 [ 0.  3.  1.]
 [ 2.  2.  0.]]
overwritten matrix:
[[-7.54901961  2.68627451 -1.          ]
 [-1.50980392  1.1372549   0.          ]
 [-0.7254902   0.15686275  0.          ]]

```

Here, if the inputs ‘mat2’ is double (float64) type and overwrite is enabled in dgetrs(), it’s overwritten with solution having double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```

import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# getrf() demo
from frovedis.linalg.scalapack import getrf,getrs

# compute lu and piv using getrf()
rf = getrf(mat1,dtype=np.float32)

# getrs() demo and specifying the dtype of output LU matrix
from frovedis.linalg.scalapack import getrs
rf = getrs(rf[0],rf[1],mat2,dtype=np.float32)
print(rf)

```

Output

```

(array([[ -7.54902    ,  2.6862745 , -1.          ],
        [-1.509804    ,  1.137255   ,  0.          ],
        [-0.7254902   ,  0.15686277 ,  0.          ]], dtype=float32), 0)

```

For example,

```

from frovedis.matrix.dense import FrovedisBlockcyclicMatrix

```

```

mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)
bcm2 = FrovedisBlockcyclicMatrix(mat2, dtype = np.float32)

# getsrs() demo and bcm1 and bcm2 are an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import getrf,getsrs
rf = getrf(bcm1,dtype = np.float32)
ri = getsrs(rf[0],rf[1],bcm2,dtype = np.float32)

# Unpacking the tuple
ri[0].debug_print()
print(ri[1])

Output

0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0

```

Note:- ‘dtype’ for the wrapper function and FrovedisBlockcyclicMatrix instance must be same during computation. Otherwise, it will raise an exception.

Return Value

1. If ‘b’ is python input such as numpy matrix and dtype = np.float32:

- It returns a tuple (x, stat) where,
- **x**: It is a numpy matrix having float (float32) type values (by default) containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetsrs.

2. If ‘b’ is python input such as numpy matrix and dtype = np.float64:

- It returns a tuple (x, stat) where,
- **x**: It is a numpy matrix having double (float64) type values (by default) containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetsrs.

3. If ‘b’ is an instance of FrovedisBlockcyclicMatrix:

- It returns a tuple (x, stat) where,
- **x**: It returns an instance of FrovedisBlockcyclicMatrix containing the solution matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

38.4.1.18 18. sgetsrs(lu, piv, b, trans = 0, overwrite_b = 0)

Parameters

lu: It accepts a python array-like input or numpy matrix having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values computed using dgetrf.

piv: It accepts an instance of GetrfResult containing server side pointer of pivot array computed using dgetrf().

b: It accepts a python array-like input or right hand side matrix of the linear equation having int, float (float32) or double (float64) type values. It also accepts FrovedisBlockcyclicMatrix instance having float (float32) type values. It should have number of rows >= the number of rows in ‘a’ and at least 1 column in it.

trans: It accepts an integer parameter indicating if transpose of ‘lu’ needs to be computed before solving linear equation. If it is not 0, then the transpose is computed. (Default: 0)

overwrite_b: It accepts an integer parameter, if set to 0, then 'b' will remain unchanged. Otherwise 'b' will be overwritten with the solution matrix. (Default: 0)

Purpose

It solves a system of linear equations, $AX = B$ with the matrix 'a' using the LU factorization computed by `sgetrf()`.

Thus before calling this function, it is required to obtain the factored matrix 'lu' by calling `sgetrf()`.

It computes the solution matrix for the system of linear equations with float precision.

This method internally uses `Scalapack.getrs()` (**present in `frovedis.matrix` module**).

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[ -9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])

# sgetrf() demo
from frovedis.linalg.scalapack import sgetrf,sgetrs

# compute lu and piv using sgetrf()
rf = sgetrf(mat1)

# sgetrs() demo
ri = sgetrs(rf[0],rf[1],mat2)
print(ri)
```

Output

```
(array([[ -7.54902    ,  2.6862745 , -1.          ],
       [-1.509804   ,  1.137255   ,  0.          ],
       [-0.7254902  ,  0.15686277 ,  0.          ]], dtype=float32), 0)
```

For example,

```
import numpy as np
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])

# sgetrf() demo
from frovedis.linalg.scalapack import sgetrf,sgetrs

# compute lu and piv using sgetrf()
rf = sgetrf(mat1)

# sgetrs() demo and overwriting of b
print('original matrix mat2: ')
print(mat2)
ri = sgetrs(rf[0],rf[1], mat2, overwrite_b = 1)
print('overwritten matrix: ')
print(mat2)
```

Output

```
original matrix mat2:
[[ -9.   3.  -1.]
 [  0.   3.   1.]
 [  2.   2.   0.]]
overwritten matrix:
```



```

[[-7.54901981  2.68627453 -1.          ]
 [-1.50980401  1.13725495  0.          ]
 [-0.72549021  0.15686277  0.          ]]

```

Here, if the inputs ‘mat2’ is double (float64) type and overwrite is enabled in `dgetrs()`, it’s overwritten with solution having double (float64) type values.

Same applies for other types (int, float (float32)) when input is a numpy matrix/array and overwrite is enabled.

For example,

```

from frovedis.matrix.dense import FrovedisBlockcyclicMatrix
mat1 = np.array([[1.,0.,2.],[-1.,5.,0.],[0.,3.,-9.]])
mat2 = np.array([[-9.,3.,-1.],[0.,3.,1.],[2.,2.,0.]])
bcm1 = FrovedisBlockcyclicMatrix(mat1,dtype = np.float32)
bcm2 = FrovedisBlockcyclicMatrix(mat2,dtype = np.float32)

# sgetrs() demo and bcm1 and bcm2 are an instance of FrovedisBlockcyclicMatrix
from frovedis.linalg.scalapack import sgetrf,sgetrs
rf = sgetrf(bcm1)
ri = sgetrs(rf[0],rf[1],bcm2)

# Unpacking the tuple
ri[0].debug_print()
print(ri[1])

```

Output

```

0
matrix:
num_row = 3, num_col = 3
node 0
node = 0, local_num_row = 3, local_num_col = 3, type = 2, descriptor = 1 1 3 3 3 3 0 0 3 3 3
val = -7.54902 -1.5098 -0.72549 2.68627 1.13725 0.156863 -1 0 0

```

Return Value

1. If ‘b’ is python input such as numpy matrix:

- It returns a tuple (**x**, **stat**) where,
- **x**: It is a numpy matrix having float (float32) type value and containing the solution matrix. In case ‘overwrite_b’ is enabled, then dtype for the matrix will depend on input ‘b’ dtype.
- **stat**: It returns an integer containing status (info) of native scalapack dgetrs.

2. If ‘b’ is an instance of **FrovedisBlockcyclicMatrix**:

- It returns a tuple (**x**, **stat**) where,
- **x**: It returns an instance of **FrovedisBlockcyclicMatrix** containing the solution matrix.
- **stat**: It returns an integer containing status (info) of native scalapack dgetri.

38.5 SEE ALSO

- **Linalg Functions**

Chapter 39

Graph

39.1 NAME

Graph - This class implements a directed or undirected frovedis graph.

39.2 SYNOPSIS

```
frovedis.graph.graph.Graph(nx_graph=None)
```

39.2.1 Public Member Functions

```
load(nx_graph)
load_csr(smat)
debug_print()
release()
clear()
number_of_edges()
number_of_nodes()
save(fname)
load_text(fname)
to_networkx_graph()
```

39.3 DESCRIPTION

Base class for undirected and directed frovedis graphs. A Frovedis Graph stores nodes and edges with optional data. Here, the graph data, edge to edge information is internally stored as and adjacency matrix data.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis public method interface is almost same as NetworkX Graph interface, but it doesn't have any dependency with NetworkX. It can be used simply even if the system doesn't have NetworkX installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for Graph() on the frovedis server. Once the graph is loaded for the given input networkx or scipy csr_matrix graph data at the frovedis server, it returns a frovedis graph instance to the client python program.

39.3.1 Detailed Description

39.3.1.1 1. Graph()

Parameters

nx_graph: Data to initialize graph. Here data provided is a scipy sparse csr_matrix or a networkx.Graph instance. It loads such data to create a frovedis graph. (Default: None)

When it is None, an empty frovedis graph is created.

Purpose

It initializes a Frovedis Graph object having nodes and edges. It can create both directed or undirected graph.

For example,

a) When loading an undirected networkx.Graph as input,

```
import frovedis.graph as fnx
import networkx as gn timer
net_graph = gn timer.read_edgelist('input/cit-Patents_10.txt')
frov_graph = fnx.Graph(net_graph)
```

b) When loading a directed networkx.Graph as input,

```
import frovedis.graph as fnx
import networkx as gn timer
net_graph = gn timer.read_edgelist('input/cit-Patents_10.txt', create_using = gn timer.DiGraph())
frov_graph = fnx.Graph(net_graph)
```

c) When loading a scipy sparse csr_matrix as input to create undirected frovedis graph,

```
#Here it creates an undirected graph
import frovedis.graph as fnx
from scipy.sparse import csr_matrix
data = [1,1,1,1,1,1,1,1,1,1,1,1,1,1,1,1,1]
row = [0,0,0,0,0,1,2,3,4,5,6,6,6,6,7,8,9,10,11,12]
col = [1,2,3,4,5,0,0,0,0,0,7,8,9,10,6,6,6,6,12,11]
csr_mat = csr_matrix((data, (row, col)), shape = (13, 13))
frov_graph = fnx.Graph(csr_mat)
```

d) Again, when loading a scipy sparse csr_matrix as input to create directed frovedis graph,

```
#Here it creates a directed graph
import frovedis.graph as fnx
from scipy.sparse import csr_matrix
data = [1,1,1,1,1,1,1,1,1,1,1]
row = [0,0,0,0,0,6,6,6,6,11]
col = [1,2,3,4,5,7,8,9,10,12]
csr_mat = csr_matrix((data, (row, col)), shape = (13, 13))
frov_graph = fnx.Graph(csr_mat)
```

Return Value

It simply returns “self” reference.

39.3.1.2 2. load(nx_graph)**Parameters**

nx_graph: Data to initialize graph. Here graph data provided is a networkx.Graph instance.

Purpose

It loads a networkx graph to create a frovedis graph. Here, the loaded graph can be directed or undirected.

For example,

```
import frovedis.graph as fnx
import networkx as gnx
net_graph = gnx.read_edgelist('input/cit-Patents_10.txt')
frov_graph = fnx.Graph().load(nx_graph = net_graph)
```

Return Value

It simply returns “self” reference.

39.3.1.3 3. load_csr(smat)**Parameters**

smat: Data to initialize graph. Here graph data provided is a scipy sparse csr_matrix.

Purpose

It loads frovedis graph from a scipy csr_matrix. Here, depending on the scipy csr_matrix data, the final graph maybe directed or undirected.

For example,

```
#Loading a scipy csr_matrix to create an undirected frovedis graph
import frovedis.graph as fnx
from scipy.sparse import csr_matrix
data = [1,1,1,1,1,1,1,1,1,1,1,1,1,1,1,1,1]
row = [0,0,0,0,0,1,2,3,4,5,6,6,6,6,7,8,9,10,11,12]
col = [1,2,3,4,5,0,0,0,0,0,7,8,9,10,6,6,6,6,12,11]
csr_mat = csr_matrix((data, (row, col)), shape = (13, 13))
frov_graph = fnx.Graph().load_csr(smat = csr_mat)
```

```
#Loading a scipy csr_matrix to create a directed frovedis graph
import frovedis.graph as fnx
from scipy.sparse import csr_matrix
data = [1,1,1,1,1,1,1,1,1,1]
row = [0,0,0,0,0,6,6,6,6,11]
col = [1,2,3,4,5,7,8,9,10,12]
csr_mat = csr_matrix((data, (row, col)), shape = (13, 13))
frov_graph = fnx.Graph().load_csr(smat = csr_mat)
```

Return Value

It simply returns “self” reference.

39.3.1.4 4. debug_print()**Purpose**

It shows graph information on the client side and server side user terminal. It is mainly used for debugging purpose.

For example,

```
frov_graph.debug_print()
```

Output on client side

```
Num of edges: 20
```

```
Num of vertices: 13
```

It displays information such as number of edges and vertices.

Output on server side

```
Num of edges: 20
```

```
Num of vertices: 13
```

```
is directed: 0
```

```
is weighted: 0
```

```
in-degree:
```

```
5 1 1 1 1 1 4 1 1 1 1 1 1
```

```
out-degree:
```

```
5 1 1 1 1 1 4 1 1 1 1 1 1
```

```
adjacency matrix:
```

```
0 1 1 1 1 1 0 0 0 0 0 0 0
```

```
1 0 0 0 0 0 0 0 0 0 0 0 0
```

```
1 0 0 0 0 0 0 0 0 0 0 0 0
```

```
1 0 0 0 0 0 0 0 0 0 0 0 0
```

```
1 0 0 0 0 0 0 0 0 0 0 0 0
```

```
1 0 0 0 0 0 0 0 0 0 0 0 0
```

```
0 0 0 0 0 0 0 1 1 1 1 0 0
```

```
0 0 0 0 0 0 1 0 0 0 0 0 0
```

```
0 0 0 0 0 0 1 0 0 0 0 0 0
```

```
0 0 0 0 0 0 1 0 0 0 0 0 0
```

```
0 0 0 0 0 0 1 0 0 0 0 0 0
```

```
0 0 0 0 0 0 0 0 0 0 0 0 1
```

```
0 0 0 0 0 0 0 0 0 0 0 1 0
```

It displays information such as number of edges and vertices, directed ('0' for No, '1' for Yes) , incoming links, outgoing links and an adjacency matrix. Currently, it shows information about an undirected frovedis graph.

Return Value

It returns nothing.

39.3.1.5 5. release()

Purpose

It can be used to release the in-memory model at frovedis server.

For example,

```
frov_graph.release()
```

This will remove the graph model, model-id present on server, along with releasing server side memory.

Return Value

It returns nothing.

39.3.1.6 6. clear()

Purpose

It can be used to release the in-memory model at frovedis server. This method is an alias to release().

For example,

```
frov_graph.clear()
```

This will remove the graph model, model-id present on server, along with releasing server side memory.

Return Value

It returns nothing.

39.3.1.7 7. number_of_edges()

Purpose

It is used to fetch the number of edges present in the directed or undirected frovedis graph.

For example,

```
edge_count = frov_graph.number_of_edges()
print('Number of edges: ', edge_count)
```

Output

```
Number of edges: 20
```

Return Value

It returns a long (int64) type value specifying the number of edges.

39.3.1.8 8. number_of_nodes()

Purpose

It is used to fetch the number of nodes/vertices present in the directed or undirected frovedis graph.

For example,

```
nodes_count = frov_graph.number_of_nodes()
print('Number of nodes: ', nodes_count)
```

Output

```
Number of nodes: 13
```

Return Value

It returns a long (int64) type value specifying the number of node/vertices.

39.3.1.9 9. save(fname)

Parameters

fname: A string object containing the name of the file on which the adjacency matrix is to be saved.

Purpose

On success, it writes the adjacency matrix information in the specified file. Otherwise, it throws an exception.

For example,

```
# To save the frovedis graph
frov_graph.save("./out/FrovGraph")
```

This will save the frovedis graph on the path '/out/FrovGraph'. It would raise exception if the directory already exists with same name.

Return Value

It returns nothing.

39.3.1.10 10. load_text(fname)**Parameters**

fname: A string object containing the name of the file having adjacency matrix information to be loaded.

Purpose

It loads a frovedis graph from the specified file path (having adjacency matrix data).

For example,

```
# To load the frovedis graph
frov_graph.load_text("./out/FrovGraph")
```

Return Value

It simply returns “self” reference.

39.3.1.11 11. to_networkx_graph()**Purpose**

It is used to convert a frovedis graph into a networkx graph.

For example,

```
nx_graph = frov_graph.to_networkx_graph()
print(nx_graph.adj)
```

Output

```
{0: {1: {'weight': 1.0}, 2: {'weight': 1.0}, 3: {'weight': 1.0}, 4: {'weight': 1.0},
5: {'weight': 1.0}}, 1: {0: {'weight': 1.0}}, 2: {0: {'weight': 1.0}},
3: {0: {'weight': 1.0}}, 4: {0: {'weight': 1.0}}, 5: {0: {'weight': 1.0}},
6: {7: {'weight': 1.0}, 8: {'weight': 1.0}, 9: {'weight': 1.0}, 10: {'weight': 1.0}},
7: {6: {'weight': 1.0}}, 8: {6: {'weight': 1.0}}, 9: {6: {'weight': 1.0}},
10: {6: {'weight': 1.0}}, 11: {12: {'weight': 1.0}}, 12: {11: {'weight': 1.0}}}
```

Return Value

It returns a networkx.Graph instance.

39.4 SEE ALSO

- Graph in Frovedis
- Breadth First Search in Frovedis
- Connected Components in Frovedis
- PageRank in Frovedis
- Single Source Shortest Path in Frovedis

Chapter 40

pagerank()

40.1 NAME

`pagerank()` - It computes a ranking of the nodes in the graph 'G' based on the structure of the incoming links. It was originally designed as an algorithm to rank web pages.

40.2 SYNOPSIS

```
frovedis.graph.Pagerank.pagerank(G, alpha=0.85, personalization=None, max_iter=100, tol=1.0e-6,
                                nstart=None, weight='weight', dangling=None, verbose=0)
```

40.3 DESCRIPTION

The PageRank algorithm ranks the nodes in a graph by their relative importance or influence. PageRank determines each node's ranking by identifying the number of links to the node and the quality of the links. The quality of a link is determined by the importance (PageRank) of the node that presents the outbound link.

The PageRank algorithm was designed at first to measure the importance of a webpage by analyzing the quantity and quality of the links that point to it.

Mathematically, PageRank (PR) is defined as:

$$PR(A) = (1 - d) + d(PR(T_1)/C(T_1) + \dots + PR(T_n)/C(T_n))$$

where **Page A** has pages **T1 to Tn** which point to it.

d is a damping factor. It corresponds to the probability that an imaginary web surfer will suddenly visit a random page on the web instead of following one of the outbound links prescribed by the web page that the surfer is currently visiting. This is useful for accounting for sinks (web pages that have inbound links but no outbound links).

Also, the **(1 - d)** in the preceding formula dampens the contribution of the incoming PageRanks from the adjacent vertices. It is as if imaginary outbound edges are added from all sink vertices to every other vertex in the graph, and to keep things fair, this same thing is done to the non-sink vertices as well.

C(A) is defined as the number of links going out of page A.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis public method interface is almost same as NetworkX pagerank public method interface, but

it doesn't have any dependency with NetworkX. It can be used simply even if the system doesn't have NetworkX installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for pagerank() on the frovedis server. Once the PageRanks are computed for the given input graph data at the frovedis server, it returns a dictionary of nodes with PageRank as value to the client python program.

40.3.1 Detailed Description

40.3.1.1 1. pagerank()

Parameters

G: An instance of networkx graph or frovedis graph on which pagerank is to be computed. The graph can be directed or undirected.

alpha: A positive integer parameter that is referred as **damping factor**. It must be in range 0 and 1. It represents a sort of minimum PageRank value. (Default: 0.85)

personalization: An unused parameter. (Default: None)

max_iter: A positive integer parameter that specifies the maximum number of iterations to run this method. Convergence may occur before max_iter value, in that case the iterations will stop. (Default: 100)

tol: A positive double (float64) parameter specifying the convergence tolerance. The PageRank iterations stop if the sum of the error values for all nodes is below this value. (Default: 1.0e-6)

nstart: An unused parameter. (Default: None)

weight: An unused parameter. (Default: 'weight')

dangling: An unused parameter. (Default: None)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method ranks the nodes present in the graph with the given parameters.

The parameters: "personalization", "nstart", "weight" and "dangling" are simply kept to make the method uniform to NetworkX pagerank method. They are not used anywhere within the frovedis implementation.

For example,

When loading an undirected networkx.graph as input from an edgelist file,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# An undirected graph loaded from edgelist file
import networkx as gnx
```

```
networkx_graph = gnz.read_edgelist('input/cit-Patents.txt')
ranks = gnz.pagerank(networkx_graph)
print ("NX PR: ", ranks)
```

Output

```
[rank 0] pagerank: converged in 8 iterations!
NX PR: {1: 0.11705684590144233, 2: 0.06889632312740385, 3: 0.06889632312740385,
4: 0.06889632312740385, 5: 0.06889632312740385, 6: 0.06889632312740385, 7: 0.10702340365685098,
8: 0.06939799523963341, 9: 0.06939799523963341, 10: 0.06939799523963341,
11: 0.06939799523963341, 12: 0.07692307692307693, 13: 0.07692307692307693}
```

When loading an undirected frovedis graph as input from an edgelist file,

FILE: cit-Patents.txt (same file)

```
# An undirected graph loaded from edgelist file
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents.txt')
ranks = fnx.pagerank(frov_graph)
print ("FROV PR: ", ranks)
```

Output

```
[rank 0] pagerank: converged in 8 iterations!
FROV PR: {1: 0.11705684590144233, 2: 0.06889632312740385, 3: 0.06889632312740385,
4: 0.06889632312740385, 5: 0.06889632312740385, 6: 0.06889632312740385, 7: 0.10702340365685098,
8: 0.06939799523963341, 9: 0.06939799523963341, 10: 0.06939799523963341,
11: 0.06939799523963341, 12: 0.07692307692307693, 13: 0.07692307692307693}
```

When loading an directed networkx.graph as input from an edgelist file,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import networkx as gnz
networkx_graph = gnz.read_edgelist('input/cit-Patents.txt', create_using = gnz.DiGraph())
ranks = gnz.pagerank(networkx_graph)
print ("NX PR: ", ranks)
```

Output

```
[rank 0] pagerank: converged in 3 iterations!
NX PR: {1: 0.06538461538461539, 2: 0.06734615384615385, 3: 0.06734615384615385,
4: 0.06734615384615385, 5: 0.06734615384615385, 6: 0.06734615384615385, 7: 0.06538461538461539,
8: 0.06783653846153846, 9: 0.06783653846153846, 10: 0.06783653846153846,
11: 0.06783653846153846, 12: 0.06538461538461539, 13: 0.0751923076923077}
```

When loading an directed frovedis graph as input from an edgelist file,

FILE: cit-Patents.txt (same file)

```
# A directed graph loaded from edgelist file
import frovedis.graph as fnx
import networkx
frov_graph = fnx.read_edgelist('input/cit-Patents.txt', create_using = networkx.DiGraph())
ranks = fnx.pagerank(frov_graph)
print ("FROV PR: ", ranks)
```

Output

```
[rank 0] pagerank: converged in 3 iterations!
FROV PR: {1: 0.06538461538461539, 2: 0.06734615384615385, 3: 0.06734615384615385,
4: 0.06734615384615385, 5: 0.06734615384615385, 6: 0.06734615384615385, 7: 0.06538461538461539,
8: 0.06783653846153846, 9: 0.06783653846153846, 10: 0.06783653846153846,
11: 0.06783653846153846, 12: 0.06538461538461539, 13: 0.0751923076923077}
```

Return Value

It returns a dictionary of nodes with PageRank as value.

40.4 SEE ALSO

- Graph in Frovedis
- Breadth First Search in Frovedis
- Connected Components in Frovedis
- Single Source Shortest Path in Frovedis

Chapter 41

Breadth First Search

41.1 NAME

Breadth First Search - Basic algorithms for breadth-first searching the nodes of a graph.

41.2 SYNOPSIS

```
frovedis.graph.traversal.bfs(G, source, depth_limit=None, opt_level=1, hyb_threshold=0.4,  
                             verbose=0)
```

```
frovedis.graph.traversal.bfs_edges(G, source, reverse=False, depth_limit=None,  
                                   sort_neighbors=None, opt_level=1, hyb_threshold=0.4,  
                                   verbose=0)
```

```
frovedis.graph.traversal.bfs_tree(G, source, reverse=False, depth_limit=None,  
                                  sort_neighbors=None, opt_level=1, hyb_threshold=0.4,  
                                  verbose=0)
```

```
frovedis.graph.traversal.bfs_predecessors(G, source, depth_limit=None, sort_neighbors=None,  
                                           opt_level=1, hyb_threshold=0.4, verbose=0)
```

```
frovedis.graph.traversal.bfs_successors(G, source, depth_limit=None, sort_neighbors=None,  
                                         opt_level=1, hyb_threshold=0.4, verbose=0)
```

```
frovedis.graph.traversal.descendants_at_distance(G, source, distance, opt_level=1,  
                                                hyb_threshold=0.4, verbose=0)
```

41.3 DESCRIPTION

Breadth-first search is an algorithm for searching a tree data structure for a node that satisfies a given property. It starts at the tree root and explores all nodes at the present depth prior to moving on to the nodes at the next depth level.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis breadth first search public method interfaces are almost same as NetworkX breadth first search

public method interface, but it doesn't have any dependency on NetworkX. It can be used simply even if the system doesn't have NetworkX installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

41.3.1 Detailed Description

41.3.1.1 1. bfs()

Parameters

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for breadth-first search. This method iterates over only those edges in the component, reachable from this node. It must be in range [1, **G.num_vertices**].

depth_limit: A positive integer parameter that specifies the maximum search depth. (Default: None)
When it is None (not specified explicitly), it will be set as maximum value for int64 datatype.

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter that is used for reducing computation times during breadth first search. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.
- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.
- **When opt_level = 2**: this is much better than '**opt_level = 0**' but slightly slower than **opt_level = 1**. It optimizes the memory usage over '**opt_level = 1**'.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization during breadth first search when the number of remaining nodes to be visited becomes less then this value. It optimizes the execution time. This parameter works only with '**opt_level = 2**'. It must be within the range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method computes bfs traversal path of a graph.

This method is not present in Networkx. It is only provided in frovedis.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
```

```
import networkx as nx
import frovedis.graph as fnx
G = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                    create_using = nx.DiGraph())
print ("Frovedis BFS : ", list(fnx.bfs(G, source = 1, depth_limit = 1)))
```

Output

```
Frovedis BFS :  [{1: [1]}, {2: [1, 2]}, {3: [1, 3]}, {4: [1, 4]}, {5: [1, 5]}, {6: [1, 6]}]
```

In case, we started from node 7 i.e source = 7 in the directed graph ‘G’

```
print ("Frovedis BFS : ", list(fnx.bfs(G, source = 7, depth_limit = 1)))
```

Output

```
Frovedis BFS :  [{7: [7]}, {8: [7, 8]}, {9: [7, 9]}, {10: [7, 10]}, {11: [7, 11]}]
```

Return Value

It returns a dictionary with keys as destination node-id and values as corresponding traversal path from the source. In case any node in input graph not reachable from the source, it would not be included in the resultant dictionary.

41.3.1.2 2. bfs_edges()

Parameters

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for breadth-first search. This method iterates over only those edges in the component, reachable from this node. It must be in range [1, **G.num_vertices**].

reverse: A boolean parameter, in general, specifies the direction of traversal (forward or backward) if ‘G’ is a directed graph. Currently, it can only traverse in forward direction (**reverse = True not supported yet**). (Default: False)

depth_limit: A positive integer parameter that specifies the maximum search depth. (Default: None)
When it is None (not specified explicitly), it will be set as maximum value for int64 datatype.

sort_neighbors: An unused parameter. (Default: None)

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter that is used for reducing computation times during breadth first search. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.

- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.

- **When opt_level = 2**: this is much better than ‘opt_level = 0’ but slightly slower than **opt_level = 1**. It optimizes the memory usage over ‘opt_level = 1’.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization during breadth first search when the number of remaining nodes to be visited becomes less than this value. It optimizes the execution time. This parameter works only with ‘opt_level = 2’. It must be within the range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method iterates over edges in a breadth-first-search starting at source.

The parameter: “sort_neighbors” is simply kept to make the method uniform to NetworkX bfs_edges method. It is not used anywhere within the frovedis implementation.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
G = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                    create_using = nx.DiGraph())
print("Frovedis BFS edges: ", list(fnx.bfs_edges(G, source = 1, depth_limit = 1)))
```

Output

```
Frovedis BFS edges: [(1, 2), (1, 3), (1, 4), (1, 5), (1, 6)]
```

In case, we started from node 7 i.e source = 7 in the directed graph ‘G’

```
print("Frovedis BFS edges: ", list(fnx.bfs_edges(G, source = 7, depth_limit = 1)))
```

Output

```
Frovedis BFS edges: [(7, 8), (7, 9), (7, 10), (7, 11)]
```

Return Value

It yields edges resulting from the breadth-first search.

41.3.1.3 3. bfs_tree()

Parameters

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for breadth-first search. This method iterates over only those edges in the component, reachable from this node. It must be in range [1, G.num_vertices].

reverse: A boolean parameter, in general, specifies the direction of traversal (forward or backward) if ‘G’ is a directed graph. Currently, it can only traverse in forward direction (**reverse = True not supported yet**). (Default: False)

depth_limit: A positive integer parameter that specifies the maximum search depth. (Default: None)
When it is None (not specified explicitly), it will be set as maximum value for int64 datatype.

sort_neighbors: An unused parameter. (Default: None)

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter that is used for reducing computation times during breadth first search. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.

- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.

- **When opt_level = 2**: this is much better than ‘opt_level = 0’ but slightly slower than opt_level = 1. It optimizes the memory usage over ‘opt_level = 1’.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization during breadth first search when the number of remaining nodes to be visited becomes less than this value. It optimizes the execution time. This parameter works only with ‘**opt_level = 2**’. It must be within the range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method constructs a directed tree from of a breadth-first-search starting at source.

The parameter: “sort_neighbors” is simply kept to make the method uniform to NetworkX bfs_tree method. It is not used anywhere within the frovedis implementation.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
G = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                    create_using = nx.DiGraph())
fnx.bfs_tree(G, source = 1, depth_limit = 1)

# To check edges in newly created networkx directed graph
print("Edges in Frovedis bfs_tree: ")
print(fnx.bfs_tree(G, source = 1, depth_limit = 1).number_of_edges())
```

Output

```
Edges in Frovedis bfs_tree:
5
```

In case, we started from node 7 i.e source = 7 in the directed graph ‘G’

```
fnx.bfs_tree(G, source = 7, depth_limit = 1)

# To check edges in newly created networkx directed graph
print("Edges in Frovedis bfs_tree: ")
print(fnx.bfs_tree(G, source = 7, depth_limit = 1).number_of_edges())
```

Output

```
Edges in Frovedis bfs_tree:
```

4

Return Value

It returns a new Networkx.DiGraph instance.

41.3.1.4 4. bfs_predecessors()**Parameters**

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for breadth-first search. This method iterates over only those edges in the component, reachable from this node. It must be in range [1, **G.num_vertices**].

depth_limit: A positive integer parameter that specifies the maximum search depth. (Default: None)
When it is None (not specified explicitly), it will be set as maximum value for int64 datatype.

sort_neighbors: An unused parameter. (Default: None)

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter that is used for reducing computation times during breadth first search. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.
- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.
- **When opt_level = 2**: this is much better than ‘opt_level = 0’ but slightly slower than **opt_level = 1**. It optimizes the memory usage over ‘opt_level = 1’.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization during breadth first search when the number of remaining nodes to be visited becomes less then this value. It optimizes the execution time. This parameter works only with ‘opt_level = 2’. It must be within the range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method provides predecessors for each node in breadth-first-search from source.

The parameter: “sort_neighbors” is simply kept to make the method uniform to NetworkX bfs_tree method. It is not used anywhere within the frovedis implementation.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
```

```
G = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                    create_using = nx.DiGraph())
print("Frovedis bfs_predecessors: ")
print(list(fnx.bfs_predecessors(G, source = 1, depth_limit = 1)))
```

Output

```
Frovedis bfs_predecessors:
[(2, 1), (3, 1), (4, 1), (5, 1), (6, 1)]
```

Here, node 2, node 3, node 4, node 5 and node 6 were having source = 1 as their predecessor.

In case, we started from node 7 i.e source = 7 in the directed graph ‘G’

```
print("Frovedis bfs_predecessors: ")
print(list(fnx.bfs_predecessors(G, source = 7, depth_limit = 1)))
```

Output

```
Frovedis bfs_predecessors:
[(8, 7), (9, 7), (10, 7), (11, 7)]
```

Here, node 8, node 9, node 10 and node 11 were having source = 7 as their predecessor.

Return Value

It yields predecessors for each node in form of **(node, predecessor)** iterator.

41.3.1.5 5. bfs_successors()

Parameters

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for breadth-first search. This method iterates over only those edges in the component, reachable from this node. It must be in range `[1, G.num_vertices]`.

depth_limit: A positive integer parameter that specifies the maximum search depth. (Default: None) When it is None (not specified explicitly), it will be set as maximum value for int64 datatype.

sort_neighbors: An unused parameter. (Default: None)

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter that is used for reducing computation times during breadth first search. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.

- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.

- **When opt_level = 2**: this is much better than ‘opt_level = 0’ but slightly slower than opt_level = 1. It optimizes the memory usage over ‘opt_level = 1’.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization during breadth first search when the number of remaining nodes to be visited becomes less than this value. It optimizes the execution time. This parameter works only with ‘opt_level = 2’. It must be within the range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method provides successors for each node in breadth-first-search from source.

The parameter: “sort_neighbors” is simply kept to make the method uniform to NetworkX bfs_tree method. It is not used anywhere within the frovedis implementation.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
G = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                    create_using = nx.DiGraph())
print("Frovedis bfs_successors: ", list(fnx.bfs_successors(G, source = 1, depth_limit = 1)))

Output
```

```
Frovedis bfs_successors:  [(1, [2, 3, 4, 5, 6])]
```

Here, for source = 1, node 2, node 3, node 4, node 5 and node 6 were its successor.

In case, we started from node 7 i.e source = 7 in the directed graph ‘G’

```
print("Frovedis bfs_successors: ", list(fnx.bfs_successors(G, source = 7, depth_limit = 1)))

Output
```

```
Frovedis bfs_successors:  [(7, [8, 9, 10, 11])]
```

Here, for source = 7, node 8, node 9, node 10 and node 11 were its successor.

Return Value

It yields successors for each node in form of **(node, successor)** iterator.

41.3.1.6 6. descendants_at_distance()

Parameters

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for breadth-first search. This method iterates over only those edges in the component, reachable from this node. It must be in range [1, G.num_vertices].

distance: A positive integer parameter that specifies the distance of the wanted nodes from source.

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter is used for reducing computation times during breadth first search. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.

- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.

- **When opt_level = 2**: this is much better than ‘opt_level = 0’ but slightly slower than opt_level = 1. It optimizes the memory usage over ‘opt_level = 1’.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization during breadth first search when the number of remaining nodes to be visited becomes less than this value. It optimizes the execution time. This parameter works only with ‘opt_level = 2’. It must be within the

range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method returns all nodes at a fixed distance from source in G.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
G = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                    create_using = nx.DiGraph())
print("Frovedis descendants at distance:")
print(fnx.descendants_at_distance(G, source = 1, distance = 1))
```

Output

```
Frovedis descendants at distance:
{2, 3, 4, 5, 6}
```

In case, we started from node 7 i.e source = 7 in the directed graph 'G'

```
print("Frovedis descendants at distance:")
print(fnx.descendants_at_distance(G, source = 7, distance = 1))
```

Output

```
Frovedis descendants at distance:
{8, 9, 10, 11}
```

Return Value

It returns an instance of Set having the descendants of source in graph 'G' at the given distance from source.

41.4 SEE ALSO

- Graph in Frovedis
- Connected Components in Frovedis

- **Single Source Shortest Path in Frovedis**
- **PageRank in Frovedis**

Chapter 42

single_source_shortest_path()

42.1 NAME

`single_source_shortest_path()` - finds the shortest path from source to all reachable nodes in graph 'G'. Here, graph 'G' maybe a directed or undirected graph.

42.2 SYNOPSIS

```
frovedis.graph.traversal.single_source_shortest_path(G, source, return_distance=False,  
                                                    verbose=0)
```

42.3 DESCRIPTION

In graph theory, the shortest path problem is the problem of finding a path between two vertices (or nodes) in a graph such that the sum of the weights of its constituent edges is minimized.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis public `single_source_shortest_path` method interface is almost same as NetworkX `single_source_shortest_path` public method interface, but it doesn't have any dependency on NetworkX. It can be used simply even if the system doesn't have NetworkX installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for `single_source_shortest_path()` on the frovedis server. Once the shortest distances are computed for the given input graph data at the frovedis server, it returns a dictionary of lists containing shortest path from source to all other nodes to the client python program.

42.3.1 Detailed Description

42.3.1.1 1. single_source_shortest_path()

Parameters

G: An instance of networkx graph or frovedis graph. The graph can be directed or undirected.

source: A positive integer parameter that specifies the starting node for the path. It must be in the range

[1, G.num_vertices].

return_distance: A boolean parameter if set to True, will return the shortest distances from source to all nodes along with traversal path. Otherwise, it will return only the traversal path. (Default: False)

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

Purpose

This method computes the shortest path between source and all other nodes reachable from source.

For example,

FILE: cit-Patents.txt

```
1 2
1 3
1 4
1 5
1 6
7 8
7 9
7 10
7 11
12 13
```

Here, the above file contains a list of edges between the nodes of graph G.

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                              create_using = nx.DiGraph())
# use verbose = 1 for detailed information
path, dist = fnx.single_source_shortest_path(frov_graph, source = 1, return_distance=True,
                                             verbose = 1)

print("Frovedis sssp traversal path: ")
print(list(path))
print("Frovedis sssp traversal distance from source: ")
print(dist)
```

Output

```
sssp computation time: 0.001 sec.
sssp res conversion time: 0.000 sec.
Frovedis sssp traversal path:
[{1: [1]}, {2: [1, 2]}, {3: [1, 3]}, {4: [1, 4]}, {5: [1, 5]}, {6: [1, 6]}]
Frovedis sssp traversal distance from source:
{1: 0.0, 2: 1.0, 3: 1.0, 4: 1.0, 5: 1.0, 6: 1.0}
```

Incase we had started with node 2 i.e source = 2, then,

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                              create_using = nx.DiGraph())
path, dist = fnx.single_source_shortest_path(frov_graph, source = 2, return_distance=True)
print("Frovedis sssp traversal path: ")
```



```
print(list(path))
print("Frovedis sssp traversal distance from source: ")
print(dist)
```

Output

```
Frovedis sssp traversal path:
[{2: [2]}, {1: [2, 1]}, {3: [2, 1, 3]}, {4: [2, 1, 4]}, {5: [2, 1, 5]}, {6: [2, 1, 6]}]
Frovedis sssp traversal distance from source:
{2: 0.0, 1: 1.0, 3: 2.0, 4: 2.0, 5: 2.0, 6: 2.0}
```

When source = 2 and return_distances = False,

```
# A directed graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int32,
                              create_using = nx.DiGraph())
# return_distances = False, by default
path= fnx.single_source_shortest_path(frov_graph, source = 2)
print("Frovedis sssp traversal path: ")
print(list(path))
```

Output

```
Frovedis sssp traversal path:
[{2: [2]}, {1: [2, 1]}, {3: [2, 1, 3]}, {4: [2, 1, 4]}, {5: [2, 1, 5]}, {6: [2, 1, 6]}]
```

Return Value

It returns a dictionary of lists containing shortest path from source to all other nodes.

42.4 SEE ALSO

- **Graph in Frovedis**
- **Breadth First Search in Frovedis**
- **Connected Components in Frovedis**
- **PageRank in Frovedis**

Chapter 43

connected_components()

43.1 NAME

`connected_components()` - generates the connected components of an undirected graph 'G'.

43.2 SYNOPSIS

```
frovedis.graph.ConnectedComponents.connected_components(G, opt_level=2, hyb_threshold=0.4,  
                                                         verbose=0, print_summary=False,  
                                                         print_limit=5)
```

43.3 DESCRIPTION

In graph theory, a component of an undirected graph is a connected subgraph that is not part of any larger connected subgraph. The components of any graph partition its vertices into disjoint sets, and are the induced subgraphs of those sets. A graph that is itself connected has exactly one component, consisting of the whole graph. Components are also sometimes called connected components.

This module provides a client-server implementation, where the client application is a normal python program. The frovedis public `connected_components` method interface is almost same as NetworkX `connected_components` public method interface, but it doesn't have any dependency on NetworkX. It can be used simply even if the system doesn't have NetworkX installed. Thus, in this implementation, a python client can interact with a frovedis server sending the required python data for training at frovedis side. Python data is converted into frovedis compatible data internally and the python ML call is linked with the respective frovedis ML call to get the job done at frovedis server.

Python side calls for `connected_components()` on the frovedis server. Once the connected components are computed for the given input graph data at the frovedis server, it returns a dictionary with keys as root-nodeid for each component and values as list of pairs of nodeid with its distance from root of the component to which the node belongs.

43.3.1 Detailed Description

43.3.1.1 1. `connected_components()`

Parameters

G: An instance of networkx graph or frovedis graph. The graph must be undirected.

opt_level: Zero or a positive integer parameter that must be in range 0 to 2. It is an optimization parameter that is used for reducing computation time while generating the connected components. (Default: 1)

- **When opt_level = 0**: this should only be used where systems have memory constraints. It is slowest.

- **When opt_level = 1**: this is fastest. It uses comparatively large amount of memory.

- **When opt_level = 2**: this is much better than '**opt_level = 0**', but slightly slower than **opt_level = 1**. It optimizes the memory usage over '**opt_level = 1**'.

hyb_threshold: A double (float64) parameter that specifies a threshold value which performs optimization in generating connected components when the number of remaining nodes to be visited becomes less than this value. It optimizes the execution time. This parameter works only with '**opt_level = 2**'. It must be within the range 0 to 1. (Default: 0.4)

For example, if it is 0.5, then optimization starts when number of remaining nodes to be visited is less than 50%.

verbose: An integer parameter specifying the log level to use. Its value is 0 by default (INFO level). But it can be set to 1 (DEBUG level) or 2 (TRACE level) for getting training time logs from frovedis server.

print_summary: A boolean parameter that specifies whether to print summary of connected components. (Default: False)

print_limit: An integer parameter that specifies the maximum number of nodes info to be printed. (Default: 5)

Purpose

This method computes connected components of an undirected graph.

For example,

```
# An undirected graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int64)
ret = fnx.connected_components(frov_graph)
for i in ret:
    print(i)
```

Output

```
{1, 2, 3, 4, 5, 6}
{7, 8, 9, 10, 11}
{12, 13}
```

When **print_summary = True** and **print_limit = 5** (by default), then,

```
# An undirected graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int64)
ret = fnx.connected_components(frov_graph, print_summary = True)
for i in ret:
    print(i)
```

Output

```

Number of connected components: 3
Root with its count of nodes in each connected component: (root_id:count)
1:6
7:5
12:2
Nodes in which cc: (node_id:root_id)
1:1
2:1
3:1
4:1
5:1
...
Nodes dist: (node:level_from_root)
1:0
2:1
3:1
4:1
5:1
...
{1, 2, 3, 4, 5, 6}
{7, 8, 9, 10, 11}
{12, 13}

```

When `print_limit = 10`, then,

```

# An undirected graph loaded from edgelist file
import numpy as np
import networkx as nx
import frovedis.graph as fnx
frov_graph = fnx.read_edgelist('input/cit-Patents_10.txt', nodetype = np.int64)
ret = fnx.connected_components(frov_graph, print_summary = True, print_limit = 10)
for i in ret:
    print(i)

```

Output

```

Number of connected components: 3
Root with its count of nodes in each connected component: (root_id:count)
1:6
7:5
12:2
Nodes in which cc: (node_id:root_id)
1:1
2:1
3:1
4:1
5:1
6:1
7:7
8:7
9:7
10:7
...
Nodes dist: (node:level_from_root)
1:0
2:1

```

```
3:1
4:1
5:1
6:1
7:0
8:1
9:1
10:1
...
{1, 2, 3, 4, 5, 6}
{7, 8, 9, 10, 11}
{12, 13}
```

Return Value

It yields an output as Sets of nodes for each connected components.

43.4 SEE ALSO

- **Graph in Frovedis**
- **Breadth First Search in Frovedis**
- **Single Source Shortest Path in Frovedis**
- **PageRank in Frovedis**

Chapter 44

FrovedisDvector

44.1 NAME

FrovedisDvector - A data structure used in modeling the in-memory dvector data of frovedis server side at client python side.

44.2 SYNOPSIS

```
class frovedis.matrix.dvector.FrovedisDvector(vec=None)
```

44.2.1 Public Member Functions

```
load (vec)  
load_numpy_array (vec)  
debug_print()  
release()
```

44.3 DESCRIPTION

FrovedisDvector is a pseudo data structure at client python side which aims to model the frovedis server side `dvector<double>` (see manual of frovedis dvector for details).

Note that the actual vector data is created at frovedis server side only. Python side FrovedisDvector contains a proxy handle of the in-memory vector data created at frovedis server, along with its size.

44.3.1 Constructor Documentation

44.3.1.1 FrovedisDvector (vec=None)

Parameters

vec: It can be any python array-like object or None. In case of None (Default), it does not make any request to server.

Purpose

This constructor can be used to construct a FrovedisDvector instance, as follows:

```
v1 = FrovedisDvector()           # empty dvector, no server request is made
v2 = FrovedisDvector([1,2,3,4]) # will load data from the given list
```

Return Type

It simply returns “self” reference.

44.3.2 Pubic Member Function Documentation

44.3.2.1 load (vec)

Parameters

vec: It can be any python array-like object (but not None).

Purpose

This function works similar to the constructor. It can be used to load a FrovedisDvector instance, as follows:

```
v = FrovedisDvector().load([1,2,3,4]) # will load data from the given list
```

Return Type

It simply returns “self” reference.

44.3.2.2 load__numpy__array (vec)

Parameters

vec: Any numpy array with values to be loaded in.

Purpose

This function can be used to load a python side numpy array data into frovedis server side dvector. It accepts a python numpy array object and converts it into the frovedis server side dvector whose proxy along size information are stored in the target FrovedisDvector object.

Return Type

It simply returns “self” reference.

44.3.2.3 size()

Purpose

It returns the size of the dvector

Return Type

An integer value containing size of the target dvector.

44.3.2.4 debug__print()

Purpose

It prints the contents of the server side distributed vector data on the server side user terminal. It is mainly useful for debugging purpose.

Return Type

It returns nothing.

44.3.2.5 release()**Purpose**

This function can be used to release the existing in-memory data at frovedis server side.

Return Type

It returns nothing.

44.3.2.6 FrovedisDvector.asDvec(vec)**Parameters**

vec: A numpy array or python array like object or an instance of FrovedisDvector.

Purpose

This static function is used in order to convert a given array to a dvector. If the input is already an instance of FrovedisDvector, then the same will be returned.

Return Type

An instance of FrovedisDvector.

Chapter 45

FrovedisCRSMatrix

45.1 NAME

FrovedisCRSMatrix - A data structure used in modeling the in-memory crs matrix data of frovedis server side at client python side.

45.2 SYNOPSIS

```
class frovedis.matrix.sparse.FrovedisCRSMatrix(mat=None)
```

45.2.1 Public Member Functions

```
load (mat)
load_scipy_matrix (mat)
load_text (filename)
load_binary (dirname)
save_text (filename)
save_binary (dirname)
debug_print()
release()
```

45.3 DESCRIPTION

FrovedisCRSMatrix is a pseudo matrix structure at client python side which aims to model the frovedis server side `crs_matrix<double>` (see manual of frovedis `crs_matrix` for details).

Note that the actual matrix data is created at frovedis server side only. Python side FrovedisCRSMatrix contains a proxy handle of the in-memory matrix data created at frovedis server, along with number of rows and number of columns information.

45.3.1 Constructor Documentation

45.3.1.1 FrovedisCRSMatrix (mat=None)

Parameters

mat: It can be a string containing filename having text data to be loaded, or any scipy sparse matrix or any python array-like object or None. In case of None (Default), it does not make any request to server.

Purpose

This constructor can be used to construct a FrovedisCRSMatrix instance, as follows:

```
mat1 = FrovedisCRSMatrix() # empty matrix, no server request is made
mat2 = FrovedisCRSMatrix("./data") # will load data from given text file
mat3 = FrovedisCRSMatrix([1,2,3,4]) # will load data from the given list
```

Return Type

It simply returns “self” reference.

45.3.2 Pubic Member Function Documentation

45.3.2.1 load (mat)

Parameters

mat: It can be a string containing filename having text data to be loaded, or any scipy sparse matrix or any python array-like object (but it can not be None).

Purpose

This works similar to the constructor.

It can be used to load a FrovedisCRSMatrix instance, as follows:

```
mat1 = FrovedisCRSMatrix().load("./data") # will load data from given text file
mat2 = FrovedisCRSMatrix().load([1,2,3,4]) # will load data from the given list
```

Return Type

It simply returns “self” reference.

45.3.2.2 load_scipy_matrix (mat)

Parameters

mat: Any scipy matrix with values to be loaded in.

Purpose

This function can be used to load a python side scipy sparse data matrix into frovedis server side crs matrix. It accepts a scipy sparse matrix object and converts it into the frovedis server side crs matrix whose proxy along with number of rows and number of columns information are stored in the target FrovedisCRSMatrix object.

Return Type

It simply returns “self” reference.

45.3.2.3 load_text (filename)

Parameters

filename: A string object containing the text file name to be loaded.

Purpose

This function can be used to load the data from a text file into the target matrix. Note that the file must be placed at server side at the given path and it should have contents stored in libSVM format, i.e., “column_index:value” at each row (see frowedis manual of make_crs_matrix_load() for more details).

Return Type

It simply returns “self” reference.

45.3.2.4 load_binary (dirname)

Parameters

dirname: A string object containing the directory name having the binary data to be loaded.

Purpose

This function can be used to load the data from the specified directory with binary data file into the target matrix. Note that the file must be placed at server side at the given path.

Return Type

It simply returns “self” reference.

45.3.2.5 save_text (filename)

Parameters

filename: A string object containing the text file name in which the data is to be saved.

Purpose

This function is used to save the target matrix as text file with the filename at the given path. Note that the file will be saved at server side at the given path.

Return Type

It returns nothing.

45.3.2.6 save_binary (dirname)

Parameters

dirname: A string object containing the directory name in which the data is to be saved as little-endian binary form.

Purpose

This function is used to save the target matrix as little-endian binary file with the filename at the given path. Note that the file will be saved at server side at the given path.

Return Type

It returns nothing.

45.3.2.7 numRows()

Purpose

It returns the number of rows in the matrix

Return Type

An integer value containing rows count in the target matrix.

45.3.2.8 numCols()**Purpose**

It returns the number of columns in the matrix

Return Type

An integer value containing columns count in the target matrix.

45.3.2.9 debug_print()**Purpose**

It prints the contents of the server side distributed matrix data on the server side user terminal. It is mainly useful for debugging purpose.

Return Type

It returns nothing.

45.3.2.10 release()**Purpose**

This function can be used to release the existing in-memory data at frovedis server side.

Return Type

It returns nothing.

45.3.2.11 FrovedisCRSMatrix.asCRS(mat)**Parameters**

mat: A scipy matrix, an instance of FrovedisCRSMatrix or any python array-like data.

Purpose

This static function is used in order to convert a given matrix to a crs matrix. If the input is already an instance of FrovedisCRSMatrix, then the same will be returned.

Return Type

An instance of FrovedisCRSMatrix.

Chapter 46

FrovedisBlockcyclicMatrix

46.1 NAME

FrovedisBlockcyclicMatrix - A data structure used in modeling the in-memory blockcyclic matrix data of frovedis server side at client python side.

46.2 SYNOPSIS

```
class frovedis.matrix.dense.FrovedisBlockcyclicMatrix(mat=None)
```

46.2.1 Overloaded Operators

```
operator= (mat)
operator+ (mat)
operator- (mat)
operator* (mat)
operator~ (mat)
```

46.2.2 Public Member Functions

```
load (mat)
load_numpy_matrix (mat)
load_text (filename)
load_binary (dirname)
save_text (filename)
save_binary (dirname)
transpose()
to_numpy_matrix ()
debug_print()
release()
```

46.3 DESCRIPTION

FrovedisBlockcyclicMatrix is a pseudo matrix structure at client python side which aims to model the frovedis server side `blockcyclic_matrix<double>` (see manual of frovedis `blockcyclic_matrix` for details).

Note that the actual matrix data is created at frovedis server side only. Python side FrovedisBlockcyclicMatrix contains a proxy handle of the in-memory matrix data created at frovedis server, along with number of rows and number of columns information.

46.3.1 Constructor Documentation

46.3.1.1 FrovedisBlockcyclicMatrix (mat=None)

Parameters

mat: It can be a string containing filename having text data to be loaded, or another FrovedisBlockcyclicMatrix instance for copy or any python array-like object or None. In case of None (Default), it does not make any request to server.

Purpose

This constructor can be used to construct a FrovedisBlockcyclicMatrix instance, as follows:

```
mat1 = FrovedisBlockcyclicMatrix() # empty matrix, no server request is made
mat2 = FrovedisBlockcyclicMatrix("./data") # will load data from given text file
mat3 = FrovedisBlockcyclicMatrix(mat2) # copy constructor
mat4 = FrovedisBlockcyclicMatrix([1,2,3,4]) # will load data from the given list
```

Return Type

It simply returns “self” reference.

46.3.2 Overloaded Operators Documentation

46.3.2.1 operator= (mat)

Parameters

mat: An existing FrovedisBlockcyclicMatrix instance to be copied.

Purpose

It can be used to copy the input matrix in the target matrix. It returns a self reference to support operator chaining.

For example,

```
m1 = FrovedisBlockcyclicMatrix([1,2,3,4])
m2 = m1 (copy operator)
m3 = m2 = m1
```

Return Type

It returns “self” reference.

46.3.2.2 operator+ (mat)

Parameters

mat: An instance of FrovedisBlockcyclicMatrix or an array-like structure.

Purpose

It can be used to perform addition between two blockcyclic matrices. If the input data is not a Frovedis-BlockcyclicMatrix instance, internally it will get converted into a FrovedisBlockcyclicMatrix instance first and then that will be added with the source matrix.

For example,

```
m1 = FrovedisBlockcyclicMatrix([1,2,3,4])
m2 = FrovedisBlockcyclicMatrix([1,2,3,4])
m3 = m2 + m1
```

Return Type

It returns the resultant matrix of the type FrovedisBlockcyclicMatrix.

46.3.2.3 operator- (mat)**Parameters**

mat: An instance of FrovedisBlockcyclicMatrix or an array-like structure.

Purpose

It can be used to perform subtraction between two blockcyclic matrices. If the input data is not a Frovedis-BlockcyclicMatrix instance, internally it will get converted into a FrovedisBlockcyclicMatrix instance first and then that will be subtracted from the source matrix.

For example,

```
m1 = FrovedisBlockcyclicMatrix([1,2,3,4])
m2 = FrovedisBlockcyclicMatrix([1,2,3,4])
m3 = m2 - m1
```

Return Type

It returns the resultant matrix of the type FrovedisBlockcyclicMatrix.

46.3.2.4 operator* (mat)**Parameters**

mat: An instance of FrovedisBlockcyclicMatrix or an array-like structure.

Purpose

It can be used to perform multiplication between two blockcyclic matrices. If the input data is not a FrovedisBlockcyclicMatrix instance, internally it will get converted into a FrovedisBlockcyclicMatrix instance first and then that will be multiplied with the source matrix.

For example,

```
m1 = FrovedisBlockcyclicMatrix([1,2,3,4])
m2 = FrovedisBlockcyclicMatrix([1,2,3,4])
m3 = m2 * m1
```

Return Type

It returns the resultant matrix of the type FrovedisBlockcyclicMatrix.

46.3.2.5 operator~ ()**Purpose**

It can be used to obtain transpose of the target matrix. If the input data is not a FrovedisBlockcyclicMatrix

instance, internally it will get converted into a FrovedisBlockcyclicMatrix instance first and then the transpose will get computed.

For example,

```
m1 = FrovedisBlockcyclicMatrix([1,2,3,4])
m2 = ~m1
```

Return Type

It returns the resultant matrix of the type FrovedisBlockcyclicMatrix.

46.3.3 Pubic Member Function Documentation

46.3.3.1 load (mat)

Parameters

mat: It can be a string containing filename having text data to be loaded, or another FrovedisBlockcyclicMatrix instance for copy or any python array-like object (but it can not be None).

Purpose

This function works similar to the constructor. It can be used to load a FrovedisBlockcyclicMatrix instance, as follows:

```
mat1 = FrovedisBlockcyclicMatrix().load("./data") # will load data from given text file
mat2 = FrovedisBlockcyclicMatrix().load(mat1)    # copy operation
mat3 = FrovedisBlockcyclicMatrix().load([1,2,3,4]) # will load data from the given list
```

Return Type

It simply returns “self” reference.

46.3.3.2 load_numpy_matrix (mat)

Parameters

mat: A numpy matrix with values to be loaded in.

Purpose

This function can be used to load a python side dense data matrix into a frovedis server side blockcyclic matrix. It accepts a numpy matrix object and converts it into the frovedis server side blockcyclic matrix whose proxy along with number of rows and number of columns information are stored in the target FrovedisBlockcyclicMatrix object.

Return Type

It simply returns “self” reference.

46.3.3.3 load_text (filename)

Parameters

filename: A string object containing the text file name to be loaded.

Purpose

This function can be used to load the data from a text file into the target matrix. Note that the file must be placed at server side at the given path.

Return Type

It simply returns “self” reference.

46.3.3.4 load_binary (dirname)

Parameters

dirname: A string object containing the directory name having the binary data to be loaded.

Purpose

This function can be used to load the data from the specified directory with binary data file into the target matrix. Note that the file must be placed at server side at the given path.

Return Type

It simply returns “self” reference.

46.3.3.5 save_text (filename)

Parameters

filename: A string object containing the text file name in which the data is to be saved.

Purpose

This function is used to save the target matrix as text file with the filename at the given path. Note that the file will be saved at server side at the given path.

Return Type

It returns nothing.

46.3.3.6 save_binary (dirname)

Parameters

dirname: A string object containing the directory name in which the data is to be saved as little-endian binary form.

Purpose

This function is used to save the target matrix as little-endian binary file with the filename at the given path. Note that the file will be saved at server side at the given path.

Return Type

It returns nothing.

46.3.3.7 transpose ()

Purpose

This function will compute the transpose of the given matrix.

Return Type

It returns the transposed blockcyclic matrix of the type FrovedisBlockcyclicMatrix.

46.3.3.8 to_numpy_matrix ()

Purpose

This function is used to convert the target blockcyclic matrix into numpy matrix.

Note that this function will request frovedis server to gather the distributed data, and send back that data in the rowmajor array form and the python client will then convert the received numpy array from frovedis server to python numpy matrix.

Return Type

It returns a two-dimensional dense numpy matrix

46.3.3.9 numRows()**Purpose**

It returns the number of rows in the matrix

Return Type

An integer value containing rows count in the target matrix.

46.3.3.10 numCols()**Purpose**

It returns the number of columns in the matrix

Return Type

An integer value containing columns count in the target matrix.

46.3.3.11 debug_print()**Purpose**

It prints the contents of the server side distributed matrix data on the server side user terminal. It is mainly useful for debugging purpose.

Return Type

It returns nothing.

46.3.3.12 release()**Purpose**

This function can be used to release the existing in-memory data at frovedis server side.

Return Type

It returns nothing.

46.3.3.13 FrovedisBlockcyclicMatrix.asBCM(mat)**Parameters**

mat: An instance of FrovedisBlockcyclicMatrix or any python array-like structure.

Purpose

This static function is used in order to convert a given matrix to a blockcyclic matrix. If the input is already an instance of FrovedisBlockcyclicMatrix, then the same will be returned.

Return Type

An instance of FrovedisBlockcyclicMatrix.

Chapter 47

pblas__wrapper

47.1 NAME

pblas__wrapper - a frovedis module provides user-friendly interfaces for commonly used pblas routines in scientific applications like machine learning algorithms.

47.2 SYNOPSIS

```
import frovedis.matrix.wrapper.PBLAS
```

47.2.1 Public Member Functions

```
PBLAS.swap (v1, v2)
PBLAS.copy (v1, v2)
PBLAS.scal (v, al)
PBLAS.axpy (v1, v2, al=1.0)
PBLAS.dot (v1, v2)
PBLAS.nrm2 (v)
PBLAS.gemv (m, v1, v2, trans=False, al=1.0, b2=0.0)
PBLAS.ger (v1, v2, m, al=1.0)
PBLAS.gemm (m1, m2, m3, trans__m1=False, trans__m2=False, al=1.0, be=0.0)
PBLAS.geadd (m1, m2, trans=False, al=1.0, be=1.0)
```

47.3 DESCRIPTION

PBLAS is a high-performance scientific library written in Fortran language. It provides rich set of functionalities on vectors and matrices. The computation loads of these functionalities are parallelized over the available processes in a system and the user interfaces of this library is very detailed and complex in nature. It requires a strong understanding on each of the input parameters, along with some distribution concepts.

Frovedis provides a wrapper module for some commonly used PBLAS subroutines in scientific applications like machine learning algorithms. These wrapper interfaces are very simple and user needs not to consider all the detailed distribution parameters. Only specifying the target vectors or matrices with some other

parameters (depending upon need) are fine. At the same time, all the use cases of a PBLAS routine can also be performed using Frovedis PBLAS wrapper of that routine.

This python module implements a client-server application, where the python client can send the python matrix data to frovedis server side in order to create blockcyclic matrix at frovedis server and then python client can request frovedis server for any of the supported PBLAS operation on that matrix. When required, python client can request frovedis server to send back the resultant matrix and it can then create equivalent python data.

The individual detailed descriptions can be found in the subsequent sections. Please note that the term “inout”, used in the below section indicates a function argument as both “input” and “output”.

47.3.1 Detailed Description

47.3.1.1 swap (v1, v2)

Parameters

v1: A FrovedisBlockcyclicMatrix with single column (inout)

v2: A FrovedisBlockcyclicMatrix with single column (inout)

Purpose

It will swap the contents of v1 and v2, if they are semantically valid and are of same length.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.2 copy (v1, v2)

Parameters

v1: A FrovedisBlockcyclicMatrix with single column (input)

v2: A FrovedisBlockcyclicMatrix with single column (output)

Purpose

It will copy the contents of v1 in v2 ($v2 = v1$), if they are semantically valid and are of same length.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.3 scal (v, al)

Parameters

v: A FrovedisBlockcyclicMatrix with single column (inout)

al: A double parameter to specify the value to which the input vector needs to be scaled. (input)

Purpose

It will scale the input vector with the provided “al” value, if it is semantically valid. On success, input vector “v” would be updated (in-place scaling).

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.4 axpy (v1, v2, al=1.0)

Parameters

v1: A FrovedisBlockcyclicMatrix with single column (input)

v2: A FrovedisBlockcyclicMatrix with single column (inout)

al: A double parameter to specify the value to which “v1” needs to be scaled (not in-place scaling) [Default: 1.0] (input/optional)

Purpose

It will solve the expression $v2 = al*v1 + v2$, if the input vectors are semantically valid and are of same length. On success, “v2” will be updated with desired result, but “v1” would remain unchanged.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.5 dot (v1, v2)

Parameters

v1: A FrovedisBlockcyclicMatrix with single column (input)

v2: A FrovedisBlockcyclicMatrix with single column (input)

Purpose

It will perform dot product of the input vectors, if they are semantically valid and are of same length. Input vectors would not get modified during the operation.

Return Value

On success, it returns the dot product result of the type double. If any error occurs, it throws an exception.

47.3.1.6 nrm2 (v)

Parameters

v: A FrovedisBlockcyclicMatrix with single column (input)

Purpose

It will calculate the norm of the input vector, if it is semantically valid. Input vector would not get modified during the operation.

Return Value

On success, it returns the norm value of the type double. If any error occurs, it throws an exception.

47.3.1.7 gemv (m, v1, v2, trans=False, al=1.0, be=0.0)

Parameters

m: A FrovedisBlockcyclicMatrix (input)

v1: A FrovedisBlockcyclicMatrix with single column (input)

v2: A FrovedisBlockcyclicMatrix with single column (inout)

trans: A boolean value to specify whether to transpose “m” or not [Default: False] (input/optional)

al: A double type value [Default: 1.0] (input/optional)

be: A double type value [Default: 0.0] (input/optional)

Purpose

The primary aim of this routine is to perform simple matrix-vector multiplication.

But it can also be used to perform any of the below operations:

$$(1) \quad v2 = al*m*v1 + be*v2$$

$$(2) \quad v2 = al*transpose(m)*v1 + be*v2$$

If *trans*=False, then expression (1) is solved. In that case, the size of “v1” must be at least the number of columns in “m” and the size of “v2” must be at least the number of rows in “m”.

If `trans=True`, then expression (2) is solved. In that case, the size of “v1” must be at least the number of rows in “m” and the size of “v2” must be at least the number of columns in “m”.

Since “v2” is used as input-output both, memory must be allocated for this vector before calling this routine, even if simple matrix-vector multiplication is required. Otherwise, this routine will throw an exception.

For simple matrix-vector multiplication, no need to specify values for the input parameters “trans”, “al” and “be” (leave them at their default values).

On success, “v2” will be overwritten with the desired output. But “m” and “v1” would remain unchanged.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.8 ger (v1, v2, m, al=1.0)

Parameters

v1: A FrovedisBlockcyclicMatrix with single column (input)

v2: A FrovedisBlockcyclicMatrix with single column (input)

m: A FrovedisBlockcyclicMatrix (inout)

al: A double type value [Default: 1.0] (input/optional)

Purpose

The primary aim of this routine is to perform simple vector-vector multiplication of the sizes “a” and “b” respectively to form an axb matrix. But it can also be used to perform the below operations:

$$m = al*v1*v2' + m$$

This operation can only be performed if the inputs are semantically valid and the size of “v1” is at least the number of rows in matrix “m” and the size of “v2” is at least the number of columns in matrix “m”.

Since “m” is used as input-output both, memory must be allocated for this matrix before calling this routine, even if simple vector-vector multiplication is required. Otherwise it will throw an exception.

For simple vector-vector multiplication, no need to specify the value for the input parameter “al” (leave it at its default value).

On success, “m” will be overwritten with the desired output. But “v1” and “v2” will remain unchanged.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.9 gemm (m1, m2, m3, trans_m1=False, trans_m2=False, al=1.0, be=0.0)

Parameters

m1: A FrovedisBlockcyclicMatrix (input)

m2: A FrovedisBlockcyclicMatrix (input)

m3: A FrovedisBlockcyclicMatrix (inout)

trans_m1: A boolean value to specify whether to transpose “m1” or not [Default: False] (input/optional)

trans_m2: A boolean value to specify whether to transpose “m2” or not [Default: False] (input/optional)

al: A double type value [Default: 1.0] (input/optional)

be: A double type value [Default: 0.0] (input/optional)

Purpose

The primary aim of this routine is to perform simple matrix-matrix multiplication.

But it can also be used to perform any of the below operations:

(1) $m3 = al*m1*m2 + be*m3$

(2) $m3 = al*transpose(m1)*m2 + be*m3$


```
(3) m3 = al*m1*transpose(m2) + be*m3
(4) m3 = al*transpose(m1)*transpose(m2) + be*m3
```

(1) will be performed, if both “trans_m1” and “trans_m2” are False.

(2) will be performed, if trans_m1=True and trans_m2 = False.

(3) will be performed, if trans_m1=False and trans_m2 = True.

(4) will be performed, if both “trans_m1” and “trans_m2” are True.

If we have four variables nrowa, nrowb, ncola, ncolb defined as follows:

```
if(trans_m1) {
    nrowa = number of columns in m1
    ncola = number of rows in m1
}
else {
    nrowa = number of rows in m1
    ncola = number of columns in m1
}

if(trans_m2) {
    nrowb = number of columns in m2
    ncolb = number of rows in m2
}
else {
    nrowb = number of rows in m2
    ncolb = number of columns in m2
}
```

Then this function can be executed successfully, if the below conditions are all true:

- (a) "ncola" is equal to "nrowb"
- (b) number of rows in "m3" is equal to or greater than "nrowa"
- (b) number of columns in "m3" is equal to or greater than "ncolb"

Since “m3” is used as input-output both, memory must be allocated for this matrix before calling this routine, even if simple matrix-matrix multiplication is required. Otherwise it will throw an exception.

For simple matrix-matrix multiplication, no need to specify the value for the input parameters “trans_m1”, “trans_m2”, “al”, “be” (leave them at their default values).

On success, “m3” will be overwritten with the desired output. But “m1” and “m2” will remain unchanged.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.3.1.10 geadd (m1, m2, trans=False, al=1.0, be=1.0)

Parameters

m1: A FrovedisBlockcyclicMatrix (input)

m2: A FrovedisBlockcyclicMatrix (input)

trans: A boolean value to specify whether to transpose “m1” or not [Default: False] (input/optional)

al: A double type value [Default: 1.0] (input/optional)

be: A double type value [Default: 1.0] (input/optional)

Purpose

The primary aim of this routine is to perform simple matrix-matrix addition. But it can also be used to perform any of the below operations:

- (1) $m2 = a1*m1 + b2*m2$
- (2) $m2 = a1*transpose(m1) + b2*m2$

If `trans=False`, then expression (1) is solved. In that case, the number of rows and the number of columns in “m1” should be equal to the number of rows and the number of columns in “m2” respectively.

If `trans=True`, then expression (2) is solved. In that case, the number of columns and the number of rows in “m1” should be equal to the number of rows and the number of columns in “m2” respectively.

If it is needed to scale the input matrices before the addition, corresponding “a1” and “b2” values can be provided. But for simple matrix-matrix addition, no need to specify values for the input parameters “trans”, “a1” and “b2” (leave them at their default values).

On success, “m2” will be overwritten with the desired output. But “m1” would remain unchanged.

Return Value

On success, it returns nothing. If any error occurs, it throws an exception.

47.4 SEE ALSO

`scalapack_wrapper`, `blockcyclic_matrix`

Chapter 48

scalapack_wrapper

48.1 NAME

scalapack_wrapper - a frovedis module provides user-friendly interfaces for commonly used scalapack routines in scientific applications like machine learning algorithms.

48.2 SYNOPSIS

```
import frovedis.matrix.wrapper.SCALAPACK
```

48.3 WRAPPER FUNCTIONS

```
SCALAPACK.getrf (m)  
SCALAPACK.getri (m, ipivPtr)  
SCALAPACK.getrs (m1, m2, ipivPtr, trans=False)  
SCALAPACK.gesv (m1, m2)  
SCALAPACK.gels (m1, m2, trans=False)  
SCALAPACK.gesvd (m, wantU=False, wantV=False)
```

48.4 DESCRIPTION

ScaLAPACK is a high-performance scientific library written in Fortran language. It provides rich set of linear algebra functionalities whose computation loads are parallelized over the available processes in a system and the user interfaces of this library is very detailed and complex in nature. It requires a strong understanding on each of the input parameters, along with some distribution concepts.

Frovedis provides a wrapper module for some commonly used ScaLAPACK subroutines in scientific applications like machine learning algorithms. These wrapper interfaces are very simple and user needs not to consider all the detailed distribution parameters. Only specifying the target vectors or matrices with some other parameters (depending upon need) are fine. At the same time, all the use cases of a ScaLAPACK routine can also be performed using Frovedis ScaLAPACK wrapper of that routine.

This python module implements a client-server application, where the python client can send the python matrix data to frovedis server side in order to create blockcyclic matrix at frovedis server and then python

client can request frovedis server for any of the supported ScaLAPACK operation on that matrix. When required, python client can request frovedis server to send back the resultant matrix and it can then create equivalent python data (see manuals for FrovedisBlockcyclicMatrix to python data conversion).

The individual detailed descriptions can be found in the subsequent sections. Please note that the term “inout”, used in the below section indicates a function argument as both “input” and “output”.

48.4.1 Detailed Description

48.4.1.1 getrf (m)

Parameters

m: A FrovedisBlockcyclicMatrix (inout)

Purpose

It computes an LU factorization of a general M-by-N distributed matrix, “m” using partial pivoting with row interchanges.

On successful factorization, matrix “m” is overwritten with the computed L and U factors. Along with the return status of native scalapack routine, it also returns the proxy address of the node local vector “ipiv” containing the pivoting information associated with input matrix “m” in the form of GetrfResult. The “ipiv” information will be useful in computation of some other routines (like getri, gets etc.)

Return Value

On success, it returns the object of the type GetrfResult as explained above. If any error occurs, it throws an exception explaining cause of the error.

48.4.1.2 getri (m, ipivPtr)

Parameters

m: A FrovedisBlockcyclicMatrix (inout)

ipiv: A long object containing the proxy of the ipiv vector (from GetrfResult) (input)

Purpose

It computes the inverse of a distributed square matrix using the LU factorization computed by getrf(). So in order to compute inverse of a matrix, first compute it’s LU factor (and ipiv information) using getrf() and then pass the factored matrix, “m” along with the “ipiv” information to this function.

On success, factored matrix “m” is overwritten with the inverse (of the matrix which was passed to getrf()) matrix. “ipiv” will be internally used by this function and will remain unchanged.

For example,

```
res = SCALAPACK.getrf(m)          // getting LU factorization of "m"
SCALAPACK.getri(m,res.ipiv()) // "m" will have inverse of the initial value
```

Return Value

On success, it returns the exit status of the scalapack routine itself. If any error occurs, it throws an exception explaining cause of the error.

48.4.1.3 gets (m1, m2, ipiv, trans=False)

Parameters

m1: A FrovedisBlockcyclicMatrix (input)

m2: A FrovedisBlockcyclicMatrix (inout)

ipiv: A long object containing the proxy of the ipiv vector (from GetrfResult) (input)

trans: A boolean value to specify whether to transpose “m1” [Default: False] (input/optional)

Purpose

It solves a real system of distributed linear equations, $AX=B$ with a general distributed square matrix (A) using the LU factorization computed by `getrf()`. Thus before calling this function, it is required to obtain the factored matrix “m1” (along with “ipiv” information) by calling `getrf()`.

For example,

```
res = SCALAPACK.getrf(m1) // getting LU factorization of "m1"
SCALAPACK.getrs(m1,m2,res.ipiv())
```

If `trans=False`, the linear equation $AX=B$ is solved.

If `trans=True`, the linear equation $\text{transpose}(A)X=B$ ($A'X=B$) is solved.

The matrix “m2” should have number of rows \geq the number of rows in “m1” and at least 1 column in it.

On entry, “m2” contains the distributed right-hand-side (B) of the equation and on successful exit it is overwritten with the distributed solution matrix (X).

Return Value

On success, it returns the exit status of the scalapack routine itself. If any error occurs, it throws an exception explaining cause of the error.

48.4.1.4 gesv (m1, m2)

Parameters

m1: A FrovedisBlockcyclicMatrix (inout)

m2: A FrovedisBlockcyclicMatrix (inout)

Purpose

It solves a real system of distributed linear equations, $AX=B$ with a general distributed square matrix, “m1” by computing its LU factors internally. This function internally computes the LU factors and ipiv information using `getrf()` and then solves the equation using `getrs()`.

The matrix “m2” should have number of rows \geq the number of rows in “m1” and at least 1 column in it.

On entry, “m1” contains the distributed left-hand-side square matrix (A), “m2” contains the distributed right-hand-side matrix (B) and on successful exit “m1” is overwritten with its LU factors, “m2” is overwritten with the distributed solution matrix (X).

Return Value

On success, it returns the exit status of the scalapack routine itself. If any error occurs, it throws an exception explaining cause of the error.

48.4.1.5 gels (m1, m2, trans=False)

Parameters

m1: A FrovedisBlockcyclicMatrix (input)

m2: A FrovedisBlockcyclicMatrix (inout)

trans: A boolean value to specify whether to transpose “m1” [Default: False] (input/optional)

Purpose

It solves overdetermined or underdetermined real linear systems involving an M-by-N distributed matrix (A) or its transpose, using a QR or LQ factorization of (A). It is assumed that distributed matrix (A) has full rank.

If `trans=False` and $M \geq N$: it finds the least squares solution of an overdetermined system.
 If `trans=False` and $M < N$: it finds the minimum norm solution of an underdetermined system.
 If `trans=True` and $M \geq N$: it finds the minimum norm solution of an underdetermined system.
 If `trans=True` and $M < N$: it finds the least squares solution of an overdetermined system.

The matrix “m2” should have number of rows $\geq \max(M, N)$ and at least 1 column.

On entry, “m1” contains the distributed left-hand-side matrix (A) and “m2” contains the distributed right-hand-side matrix (B). On successful exit, “m1” is overwritten with the QR or LQ factors and “m2” is overwritten with the distributed solution matrix (X).

Return Value

On success, it returns the exit status of the scalapack routine itself. If any error occurs, it throws an exception explaining cause of the error.

48.4.1.6 gesvd (m, wantU=False, wantV=False)

Parameters

m: A FrovedisBlockcyclicMatrix (inout)

wantU: A boolean value to specify whether to compute U matrix [Default: False] (input)

wantV: A boolean value to specify whether to compute V matrix [Default: False] (input)

Purpose

It computes the singular value decomposition (SVD) of an M-by-N distributed matrix.

On entry “m” contains the distributed matrix whose singular values are to be computed.

If `wantU = wantV = False`, then it computes only the singular values in sorted order, so that $\text{sval}(i) \geq \text{sval}(i+1)$. Otherwise it also computes U and/or V (left and right singular vectors respectively) matrices.

On successful exit, the contents of “m” is destroyed (internally used as workspace).

Return Value

On success, it returns the object of the type GesvdResult containing the singular values and U and V components (based on the requirement) along with the exit status of the native scalapack routine. If any error occurs, it throws an exception explaining cause of the error.

48.5 SEE ALSO

blockcyclic_matrix, pblas_wrapper, arpack_wrapper, getrf_result, gesvd_result

Chapter 49

getrf__result

49.1 NAME

getrf__result - a structure to model the output of frovedis wrapper of scalapack getrf routine.

49.2 SYNOPSIS

```
import frovedis.matrix.results.GetrfResult
```

49.2.1 Public Member Functions

```
release()  
ipiv()  
stat()
```

49.3 DESCRIPTION

GetrfResult is a client python side pseudo result structure containing the proxy of the in-memory scalapack getrf result (node local ipiv vector) created at frovedis server side.

49.3.1 Public Member Function Documentation

49.3.1.1 release()

Purpose

This function can be used to release the in-memory result component (ipiv vector) at frovedis server.

Return Type

It returns nothing.

49.3.1.2 ipiv()

Purpose

This function returns the proxy of the node_local “ipiv” vector computed during getrf calculation. This value will be required in other scalapack routine calculation, like getri, getrs etc.

Return Type

A long value containing the proxy of ipiv vector.

49.3.1.3 stat()

Purpose

This function returns the exit status of the scalapack native getrf routine on calling of which the target result object was obtained.

Return Type

It returns an integer value.

Chapter 50

gesvd__result

50.1 NAME

gesvd__result - a structure to model the output of frovedis singular value decomposition methods.

50.2 SYNOPSIS

```
import frovedis.matrix.results.GesvdResult
```

50.2.1 Public Member Functions

```
to__numpy__results()
save(svec, umat=None, vmat=None)
save__binary(svec, umat=None, vmat=None)
load (svec, umat=None, vmat=None, mtype='B')
load__binary (svec, umat=None, vmat=None, mtype='B')
debug__print()
release()
stat()
getK()
```

50.3 DESCRIPTION

GesvdResult is a python side pseudo result structure containing the proxies of the in-memory SVD results created at frovedis server side. It can be used to convert the frovedis side SVD result to python equivalent data structures.

50.3.1 Public Member Function Documentation

50.3.1.1 to__numpy__results()

Purpose

This function can be used to convert the frovedis side SVD results to python numpy result structures.

If U and V both are computed, it returns: (numpy matrix, numpy array, numpy matrix) indicating (umatrix, singular vector, vmatrix).

When U is calculated, but not V, it returns: (numpy matrix, numpy array, None)

When V is calculated, but not U, it returns: (None, numpy array, numpy matrix)

When neither U nor V is calculated, it returns: (None, numpy array, None)

Return Type

It returns a tuple as explained above.

50.3.1.2 save(svec, umat=None, vmat=None)

Parameters

svec: A string object containing name of the file to save singular vectors as text data. (mandatory)

umat: A string object containing name of the file to save umatrix as text data. (optional)

vmat: A string object containing name of the file to save vmatrix as text data. (optional)

Purpose

This function can be used to save the result values in different text files at server side. If saving of U and V components are not required, “umat” and “vmat” can be None, but “svec” should have a valid filename.

Return Type

It returns nothing.

50.3.1.3 save_binary(svec, umat=None, vmat=None)

Parameters

svec: A string object containing name of the file to save singular vectors as binary data. (mandatory)

umat: A string object containing name of the file to save umatrix as binary data. (optional)

vmat: A string object containing name of the file to save vmatrix as binary data. (optional)

Purpose

This function can be used to save the result values in different files as little-endian binary data at server side. If saving of U and V components are not required, “umat” and “vmat” can be None, but “svec” should have a valid filename.

Return Type

It returns nothing.

50.3.1.4 load(svec, umat=None, vmat=None, mtype='B')

Parameters

svec: A string object containing name of the file from which to load singular vectors as text data for the target result. (mandatory)

umat: A string object containing name of the file from which to load umatrix as text data for the target result. (optional)

vmat: A string object containing name of the file from which to load vmatrix as text data for the target result. (optional)

mtype: A character value, can be either ‘B’ or ‘C’. (optional)

Purpose

This function can be used to load the result values in different text files at server side. If loading of U and V components are not required, “umat” and “vmat” can be None, but “svec” should have a valid filename.

If mtype = ‘B’ and umat/vmat is to be loaded, then they will be loaded as blockcyclic matrices at server side.

If `mtype = 'C'` and `umat/vmat` is to be loaded, then they will be loaded as colmajor matrices at server side.

Return Type

It returns nothing.

50.3.1.5 `load_binary(svec, umat=None, vmat=None, mtype='B')`

Parameters

svec: A string object containing name of the file from which to load singular vectors as binary data for the target result. (mandatory)

umat: A string object containing name of the file from which to load umatrix as binary data for the target result. (optional)

vmat: A string object containing name of the file from which to load vmatrix as binary data for the target result. (optional)

mtype: A character value, can be either 'B' or 'C'. (optional)

Purpose

This function can be used to load the result values in different little-endian binary files at server side. If loading of U and V components are not required, "umat" and "vmat" can be None, but "svec" should have a valid filename.

If `mtype = 'B'` and `umat/vmat` is to be loaded, then they will be loaded as blockcyclic matrices at server side.

If `mtype = 'C'` and `umat/vmat` is to be loaded, then they will be loaded as colmajor matrices at server side.

Return Type

It returns nothing.

50.3.1.6 `debug_print()`

Purpose

This function can be used to print the result components at server side user terminal. This is useful in debugging purpose.

Return Type

It returns nothing.

50.3.1.7 `release()`

Purpose

This function can be used to release the in-memory result components at frovedis server.

Return Type

It returns nothing.

50.3.1.8 `stat()`

Purpose

This function returns the exit status of the scalapack native gesvd routine on calling of which the target result object was obtained.

Return Type

An integer value.

50.3.1.9 getK()**Purpose**

This function returns the number of singular values computed.

Return Type

An integer value.

Chapter 51

DataFrame Introduction

51.1 NAME

DataFrame - here refers to a Frovedis dataframe. It is a two-dimensional, size-mutable, potentially heterogeneous tabular data structure.

51.2 DESCRIPTION

Dataframe is a two-dimensional data structure, i.e., data is aligned in a tabular fashion in rows and columns. This data structure also contains labeled axes (rows and columns). Arithmetic operations align on both row and column labels. It can be thought of as a dict-like container for Series objects or it can be thought of as an SQL table or a spreadsheet data representation.

Features of DataFrame:

- Columns can be of different types.
- DataFrame is mutable i.e. the number of rows and columns can be increased or decreased.
- DataFrame supports indexing and labeled columns name.
- Supports arithmetic operations on rows and columns.

Frovedis contains dataframe implementation over a client server architecture (which uses Vector Engine on server side to perform fast computations) where pandas dataframe will be converted to frovedis dataframe and then will be used to perform dataframe related operations.

In frovedis, currently dataframe can only support atomic data i.e. a column in a frovedis dataframe can only have primitive type values such as string, integer, float, double, boolean).

51.2.1 Detailed description

51.2.1.1 1. Creating frovedis DataFrame instance

a) Using DataFrame constructor

`DataFrame(df = None)`

Parameters

df: A pandas DataFrame instance or pandas Series instance. (Default: None)

When this parameter is not None (specified explicitly), it will load the pandas dataframe (or series) to perform conversion into frovedis dataframe.

Purpose

It is used to create a Frovedis dataframe from the given pandas dataframe or series.

Constructing frovedis DataFrame from pandas DataFrame:

For example,

```
# a pandas dictionary having key and values pairs
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe from key value pair
pdf1 = pd.DataFrame(peopleDF)

# creating frovedis dataframe
fdf1 = fd.DataFrame(pdf1)

# display created frovedis dataframe
fdf1.show()
```

Output

index	Age	Country	Ename	isMale
0	29	USA	Michael	0
1	30	England	Andy	0
2	27	Japan	Tanaka	0
3	19	France	Raul	0
4	31	Japan	Yuta	1

Return Value

It returns self reference.

Attributes

columns: It returns a list of string having the column labels of the frovedis dataframe.

For example,

```
# fdf1 is same from above example
print(fdf1.columns)
```

Output

```
['Age', 'Country', 'Ename', 'isMale']
```

count: It returns an integer value having the number of rows in the frovedis dataframe

For example,

```
# fdf1 is same from above example
print(fdf1.count)
```

Output

```
5
```

dtypes: It returns the dtypes of all columns in the frovedis dataframe. It returns a Series object with the data type of each column.

For example,

```
# fdf1 is same from above example
print(fdf1.dtypes)
```

Output

```
Age          int64
Country      object
Ename        object
isMale       bool
dtype: object
```

ndim: It returns an integer representing the number of axes / array dimensions.

For example,

```
# fdf1 is same from above example
print(fdf1.ndim)
```

Output

```
2
```

shape: It returns a tuple representing the dimensions of the frovedis dataframe in form: **(nrows, ncols)**

For example,

```
# fdf1 is same from above example
print(fdf1.shape)
```

Output

```
(5, 4)
```

values: It returns a numpy representation of the frovedis dataframe.

For example,

```
# fdf1 is same from above example
print(fdf1.values)
```

Output

```
[[ 'Michael' '29' 'USA' '0']
 [ 'Andy'    '30' 'England' '0']
 [ 'Tanaka'  '27' 'Japan'  '0']
 [ 'Raul'    '19' 'France'  '0']
 [ 'Yuta'    '31' 'Japan'   '1']]
```

b) By loading data using read_csv() method

`read_csv(filepath_or_buffer, sep = ',', delimiter = None, header = "infer", names = None, index_col = None, usecols = None, squeeze = False, prefix = None, mangle_dupe_cols = True, dtype = None, na_values = None, verbose = False, comment = None, low_memory = True, rows_to_see = 1024, separate_mb = 1024)`

Parameters

filepath_or_buffer: It accepts any valid string value that contains the name of the file to access as parameter. The string path can be a URL as well. In case, a file is to be read, then the pathname can be absolute path or relative path to the current working directory of the file to be opened.

sep: It accepts string value as paramter that specifies delimiter to use. (Default: ',')

Currently, it supports only single character delimiter.

When it is None (if specified explicitly), it will use ',' as delimiter.

delimiter: It accepts string value as paramter. It is an alias for ‘sep’ parameter. (Default: None)
When it is None (not specified explicitly), it will set ‘,’ as the delimiter.

header: It accepts integer or string value that specifies the row number(s) to use as the column names, and the start of the data. (Default: ‘infer’)

Currently, frovedis supports only ‘infer’, 0 or None as values for this parameter.

The default behavior is to ‘infer’ the column names:

- **If no column names are passed**, then the behavior is identical to **header = 0** and column names are inferred from the first line of the file.
- **If column names are passed explicitly**, then the behavior is identical to **header = None** and column names are inferred from the explicitly specified list.

names: It accepts array like input as parameter that specifies the list of column names to use. (Default: None)

When it is None (not specified explicitly), it will get the column names from the first line of the file. Otherwise, it will override with the given list of column names. Since, **header = 0 or None is only supported in frovedis**, the first line of the file will be used as column names. **Duplicates in this list of column names are not allowed.**

index_col: It accepts boolean and integer type values as parameter. It specifies the column(s) to use as the row labels of the frovedis dataframe. **Currently, multi-index is not supported in frovedis.** (Default: None)

- If boolean type value is passed, then currently it can only be set as False. Also, when **index_col = False**, it can be used to force to not use the first column as the index. **Also, ‘names’ parameter must be provided.**
- If integer type value is passed, then it must be in the range **(0, nCols - 1)**, where **nCols** is the number of columns present in the input file.

usecols: It accepts currently a list-like input of integer values as parameter. It specifies the subset of the columns to be used while loading data into frovedis dataframe. (Default: None)

When it is None (not specified explicitly), it will use all columns while creating frovedis dataframe.

Also, if the column names are provided and do not match the number of columns in the input file, then the list of names must be same in length as the list of ‘usecols’ parameter values. Otherwise, it raises an exception.

squeeze: An unused parameter. (Default: False)

prefix: It accepts a string value as parameter. It specifies the prefix which needs to be added to the column numbers. (Default: None)

For example, ‘X’ for X0, X1, X2, ...

This is an optional paramter which will be used when column names are not provided and header is None (no header).

When it is None (not specified explicitly), it will create list of numbers as column labels of length N - 1, where ‘N’ is the number of columns in the input file.

mangle_dupe_cols: It accepts boolean value as parameter that specifies the action taken for duplicate columns present in the input file. (Default: True)

- **When set to True (not specified explicitly) and if duplicate columns are present**, they will be specified as ‘X’, ‘X.1’, ... ‘X.N’, rather than ‘X’... ‘X’.
- **When set to False and duplicate columns are present**, it will raise an exeption (**not supported in frovedis yet**).

dtype: It accepts datatype or dictionary of columns which is used for explicitly setting type for data or columns in frovedis dataframe. (Default: None)

For boolwan type, frovedis supports:

True/False, On/OFF, 0/1, yes/No, y/n

User must provide this, then frovedis can convert to boolean type. **Make sure user provides correct dtype , otherwise it will cause issues while creating frovedis dataframe.**

na_values: It accepts scalar value or array-like input as parameter that specifies the additional strings to recognize as missing values (NA/NaN). Currently, it doesn't accept dictionary as value for this parameter. (Default: None)

When it is None (not specified explicitly), it will interpret the list of following values as missing values:

```
['null', 'NULL', 'nan', '-nan', 'NaN', '-NaN', 'NA', 'N/A', 'n/a']
```

In case we want some other value to be interpreted as missing value, it is explicitly provided.

verbose: It accepts a boolean values as parameter that specifies the log level to use. Its value is **False by default (for INFO mode)** and it can be set to **True (for DEBUG mode)**. This is used for getting the loading time logs from frovedis server. It is useful for debugging purposes. (Default: False)

comment: This is an unused parameter. (Default: None)

low_memory: It accepts boolean value as parameter that specifies if the dataframe is to be loaded chunk wise. The chunk size then would be specified by the 'separate_mb' parameter. (Default: True)

rows_to_see: It accepts integer value as parameter. It is useful when creating frovedis dataframe by loading from large files. While using big files, only some rows would be used to infer the data types of columns. This parameter specifies the number of such rows which will be used to infer data types. (Default: 1024)

separate_mb: It accepts double (float64) value as parameter. It is a memory optimization parameter. (Default: 1024)

- If **low_memory = True**, then internally frovedis dataframes of size '**separate_mb**' will be loaded separately and would be combined later into a single frovedis dataframe.
- If **low_memory = False**, then complete frovedis dataframe will be loaded at once.

Purpose

This method reads a comma-separated values (csv) file into frovedis dataframe.

Note:- While loading file into frovedis dataframe, the file should not contain blank line at beginning.

The parameters: "squeeze" and "comment" are simply kept in to to make the interface uniform to the pandas read_csv() method. These are not used internally within the frovedis implementation.

Using read_csv() to load data from given csv file path into frovedis dataframe:

File: numbers.csv

```
10,10.23,F,0
12,12.20,nan,0
13,34.90,D,1
15,100.12,A,2
```

In order to load csv file will into frovedis dataframe,

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv")
df.show()
```

Output

```
index    10      10.23   F      0
0         12      12.2   NULL   0
```

```

1      13      34.8999 D      1
2      15      100.12 A      2

```

Here, by default it uses ‘,’ delimiter while loading csv data into frovedis dataframe.

Also, since no column names were provided and header = ‘infer’, so the newly created frovedis dataframe contains column names inferred from the first line of input file.

Using read_csv() with header = None and no column name is provided:

For example,

```

import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", header = None)
df.show()

```

Output

```

index  0      1      2      3
0      10     10.23  F      0
1      12     12.2   NULL   0
2      13     34.8999 D      1
3      15     100.12 A      2

```

Here, it creates list of column numbers which is used as column names in newly created frovedis dataframe.

Using read_csv() with header = None. Also, column names are provided:

For example,

```

import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", names = ['one', 'two', 'three', 'four'], header = None)
df.show()

```

Output

```

index  one     two     three  four
0      10     10.23  F      0
1      12     12.2   NULL   0
2      13     34.8999 D      1
3      15     100.12 A      2

```

Here, given list of names will be used as column names in the newly generated frovedis dataframe.

Also, the input file contained no duplicate column names in its header.

Using read_csv() with prefix = True:

For example,

```

import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", header = None, prefix = "X")
df.show()

```

Output

```

index  X0      X1      X2      X3
0      10     10.23  F      0
1      12     12.2   NULL   0
2      13     34.8999 D      1
3      15     100.12 A      2

```

Here, ‘prefix’ is an optional parameter which will be in use when header = None and names = None. It will use column numbers by default when prefix = None.

To load specific columns in frovedis dataframe, usecols parameter will be used:

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", names=['one', 'two', 'three', 'four'], usecols = [1,2])
df.show()
```

Output

index	two	three
0	10.23	F
1	12.2	NULL
2	34.8999	D
3	100.12	A

Using read_csv() with dtype parameter:

File: numbers.csv

```
10,10.23,E,TRUE
12,12.20,nan,TRUE
13,34.90,D,FALSE
15,100.12,A,FALSE
```

In above data, **TRUE/FALSE** in python can be converted to boolean (0/1) type values using dtype parameter.

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", names = ['one', 'two', 'three', 'four'],
                dtype = {'four': bool})
df.show()
```

Output

index	one	two	three	four
0	10	10.23	E	1
1	12	12.2	NULL	1
2	13	34.8999	D	0
3	15	100.12	A	0

In order to use second column as index, index_col parameter will be used:

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", names = ['one', 'two', 'three', 'four'], index_col = 1)
df.show()
```

Output

two	one	three	four
10.23	10	F	0
12.2	12	NULL	0
34.8999	13	D	1
100.12	15	A	2

When index_col = False, it will force to not use the first column as the index:

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", names = ['one', 'two', 'three', 'four'], index_col = False)
df.show()
```

Output

index	one	two	three	four
0	10	10.23	E	0
1	12	12.2	NULL	0
2	13	34.8999	D	1
3	15	100.12	A	2

Using `mangle_dupe_cols = True`, when input file contains duplicate columns in header:

File: numbers_dupl.csv

```
score,score,grade,pass
11.2,10.23,F,0
21.2,12.20,nan,0
43.6,34.90,D,1
75.1,100.12,A,1
```

The input file contains duplicate column labels. To make sure while loading data into frovedis dataframe, `mangle_dupe_cols` will be set as `True`.

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers_dupl.csv",mangle_dupe_cols = True)
df.show()
```

Output

index	score	score.1	grade	pass
0	10	10.23	F	0
1	12	12.2	NULL	0
2	13	34.8999	D	1
3	15	100.12	A	1

Using `read_csv()` with `na_values` parameter:

File: numbers.csv

```
11,10.23,E,0,1
12,12.20,F,0,1
10,34.90,D,1,0
15,10,A,10,0
10,10.23,B,0,1
10,10.23,B,0,1
```

When using `na_values = None` (by default), it will interpret the values `nan`, `Nan`, `NULL`, etc. as missing values (explained in parameter description)

```
# demo for read_csv() with na_values = None by default
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", na_values = None)
df.show()
```

Output

index	11	10.23	E	0	1
0	12	12.2	F	0	1

1	10	34.8999	D	1	0
2	15	10	A	10	0
3	10	10.23	B	0	1
4	10	10.23	B	0	1

When list of strings are provided to be set as missing values:

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", na_values = ['10','1'])
df.show()
```

Output

index	11	10.23	E	0	1
0	12	12.2	F	0	NULL
1	NULL	34.8999	D	NULL	0
2	15	NULL	A	NULL	0
3	NULL	10.23	B	0	NULL
4	NULL	10.23	B	0	NULL

It will interpret the list of values ['10','1'] as missing values (explicitly provided) while loading input file into frovedis dataframe.

When a string is to be set as missing value:

For example,

```
import frovedis.dataframe as fdf
df = fdf.read_csv("./input/numbers.csv", na_values = '10')
df.show()
```

Output

index	11	10.23	E	0	1
0	12	12.2	F	0	1
1	NULL	34.8999	D	1	0
2	15	NULL	A	NULL	0
3	NULL	10.23	B	0	1
4	NULL	10.23	B	0	1

It will interpret the string '10' as missing value (explicitly provided) while loading input file into frovedis dataframe.

Return Value

It returns a frovedis DataFrame instance.

51.2.1.2 2. Destructing the frovedis DataFrame instance

Using release() method

Purpose

This method acts like a destructor.

It is used to release dataframe pointer from server heap and it resets all its attributes to None.

Releasing dataframe pointers from frovedis:

For example,

```
# fdf1 is same from above example
fdf1.release()
```

Return Value

It returns nothing.

51.2.1.3 3. Public Member Functions of DataFrame

DataFrame provides a lot of utilities to perform various operations. For simplicity we have categorized them into three lists: *Basic functions*, *Application*, *groupby functions*, *Aggregate functions* and *Binary operator functions*.

51.2.1.3.1 a) List of Basic Functions Basic functions are further categorized into two sub parts - *conversion and sorting functions* and *selection and combination functions*.

In the basic functions, we will discuss the common and essential functionalities of dataframe like conversion of dataframes, sorting of data, selection of specified data and combining two or more data.

51.2.1.3.1.1 Conversion Functions: Conversion functions are essential part of DataFrame which are basically used to perform conversions to other types and to narrow down the data as per specification.

1. **asDF()** - Returns a Frovedis DataFrame after suitable conversion from other DataFrame types.
2. **to_dict()** - Convert the dataframe to a dictionary.
3. **to_numpy()** - Converts a frovedis dataframe to numpy array.
4. **to_pandas()** - Returns a pandas dataframe object from frovedis dataframe.
5. **to_frovedis_rowmajor_matrix()** - Converts a frovedis dataframe to FrovedisRowmajorMatrix.
6. **to_frovedis_colmajor_matrix()** - Converts a frovedis dataframe to FrovedisColmajorMatrix.
7. **to_frovedis_crs_matrix()** - Converts a frovedis dataframe to FrovedisCRSMatrix.
8. **to_frovedis_crs_matrix_using_info()** - Converts a frovedis dataframe to FrovedisCRSMatrix provided an info object of df_to_sparse_info class.

51.2.1.3.1.2 Sorting Functions: Sorting functions are essential part of DataFrame.

1. **nlargest()** - Return the first n rows ordered by columns in descending order.
2. **nsmallest()** - Return the first n rows ordered by columns in ascending order.
3. **sort()** - Sort by the values on a column.
4. **sort_index()** - Sort dataframes according to index.
5. **sort_values()** - Sort by the values along either axis.

51.2.1.3.1.3 Generic Functions: DataFrame provides various facilities to easily select and combine together specified values and support join/merge operations.

1. **add_index()** - Adds index column to the dataframe in-place.

2. **append()** - Union of dataframes according to rows.
3. **apply()** - Apply a function along an axis of the DataFrame.
4. **astype()** - Cast a selected column to a specified dtype.
5. **between()** - Filters rows according to the specified bound over a single column at a time.
6. **copy()** - Make a copy of this object's indices and data.
7. **countna()** - Count NA values for each column/row.
8. **describe()** - Generate descriptive statistics.
9. **drop()** - Drop specified labels from rows or columns.
10. **drop_duplicates()** - Return DataFrame with duplicate rows removed.
11. **dropna()** - Remove missing values.
12. **fillna()** - Fill NA/NaN values using specified values.
13. **filter()** - Subset the dataframe rows or columns according to the specified index labels.
14. **get_index_loc()** - Returns integer location, slice or boolean mask for specified value in index column.
15. **head()** - Return the first n rows.
16. **insert()** - Insert column into DataFrame at specified location.
17. **isna()** - Detect missing values.
18. **isnull()** - Is an alias of `isna()`.
19. **join()** - Join columns of another DataFrame.
20. **merge()** - Merge dataframes according to specified parameters.
21. **rename()** - Used to rename column.
22. **rename_index()** - Renames index field (inplace).
23. **reset_index()** - Reset the index.
24. **set_index()** - Set the DataFrame index using existing columns.
25. **show()** - Displays the selected dataframe values on console.
26. **tail()** - Return the last n rows.
27. **update_index()** - Updates/sets index values.

51.2.1.3.2 b) List of Aggregate Functions Aggregate functions of dataframe help to perform computations on the specified values and helps with efficient summarization of data. The calculated values gives insight into the nature of potential data.

1. **agg()** - Aggregate using the specified functions and columns.
2. **cov()** - Returns the pairwise covariance of columns, excluding missing values.
3. **mad()** - Returns the mean absolute deviation of the values over the requested axis.
4. **max()** - Returns the maximum of the values over the requested axis.
5. **mean()** - Returns the mean of the values over the requested axis.
6. **median()** - Returns the median of the values over the requested axis.
7. **min()** - Returns the minimum of the values over the requested axis.
8. **mode()** - Returns the mode(s) of each element along the selected axis.
9. **sem()** - Returns the unbiased standard error of the mean over requested axis.
10. **std()** - Returns the sample standard deviation over requested axis.
11. **sum()** - Returns the sum of the values over the requested axis.
12. **var()** - Returns unbiased variance over requested axis.

51.2.1.3.3 c) List of Math Functions DataFrame has methods for carrying out binary operations like `add()`, `sub()`, etc and related functions like `radd()`, `rsub()`, etc. for carrying out reverse binary operations.

1. **abs()** - Return a DataFrame with absolute numeric value of each element.
2. **add()** - Get addition of dataframe and other specified value. It is equivalent to `dataframe + other`.
3. **div()** - Get floating division of dataframe and other specified value. It is equivalent to `dataframe / other`.
4. **floordiv()** - Get integer division of dataframe and other specified value. It is equivalent to `dataframe // other`.
5. **mod()** - Get modulo of dataframe and other specified value. It is equivalent to `dataframe % other`.
6. **mul()** - Get multiplication of dataframe and other specified value. It is equivalent to `dataframe * other`.
7. **pow()** - Get exponential power of dataframe and other specified value. It is equivalent to `dataframe ** other`.
8. **sub()** - Get subtraction of dataframe and other specified value. It is equivalent to `dataframe - other`.
9. **truediv()** - Get floating division of dataframe and other specified value. It is equivalent to `dataframe / other`.
10. **radd()** - Get addition of other specified value and dataframe. It is equivalent to `other + dataframe`.

11. **rdiv()** - Get floating division of other specified value and dataframe. It is equivalent to other / dataframe.
12. **rfloordiv()** - Get integer division of other specified value and dataframe. It is equivalent to other // dataframe.
13. **rmod()** - Get modulo of other specified value and dataframe. It is equivalent to other % dataframe.
14. **rmul()** - Get multiplication of other specified value and dataframe. It is equivalent to other * dataframe.
15. **rpow()** - Get exponential power of other specified value and dataframe. It is equivalent to other ** dataframe.
16. **rsub()** - Get subtraction of other specified value and dataframe. It is equivalent to other - dataframe.
17. **rtruediv()** - Get floating division of other specified value and dataframe. It is equivalent to other / dataframe.

51.2.1.4 Indexing in frovedis Dataframe

Indexing means simply selecting particular rows and columns of data from a dataframe. Indexing can also be known as Subset Selection.

There are a lot of ways to pull the rows and columns from a dataframe. There are some indexing method which help in getting an element from a dataframe. These indexing methods appear very similar but behave very differently. **Currently, frovedis supports only one type of indexing.** It is:

- **Dataframe.[]**: This function also known as indexing operator. It helps in filtering rows and columns from a dataframe. Also, it returns a pandas dataframe as output.

Indexing using []

Indexing operator is used to refer to the square brackets following an object.

In order to select a single column, we simply put the name of the column in-between the brackets.

Selecting a single column:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a pandas dataframe from key value pair
pdf2 = pd.DataFrame({ "Last Bonus": [5, 2, 2, 4],
                      "Bonus": [5, 2, 2, 4],
                      "Last Salary": [58, 59, 63, 58],
                      "Salary": [60, 60, 64, 59]
                      }, index= ["John", "Marry", "Sam", "Jo"]
)

# creating frovedis dataframe
fdf1 = fdf.DataFrame(pdf2)

# display created frovedis dataframe
fdf1.show()

print('selecting single column: ')
fdf1['Bonus'].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
John	5	5	58	60
Marry	2	2	59	60
Sam	2	2	63	64
Jo	4	4	58	59

selecting single column:

index	Bonus
John	5
Marry	2
Sam	2
Jo	4

Note:- Frovedis also supports use of attribute operators to filter/select column(s) in dataframe

In previous example, **Bonus** column can also be selected from the dataframe as follows:

For example,

```
fdf1.Bonus
```

This returns a FrovedisColumn instance.

To select multiple columns:

For example,

```
# selecting multiple columns
fdf1[['Bonus','Salary']].show()
```

Output

index	Bonus	Salary
John	5	60
Marry	2	60
Sam	2	64
Jo	4	59

Here, list of columns are passed in the indexing operator.

Filtering dataframe using slice operation with row numbers:

For example,

```
fdf1[1:2].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
Marry	2	2	59	60

Filtering dataframe using slice operation with row labels:

For example,

```
fdf1['John':'Sam'].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
Marry	2	2	59	60

Filtering can be done with help of attribute operators:

For example,

```
# filtering data using given condition
fdf1[fdf1.Bonus == 2].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
Marry	2	2	59	60
Sam	2	2	63	64

For example,

```
# filtering data using '>' operator
fdf1[fdf1.Bonus > 2].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
John	5	5	58	60
Jo	4	4	58	59

For example,

```
# filtering data using '<' operator
fdf1[fdf1.Bonus < 5].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
Marry	2	2	59	60
Sam	2	2	63	64
Jo	4	4	58	59

For example,

```
# filtering data using '!=' operator
fdf1[fdf1.Bonus != 2].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
John	5	5	58	60
Jo	4	4	58	59

For example,

```
# using '&' operation to filter data
fdf1[(fdf1.Bonus == 5) & (fdf1.Salary == 60)].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
John	5	5	58	60

For example,

```
# using '|' operation to filter data
fdf1[(fdf1.Bonus == 5) | (fdf1.Salary == 60)].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
John	5	5	58	60
Marry	2	2	59	60

For example,

```
# using '~' operation to filter data
fdf1[~(fdf1.Bonus == 5)].show()
```

Output

index	Last Bonus	Bonus	Last Salary	Salary
Marry	2	2	59	60
Sam	2	2	63	64
Jo	4	4	58	59

51.3 SEE ALSO

- **DataFrame - Generic Functions**
- **DataFrame - Conversion Functions**
- **DataFrame - Sorting Functions**
- **DataFrame - Math Functions**
- **DataFrame - Aggregate Functions**

Chapter 52

DataFrame Generic Functions

52.1 NAME

DataFrame Generic Functions - it contains list of all generally used functions with frovedis dataframe.

52.1.1 DESCRIPTION

Frovedis dataframe provides various functions which are generally/frequently used with it to perform operations:

- Selecting / Filtering / Modifying data in frovedis dataframe.
- Combining two frovedis dataframes.
- Dropping data from rows or columns in a frovedis dataframe.
- Display frovedis dataframe (full/partially)

52.1.2 Public Member Functions

1. `append(other, ignore_index = False, verify_integrity = False, sort = False)`
2. `apply()`
3. `astype(dtype, copy = True, errors = 'raise', check_bool_like_string = False)`
4. `between(left, right, inclusive="both")`
5. `copy(deep = True)`
6. `countna()`
7. `describe()`
8. `drop(labels = None, axis = 0, index = None, columns = None, level = None, inplace = False, errors = 'raise')`
9. `drop_duplicates(subset = None, keep = 'first', inplace = False, ignore_index = False)`
10. `dropna(axis = 0, how = 'any', thresh = None, subset = None, inplace = False)`
11. `fillna(value = None, method = None, axis = None, inplace = False, limit = None, downcast = None)`
12. `filter(items = None, like = None, regex = None, axis = None)`
13. `get_index_loc(value)`
14. `head(n = 5)`
15. `insert(loc, column, value, allow_duplicates = False)`
16. `isna()`
17. `isnull()`
18. `join(right, on, how = 'inner', lsuffix = '_left', rsuffix = '_right', sort = False, join_type = 'bcast')`

```

19. merge(right, on = None, how = 'inner', left_on = None, right_on = None,
        left_index = False, right_index = False, sort = False, suffixes = ('_x', '_y'),
        copy = True, indicator = False, join_type = 'bcast')
20. rename(columns, inplace = False)
21. reset_index(drop = False, inplace = False)
22. set_index(keys, drop = True, append = False, inplace = False, verify_integrity = False)
23. show()
24. tail(n = 5)

```

52.1.3 Detailed Description

52.1.3.1 1. DataFrame.append(other, ignore_index = False, verify_integrity = False, sort = False)

Parameters

other: It accepts a Frovedis DataFrame instance or a Pandas DataFrame instance or a list of Frovedis DataFrame instances which are to be appended.

ignore_index: It accepts a boolean type parameter. If True, old index axis is ignored and a new index axis is added with values 0 to n - 1, where n is the number of rows in the DataFrame. (Default: False)

verify_integrity: It accepts a boolean type as parameter. If it is set to True, it checks 'index' label for duplicate entries before appending and when there are duplicate entries in the DataFrame, it doesn't append. Otherwise duplicate entries in the 'index' label will be appended. It will also append duplicate entries when 'ignore_index' = True. (Default: False)

sort: It accepts a boolean type as parameter. It sorts the columns, if the columns of the given DataFrame and other DataFrame are not aligned. (Default: False)

Purpose

It is used to append entries of dataframe at the end of another dataframe. The columns of other DataFrame instance that are not in the calling DataFrame instance are added as new columns.

Creating two frovedis dataframes:

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame([[1, 2], [3, 4]], columns = list('AB'), index = ['x', 'y'])

# create a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display frovedis dataframe
fd_df1.show()

# create another pandas dataframe
pd_df2 = pd.DataFrame([[5, 6], [7, 8]], columns = list('AB'), index = ['x', 'y'])

# create another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)

# display other frovedis dataframe
fd_df2.show()

```

Output

index	A	B
x	1	2
y	3	4

index	A	B
x	5	6
y	7	8

Appending fd_df2 to fd_df1:

For example,

```
# example of using append method with default values
# appending fd_df2 to fd_df1
fd_df1.append(fd_df2).show()
```

Output

index	A	B
x	1	2
y	3	4
x	5	6
y	7	8

With ignore_index set to True:

For example,

```
# append() demo With ignore_index = True
fd_df1.append(fd_df2, ignore_index = True).show()
```

Output

index	A	B
0	1	2
1	3	4
2	5	6
3	7	8

Creating two frovedis dataframes with no common columns:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame([[1, 2], [3, 4]], columns = list('AB'), index = ['x', 'y'])

# create a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display frovedis dataframe
fd_df1.show()

# create another pandas dataframe
pd_df2 = pd.DataFrame([[5, 6], [7, 8]], columns = list('CD'), index = ['x', 'y'])

# create another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)
```

```
# display other frovedis dataframe
fd_df2.show()
```

Output

```
index  A      B
x      1      2
y      3      4
```

```
index  C      D
x      5      6
y      7      8
```

Append fd_df2 to fd_df1 when there are no common columns in frovedis dataframes:

For example,

```
fd_df1.append(fd_df2).show()
```

Output

```
index  A      B      D      C
x      1      2      NULL    NULL
y      3      4      NULL    NULL
x      NULL    NULL    6      5
y      NULL    NULL    8      7
```

Creating two frovedis dataframes where both have few duplicate entries:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame([[1, 2], [3, 4]], columns=list('AB'), index=['x', 'y'])

# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display a frovedis dataframe
fd_df1.show()

# creating another pandas dataframe
pd_df2 = pd.DataFrame([[1, 2], [7, 8]], columns=list('AB'), index=['u', 'v'])

# creating another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)

# display other frovedis dataframe
fd_df2.show()
```

Output

```
index  A      B
x      1      2
y      3      4
```

```
index  A      B
```


u	1	2
v	7	8

Appending fd_df2 to fd_df1 with verify_integrity = True:

For example,

```
fd_df1.append(fd_df2, verify_integrity = True).show()
```

Output

index	A	B
x	1	2
y	3	4
u	1	2
v	7	8

Creating two frovedis dataframes to perform append and using sort=True:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame([[1, 2], [3, 4]], columns=list('YZ'), index=['x', 'y'])

# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display a frovedis dataframe
fd_df1.show()

# creating another pandas dataframe
pd_df2 = pd.DataFrame([[1, 2], [7, 8]], columns=list('AB'), index=['x', 'y'])

# creating another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)

# display other frovedis dataframe
fd_df2.show()
```

Output

index	Y	Z
x	1	2
y	3	4

index	A	B
x	1	2
y	7	8

Appending fd_df2 with fd_df1 with sort = True:

For example,

```
fd_df1.append(fd_df2, sort = True).show()
```

Output

index	A	B	Y	Z
-------	---	---	---	---

x	NULL	NULL	1	2
y	NULL	NULL	3	4
x	1	2	NULL	NULL
y	7	8	NULL	NULL

Return Value

It returns a new Frovedis DataFrame consisting of the rows of original DataFrame object and the rows of other DataFrame object.

52.1.3.2 2. DataFrame.apply(func, axis = 0, raw = False, result_type = None, args = (), **kwds)

Parameters

func: Names of functions to be applied on the data. The input to be used with the function must be a frovedis DataFrame instance having atleast one numeric column.

Accepted combinations for this parameter are:

- A string function name such as 'max', 'min', etc.
 - list of functions and/or function names, For example, ['max', 'mean'].
 - dictionary with keys as column labels and values as function name or list of such functions.
- For Example, {'Age': ['max', 'min', 'mean'], 'Ename': ['count']}

axis: It accepts an integer as parameter. It is used to decide whether to perform aggregate operation along the columns or rows. (Default: 0)

raw: It accepts boolean as parameter. When set to True, the row/column will be passed as an ndarray. (Default: False)

result_type: It accepts string object as parameter. It specifies how the result will be returned. (Default: None)

These only act when **axis = 1 (columns)**:

- **expand** : list-like results will be turned into columns.
- **reduce** : returns a Series if possible rather than expanding list-like results. This is the opposite of 'expand'.
- **broadcast** : results will be broadcast to the original shape of the DataFrame, the original index and columns will be retained.

The default behaviour (None) depends on the return value of the applied function. List-like results will be returned as a Series of those. However if the apply function returns a Series these are expanded to columns.

args: Positional arguments to pass to 'func'. (Default: ())

****kwds**: This is an unused parameter.

Purpose

Apply a function along an axis of the DataFrame.

The parameter: ****kwds** is simply kept in to make the interface uniform to the pandas DataFrame.apply(). This is not used anywhere within the frovedis implementation.

It actually uses pandas.apply() internally.

Basically, it converts the pandas dataframe into frovedis dataframe and after performing apply(), it returns the result back as pandas dataframe.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf
```

```
# a dictionary
```

```

peopleDF = {
    'Name':['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age':[27, 24, 22, 32, 33, 36, 27, 32],
    'City':['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification':['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Apply function along the rows (by default) and where func is string name:

For example,

```
print(fdf1.apply('max'))
```

Output

```

Name          Princi
Age              36
City          Nagpur
Qualification   Phd
Score           52
dtype: object

```

Apply function along the columns and where func is string name:

For example,

```
print(fdf1.apply('max', axis = 1))
```

Output

```

index
0    27.0
1    34.0
2    35.0
3    45.0
4    33.0
5    50.0

```

```
6    52.0
7    32.0
dtype: float64
```

Apply multiple functions along the rows (by default) and where func is a list of functions:

For example,

```
print(fdf1.apply(['max', 'min']))
```

Output

	Name	Age	City	Qualification	Score
max	Princi	36	Nagpur	Phd	52.0
min	Abhi	22	Allahabad	B.Tech	23.0

Return Value

1. If only one ‘func’ provided:

- It returns a pandas Series instance with numeric column(s) only, after aggregation function is completed.

2. If more than one ‘func’ provided:

- It returns a pandas DataFrame instance with numeric column(s) only, after aggregation function is completed.

52.1.3.3 3. DataFrame.astype(dtype, copy = True, errors = ‘raise’, check_bool_like_string = False)

Parameters

dtype: It accepts a string, numpy.dtype or a dict of column labels to cast entire DataFrame object to same type or one or more columns to column-specific types.

copy: It accepts a boolean parameter and returns a new DataFrame object when it is True. Currently this parameter will always return a copy. The original DataFrame is not modified. (Default: True)

errors: This is an unused parameter. (Default: ‘raise’)

check_bool_like_string: A boolean parameter which when set to True will cast string columns having boolean like case-insensitive strings (True, False, yes, No, On, Off, Y, N, T, F) to boolean columns. (Default: False)

Purpose

It cast an entire Frovedis DataFrame or selected columns of Frovedis DataFrame to the specified dtype.

The parameter: “errors” is simply kept in to make the interface uniform to the pandas DataFrame.astype(). This is not used anywhere within the frovedis implementation.

Creating a frovedis DataFrame from pandas DataFrame and display all column dtypes:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame({'col1': [1, 2], 'col2': [3, 4]})

# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df)

# displaying a frovedis dataframe dtype
```

```
print(fd_df1.dtypes)
```

Output

```
col1    int64
col2    int64
dtype: object
```

Cast all columns to int32 type:

For example,

```
fd_df2 = fd_df1.astype('int32')
```

```
# display dataframe after conversion
print(fd_df2.dtypes)
```

Output

```
col1    int32
col2    int32
dtype: object
```

Creating a new frovedis DataFrame and display all column dtypes:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# converting to pandas dataframe
pd_df = pd.DataFrame(peopleDF)

# converting to frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display the datatype of object
print(fd_df.dtypes)
```

Output

```
Ename    object
Age       int64
Country   object
isMale    bool
dtype: object
```

Cast ‘Age’ column to int32 type using a dictionary:

For example,

```
print(fd_df.astype({'Age': 'int32'}).dtypes)
```

Output

```
Ename      object
Age        int32
Country    object
isMale     int32
dtype: object
```

Creating a new frovedis DataFrame and using `check_bool_like_string` parameter:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': ['F', 'No', 'Off', False, 'Y']
}

# converting to pandas dataframe
pd_df = pd.DataFrame(peopleDF)

# converting to frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display the dataframes
fd_df.show()

# display the datatype
print(fd_df.dtypes)
```

Output

```
index  Ename  Age  Country isMale
0      Michael  29    USA      F
1      Andy    30  England  No
2      Tanaka  27    Japan  Off
3      Raul    19   France False
4      Yuta    31    Japan   Y
```

```
Ename      object
Age        int64
Country    object
isMale     object
dtype: object
```

With `check_bool_like_string = True`:

For example,

```
# astype() demo with check_bool_like_string = True
fd_df.astype({'isMale':'bool'}, check_bool_like_string = True).show()

# display the datatype
print(fd_df.astype({'isMale':'bool'}, check_bool_like_string = True).dtypes)
```

Output

index	Ename	Age	Country	isMale
0	Michael	29	USA	0
1	Andy	30	England	0
2	Tanaka	27	Japan	0
3	Raul	19	France	0
4	Yuta	31	Japan	1

```

Ename      object
Age        int64
Country    object
isMale     bool
dtype: object

```

Return Value

It returns a new DataFrame instance with dtype converted as specified.

52.1.3.4 4. DataFrame.between(left, right, inclusive = “both”)

Parameters

left: It accepts scalar values as parameter.

right: It accepts scalar values as parameter.

inclusive: It accepts string object as parameter tha specifies which boundaries to include, whether to set bounds as open or closed. (Default: ‘both’)

- ‘left’: left boundary value is included.
- ‘right’: right boundary value is included.
- ‘both’: boundary values are included.
- ‘neither’: boundary values are excluded.

Purpose

This method performs filtering of rows according to the specified bound over a single column at a time.

Currently, this method filters data for numeric column data only.

Creating a frovedis DataFrame from pandas DataFrame:

For example,

```

import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

```

```
# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Using `between` function where both boundaries are provided:

For example,

```
# between() demo used with given column of dataframe
# Also, both boundaries are included
fdf1['Score'].between(40,50)
```

Unlike pandas, it returns an instance of `dfoperator` (**present in frovedis only**) which contains masked boolean data as result of `between` operation. In order to unmask the data, `to_mask_array()` must be used.

```
fdf1['Score'].between(40,50).to_mask_array
```

Output

```
[False False False  True False  True False False]
```

This returns a boolean array containing `True` wherever the corresponding element is between the boundary values 'left' and 'right'. The missing values are treated as `False`.

Also, this method can be used in the form of given expression below:

For example,

```
# between() demo used with given column of dataframe
# Also, both boundaries are included
res = fdf1[fdf1['Score'].between(40,50)]
print(res)
```

Output

index	Name	Age	City	Qualification	Score
3	Princi	32	Kannuaj	Phd	45
5	Anuj	36	Kanpur	B.Tech	50

It can also be expressed as follows:

For example,

```
print(fdf1[fdf1.Score.between(40,50)])
```

Output

index	Name	Age	City	Qualification	Score
3	Princi	32	Kannuaj	Phd	45
5	Anuj	36	Kanpur	B.Tech	50

With inclusive='left':

For example,

```
# between() demo and only left boundary included
print(fdf1['Score'].between(40,50,inclusive = 'left').to_mask_array())
```

Output

```
[False False False  True False False False False]
```

With 'inclusive='right':

For example,

```
# between() demo and only right boundary included
print(fdf1['Score'].between(40,50,inclusive = 'right').to_mask_array())
```

Output

```
[False False False  True False  True False False]
```

With 'inclusive='neither':

For example,

```
# between() demo and both boundaries excluded
print(fdf1['Score'].between(40,50,inclusive = 'neither').to_mask_array())
```

Output

```
[False False False  True False False False False]
```

Return Value

It returns a dfoperator instance.

52.1.3.5 5. DataFrame.copy(deep = True)**Parameters**

deep: A boolean parameter to decide the type of copy operation. (Default: True)

When it is True (not specified explicitly), it creates a deep copy i.e. the copy includes copy of the original DataFrame's data and the indices. **Currently this parameter doesn't support shallow copy (deep = False).**

Purpose

It creates a deep copy of the Frovedis DataFrame object's indices and data.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe
pd_df = pd.DataFrame(peopleDF)
```

```
#creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

print('displaying original dataframe object')
fd_df.show()
```

Output

```
displaying original dataframe object
index  Ename  Age    Country isMale
0      Michael 29     USA      0
1      Andy   30    England  0
2      Tanaka 27     Japan   0
3      Raul   19    France   0
4      Yuta   31     Japan   1
```

Creating a deep copy:

For example,

```
fd_df_copy = fd_df.copy()

print('displaying copied dataframe object')
fd_df_copy.show()
```

Output

```
displaying copied dataframe object
index  Ename  Age    Country isMale
0      Michael 29     USA      0
1      Andy   30    England  0
2      Tanaka 27     Japan   0
3      Raul   19    France   0
4      Yuta   31     Japan   1
```

In order to check changes done in copy are reflected in original, we can do the following:

For example,

```
# changing column Age to Age2 for the copied object
fd_df_copy.rename({'Age': 'Age2', inplace = True})

print('displaying copied dataframe object')
fd_df_copy.show()
```

Output

```
displaying copied dataframe object
index  Ename  Age2    Country isMale
0      Michael 29     USA      0
1      Andy   30    England  0
2      Tanaka 27     Japan   0
3      Raul   19    France   0
4      Yuta   31     Japan   1
```

NOTE: changes are reflected only in copied DataFrame instance but not in original DataFrame instance.

Return Value

It returns a deep copy of the DataFrame instance of the same type.

52.1.3.6 6. DataFrame.countna(axis = 0)**Parameters**

axis: It accepts an integer or string object as parameter. It is used to decide whether to count missing values along the indices or by column labels. (Default: 0)

- 0 or 'index': count missing values along the indices.

- 1 or 'columns': count missing values along the columns.

Purpose

It counts number of missing values in the given axis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Counting number of missing values along the rows:

For example,

```
# countna() demo
fdf1.countna().show()
```

Output

```

index    count
Name     0
Age      0
City     0
Qualification    0
Score     2

```

Counting number of missing values along the columns:

For example,

```

# countna() demo using axis = 1
fdf1.countna(axis = 1).show()

```

Output

```

index    count
0         0
1         0
2         0
3         0
4         1
5         0
6         0
7         1

```

Return Value

It returns a frovedis DataFrame instance.

52.1.3.7 7. DataFrame.describe()

Purpose

It generates descriptive statistics. Descriptive statistics include count, mean, median, etc, excluding missing values.

Currently, it only numeric fields are returned.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```

import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

```

```
# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Describing a DataFrame. Only numeric fields are returned:

For example,

```
print(fdf1.describe())
```

Output

	Age	Score
count	8.000000	6.000000
mean	29.125000	39.833333
median	29.500000	40.000000
var	23.553571	123.766667
mad	4.125000	9.166667
std	4.853202	11.125047
sem	1.715866	4.541781
sum	233.000000	239.000000
min	22.000000	23.000000
max	36.000000	52.000000

Return Value

It returns a pandas DataFrame instance with the result of the specified aggregate operation.

52.1.3.8 8. DataFrame.drop(labels = None, axis = 0, index = None, columns = None, level = None, inplace = False, errors = 'raise')

Parameters

labels: It takes an integer or string type as argument. It represents column labels and integer represent index values of rows to be dropped. If any of the label is not found in the selected axis, it will raise an exception. (Default: None)

When it is None (not specified explicitly), 'index' or 'columns' parameter must be provided. Otherwise it will drop specific rows of the DataFrame.

axis: It accepts an integer type parameter where the value indicates the direction of drop operation according to below conditions:

- 0 : The corresponding index labels will be dropped.
- 1 : The corresponding columns labels will be dropped.

index: It accepts an integer or string object as parameter. It is equivalent to dropping indices along the axis = 0. (Default: None)

When it is None (not specified explicitly), 'columns' or 'labels' must be provided.

columns: It accepts a string object or list of strings. It is equivalent to dropping columns along the axis = 1. (Default: None)

When it is None (not specified explicitly), 'index' or 'labels' must be provided.

level: This is an unused parameter. (Default: None)

inplace: It accepts a boolean as a parameter. It return a copy of DataFrame instance by default but when explicitly set to True, it performs operation on original DataFrame. (Default: False)

errors: This is an unused parameter. (Default: 'raise')

Purpose

It is used to drop specified labels from rows or columns.

Rows or columns can be removed by specifying label names and corresponding axis, or by specifying index or column names.

Creating a frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

pd_df = pd.DataFrame(np.arange(12).reshape(3, 4), columns = ['A', 'B', 'C', 'D'])

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df).

# display frovedis dataframe
fd_df.show()
```

Output

index	A	B	C	D
0	0	1	2	3
1	4	5	6	7
2	8	9	10	11

Drop columns using axis parameter:

For example,

```
# drop() demo with label and axis parameter
fd_df.drop(labels = ['B', 'C'], axis = 1).show()
```

Output

index	A	D
0	0	3
1	4	7
2	8	11

Other way to drop columns:

For example,

```
# drop() demo with columns parameter
fd_df.drop(columns = ['B', 'C']).show()
```

Output

index	A	D
0	0	3
1	4	7

```
2      8      11
```

Drop a row by index:

For example,

```
fd_df.drop(labels = [0, 1]).show()
```

Output

index	A	B	C	D
2	8	9	10	11

Return Value

- It returns a new of Frovedis DataFrame having remaining entries.
- It returns None when parameter 'inplace' = True.

52.1.3.9 9. DataFrame.drop_duplicates(subset = None, keep = 'first', inplace = False, ignore_index = False)

Parameters

subset: It accepts a string object or a list of strings which only consider certain columns for identifying duplicates. (Default: None)

When it is None (not specified explicitly), it will consider all of the columns.

keep: It accepts a string object which is used to determine which duplicates values to keep. (Default: 'first')

1. 'first' : Drop duplicates except for the first occurrence.
2. 'last' : Drop duplicates except for the last occurrence.

inplace: It accepts a boolean as a parameter. It return a copy of DataFrame instance by default but when explicitly set to True, it performs operation on original DataFrame. (Default: False)

ignore_index: It accepts a boolean type parameter. If True, old index axis is ignored and a new index axis is added with values 0 to n - 1, where n is the number of rows in the DataFrame. (Default: False)

Purpose

It is used to remove duplicate rows.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# Consider dataset containing ramen rating.
pd_df = pd.DataFrame({
    'brand': ['Yum Yum', 'Yum Yum', 'Indomie', 'Indomie', 'Indomie'],
    'style': ['cup', 'cup', 'cup', 'pack', 'pack'],
    'rating': [4, 4, 3.5, 15, 5]})

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
fd_df.show()
```

Output

index	brand	style	rating
0	Yum Yum	cup	4
1	Yum Yum	cup	4
2	Indomie	cup	3.5
3	Indomie	pack	15
4	Indomie	pack	5

Drop duplicates along the rows (by default):

For example,

```
# drop_duplicates() demo
fd_df.drop_duplicates()
```

Output

index	brand	style	rating
0	Yum Yum	cup	4
2	Indomie	cup	3.5
3	Indomie	pack	15
4	Indomie	pack	5

By default, it removes duplicate rows based on all columns

To remove duplicates on specific column(s), use subset parameter:

For example,

```
# drop_duplicates() demo with subset parameter
fd_df.drop_duplicates(subset = ['brand']).show()
```

Output

index	brand	style	rating
0	Yum Yum	cup	4
2	Indomie	cup	3.5

To remove duplicates and keep last occurrences, use keep parameter:

For example,

```
# drop_duplicates() demo with parameters: subset and keep = 'last'
fd_df.drop_duplicates(subset = ['brand', 'style'], keep = 'last').show()
```

Output

index	brand	style	rating
1	Yum Yum	cup	4
2	Indomie	cup	3.5
4	Indomie	pack	5

Return Value

- It returns a new of Frovedis DataFrame having remaining entries.
- It returns None when parameter 'inplace' = True.

52.1.3.10 10. DataFrame.dropna(axis = 0, how = 'any', thresh = None, subset = None, inplace = False)

Parameters

axis: It accepts an integer value that can be 0 or 1. This parameter is used to determine whether rows or

columns containing missing values are to be removed. (Default: 0)

- 0 : Drop rows which contain missing values.
- 1 : Drop columns which contain missing values.

how: It accepts a string object to determine if row or column is removed from DataFrame, when we have at least one 'NaN' or all 'NaN'. (Default: 'any')

- 'any' : If any NaN values are present, drop that row or column.
- 'all' : If all values are NaN, drop that row or column.

thresh: It accepts an integer as parameter which is the number of NaN values required for rows/columns to be dropped. (Default: None)

subset: It accepts a python ndarray. It is the name of the labels along the other axis which is being considered.

For example, if you are dropping rows, then these would be a list of columns. (Default: None)

inplace: This parameter accepts a boolean value. When it is set to True, then it performs operation on the original Frovedis DataFrame object itself otherwise operation is performed on a new Frovedis DataFrame object. (Default: False)

Purpose

It is used to remove missing values from the Frovedis DataFrame.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame({"name": ['Alfred', 'Batman', 'Catwoman'],
                        "toy": [np.nan, 'Batmobile', 'Bullwhip'],
                        "born": [np.nan, "1940-04-25", np.nan]})

# creating a frovedis dataframe from pandas dataframe
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
print("Before dropping nan values")
fd_df.show()
```

Output

```
Before dropping nan values
index   name      toy      born
0      Alfred    NULL    NULL
1      Batman    Batmobile  1940-04-25
2      Catwoman   Bullwhip   NULL
```

Drop the rows where at least one missing value is present:

For example,

```
print("After dropping nan values")
fd_df.dropna().show()
```

Output

After dropping nan values

index	name	toy	born
1	Batman	Batmobile	1940-04-25

Drop the columns where at least one missing value is present:

For example,

```
# display frovedis dataframe
print("Before dropping nan values")
fd_df.show()
```

```
print("After dropping nan values")
fd_df.dropna(axis=1)
```

Output

Before dropping nan values

index	name	toy	born
0	Alfred	NULL	NULL
1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

After dropping nan values

index	name
0	Alfred
1	Batman
2	Catwoman

Drop the rows where all elements are missing:

For example,

```
# display frovedis dataframe
print("Before dropping nan values")
fd_df.show()
```

```
# drop the rows where all elements are missing (how='all')
fd_df.dropna(how='all').show()
```

Output

Before dropping nan values

index	name	toy	born
0	NULL	NULL	NULL
1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

index	name	toy	born
1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

Keep only the rows with at least 2 non-NA values:

For example,

```
# display frovedis dataframe
print("Before dropping nan values")
fd_df.show()
```

```
# example of using drop with parameter 'thresh'
# dropna() demo with only rows with at least 2 non-NA values (thresh = 2) to keep
fd_df.dropna(thresh=2).show()
```

Output

Before dropping nan values

index	name	toy	born
0	Alfred	NULL	NULL
1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

index	name	toy	born
1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

NOTE: Since 0th index had 2 missing values, so it was dropped, but other rows are not dropped.

To specify the columns to look for missing values:

For example,

```
# display frovedis dataframe
print("Before dropping nan values")
fd_df.show()

# to drop missing values only from specified columns
fd_df.dropna(subset=[1, 2], axis = 1).show()
```

Output

Before dropping nan values

index	name	toy	born
0	Alfred	NULL	NULL
1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

index	name	toy
0	Alfred	NULL
1	Batman	Batmobile
2	Catwoman	Bullwhip

To keep the dataframe with valid entries in the same variable:

For example,

```
# display frovedis dataframe
print("Before dropping nan values")
fd_df.show()

# dropna() demo with inplace = True
fd_df.dropna(inplace=True)
fd_df.show()
```

Output

Before dropping nan values

index	name	toy	born
0	Alfred	NULL	NULL

1	Batman	Batmobile	1940-04-25
2	Catwoman	Bullwhip	NULL

index	name	toy	born
1	Batman	Batmobile	1940-04-25

Return Value

- If `inplace = False`, it returns a new Frovedis DataFrame with NA entries dropped.
- If `inplace = True`, it returns `None`.

52.1.3.11 11. DataFrame.fillna(value = None, method = None, axis = None, inplace = False, limit = None, downcast = None)

Parameters

value: It accepts a numeric type as parameter which is used to replace all NA values (e.g. 0, NULL). (Default: `None`)

When it is `None` (not specified explicitly), it must be NaN value or numeric, non-numeric value otherwise it will raise an exception.

method: This is an unused parameter. (Default: `None`)

axis: It accepts an integer or string object as parameter. It decides the axis along which missing values will be filled. (Default: `None`)

- 0 or 'index': Operation will be carried out on rows. **Currently only axis = 0 is supported in Frovedis DataFrame.**

inplace: This parameter accepts a boolean value. When it is set to `True`, then it performs operation on the original Frovedis DataFrame object itself otherwise operation is performed on a new Frovedis DataFrame object. (Default: `False`)

limit: This is an unused parameter. (Default: `None`)

downcast: This is an unused parameter. (Default: `None`)

Purpose

It replaces NA/NaN values with the specified value provided in 'value' parameter in the Frovedis DataFrame.

The parameters: "method", "limit" and "downcast" are simply kept in to make the interface uniform to the pandas DataFrame.fillna().

This is not used anywhere within the frovedis implementation.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame([[np.nan, 2, np.nan, 0],
                      [3, 4, np.nan, 1],
                      [np.nan, np.nan, np.nan, 5],
                      [np.nan, 3, np.nan, 4]],
                      columns=list("ABCD"))

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
fd_df.show()
```

Output

index	A	B	C	D
0	NULL	2	NULL	0
1	3	4	NULL	1
2	NULL	NULL	NULL	5
3	NULL	3	NULL	4

Replace all missing values with 0:

For example,

```
fd_df.fillna(0).show()
```

Output

index	A	B	C	D
0	0	2	0	0
1	3	4	0	1
2	0	0	0	5
3	0	3	0	4

Replace all missing values with 1 and keep the dataframe in the same variable:

For example,

```
# replace all NaN elements with -1s and inplace = True
fd_df.fillna(-1, inplace = True)
```

```
# display after modifying the original object
fd_df.show()
```

Output

index	A	B	C	D
0	NULL	2	NULL	0
1	3	4	NULL	1
2	NULL	NULL	NULL	5
3	NULL	3	NULL	4

index	A	B	C	D
0	-1	2	-1	0
1	3	4	-1	1
2	-1	-1	-1	5
3	-1	3	-1	4

Return Value

- It returns a Frovedis DataFrame object with missing values replaced when 'inplace' parameter is False.
- It returns None when 'inplace' parameter is set to True.

52.1.3.12 12. DataFrame.filter(items = None, like = None, regex = None, axis = None)

Parameters

items: It accepts a list of string as parameter. It filters only those labels which are mentioned. (Default: None)

When it is None (not specified explicitly), 'like' or 'regex' must be provided.

like: It accepts a string object parameter. It keeps the column labels if 'like in label == True'. (Default: None)

When it is None (not specified explicitly), 'items' or 'regex' must be provided.

regex: It accepts a regular expression as a string parameter. It keeps the column labels if 're.search(regex, label) == True'. (Default: None)

When it is None (not specified explicitly), 'items' or 'like' must be provided.

axis: It accepts an integer or string type parameter. It specifies the axis on which filter operation will be performed. (Default: None)

- 0 or 'index' : The corresponding index labels will be filtered. **Currently this is not supported in Frovedis.**

- 1 or 'columns': The corresponding columns labels will be filtered.

When it is None (not specified explicitly), it will filter along axis = 1 or 'columns'.

Purpose

It is used to filter the DataFrame according to the specified column labels.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

pd_df = pd.DataFrame(np.array([[1, 2, 3], [4, 5, 6]]),
                      index=['mouse', 'rabbit'],
                      columns=['one', 'two', 'three'])
fd_df = fdf.DataFrame(pd_df)
```

```
# display the frovedis dataframe
fd_df.show()
```

Output

index	one	two	three
mouse	1	2	3
rabbit	4	5	6

Select columns by name:

For example,

```
# filter() demo with items parameter
# select columns by name
fd_df.filter(items = ['one', 'three']).show()
```

Output

index	one	three
mouse	1	3
rabbit	4	6

Select columns by regular expression:

For example,

```
fd_df.filter(regex='e$', axis=1).show()
```

Output

index	one	three
mouse	1	3
rabbit	4	6

The output displayed only those columns whose label ends with 'e'.

Select column containing 'hre':

For example,

```
fd_df.filter(like='hre', axis=1)
```

Output

```
index    three
mouse     3
rabbit     6
```

Return Value

It returns a new Frovedis DataFrame instance with the column labels that matches the given conditions.

52.1.3.13 13. DataFrame.get_index_loc(value)

Parameters

value: It accepts an integer or string parameter. It is the index value whose location is to be determined.

Purpose

It provides integer location, slice or boolean mask for requested label.

It is present only in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

pd_df = pd.DataFrame(peopleDF)
pd_df.index = ['a', 'b', 'c', 'd', 'e']
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
fd_df.show()
```

Output

```
index  Ename  Age  Country  isMale
a      Michael  29    USA      0
b       Andy   30  England  0
c     Tanaka   27   Japan   0
d       Raul   19  France   0
e       Yuta   31   Japan   1
```

Get index location when there are unique indices:

For example,

```
# getting index location of 'd' index value
fd_df.get_index_loc('d').show()
```

Output

3

Creating a new frovedis DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe
pd_df = pd.DataFrame(peopleDF)
pd_df.index = ['a', 'a', 'd', 'd', 'e']

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display a frovedis dataframe
fd_df.show()
```

Output

index	Ename	Age	Country	isMale
a	Michael	29	USA	0
a	Andy	30	England	0
d	Tanaka	27	Japan	0
d	Raul	19	France	0
e	Yuta	31	Japan	1

Get index location when there are consecutive duplicate indices:

For example,

```
# getting index location of 'd' index value
fd_df.get_index_loc('d').show()
```

Output

```
slice(2, 4, None)
```

Creating another frovedis DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
```



```

        'isMale': [False, False, False, False, True]
    }
# creating a pandas dataframe
pd_df = pd.DataFrame(peopleDF)
pd_df.index = ['a', 'b', 'c', 'd', 'a']

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display a frovedis dataframe
fd_df.show()

```

Output

index	Ename	Age	Country	isMale
a	Michael	29	USA	0
b	Andy	30	England	0
c	Tanaka	27	Japan	0
d	Raul	19	France	0
a	Yuta	31	Japan	1

Get index location when there are duplicate indices randomly:

For example,

```

# getting index location of 'a' index value
fd_df.get_index_loc('a').show()

```

Output

```
[ True False False False  True]
```

Return Value

It returns the following values:

1. **integer:** when there is a unique index.
2. **slice:** when there is a monotonic index i.e. repetitive values in index.
3. **mask:** it returns a list of boolean values.

52.1.3.14 14. DataFrame.head(n = 5)

Parameters

n: It accepts an integer parameter which represents the number of rows to select. (Default: 5)

Purpose

This function returns the first n rows for the object based on position. It is useful for quickly testing if your object has the right type of data in it.

For negative values of n, this function returns all rows except the last n rows, equivalent to df[:-n].

Creating frovedis DataFrame from pandas DataFrame:

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

# creating the dataframe
pd_df = pd.DataFrame({'animal': ['alligator', 'bee', 'falcon', 'lion',
                                'monkey', 'parrot', 'shark', 'whale', 'zebra']})

```

```
})
```

```
#creating frovedis dataframe  
fd_df = fdf.DataFrame(pd_df)
```

```
# display frovedis dataframe  
fd_df.show()
```

Output

```
index  animal  
0      alligator  
1       bee  
2      falcon  
3       lion  
4      monkey  
5      parrot  
6       shark  
7       whale  
8       zebra
```

Viewing the first 5 lines (by default n=5):

For example,

```
# head() demo with default n value  
fd_df.head().show()
```

Output

```
index  animal  
0      alligator  
1       bee  
2      falcon  
3       lion  
4      monkey
```

Viewing the first 2 lines (n=2):

For example,

```
# head() demo with n = 2  
fd_df.head(2).show()
```

Output

```
index  animal  
0      alligator  
1       bee
```

For negative values of n:

For example,

```
# head() demo with n = -3  
fd_df.head(-3).show()
```

Output

```
index  animal  
0      alligator  
1       bee
```

```

2      falcon
3      lion
4      monkey
5      parrot

```

Return Value

- If *n* is positive integer, it returns a new DataFrame with the first *n* rows.
- If *n* is negative integer, it returns a new DataFrame with all rows except last *n* rows.

52.1.3.15 15. DataFrame.insert(loc, column, value, allow_duplicates = False)

Parameters

loc: It accepts an integer as parameter which represents the Insertion index. It must be in range (0, *n* - 1) where *n* is number of columns in dataframe.

column: It accepts a string object as parameter. It is the label of the inserted column.

value: It accepts an integer or a pandas Series instance or python ndarray as parameter. These are the values to be inserted in the specified 'column'.

allow_duplicates: It accepts a boolean value as parameter. Currently, frovedis does not support duplicate column names. (Default: False)

Purpose

It is used to insert column into DataFrame at specified location.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

pd_df = pd.DataFrame({'col1': [1, 2], 'col2': [3, 4]})
fd_df = fdf.DataFrame(pd_df)
fd_df.show()

```

Output

```

index  col1  col2
0      1    3
1      2    4

```

Inserting [99, 99] at 1st index in 'newcol' column:

For example,

```

fd_df.insert(1, "newcol", [99, 99])

# display frovedis dataframe after insertion
fd_df.show()

```

Output

```

index  col1  newcol  col2
0      1    99      3
1      2    99      4

```

Inserting a Series object having values [5,6] at 0th index in "col0" column:

For example,

```
fd_df.insert(0, "col0", pd.Series([5, 6])).show()
```

Output

index	col0	col1	col2
0	5	1	3
1	6	2	4

Return Value

It returns a self reference.

52.1.3.16 16. DataFrame.isna()

Purpose

This method is used to detect missing values in the frovedis dataframe.

It returns a boolean same-sized object indicating if the values are NA. NA values, such as None or numpy.NaN, gets mapped to True values. Everything else gets mapped to False values. Characters such as empty strings " " or numpy.inf are not considered NA values.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# show which entries in a DataFrame are NA.
peopleDF = {
    'Ename' : ['Michael', None, 'Tanaka', 'Raul', ''],
    'Age' : [29, 30, 27, 19, 0],
    'Country' : ['USA', np.inf, 'Japan', np.nan, 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe
pd_df = pd.DataFrame(peopleDF)

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# to display frovedis dataframe
fd_df.show()
```

Output

index	Ename	Age	Country	isMale
0	Michael	29	USA	0
1	NULL	30	inf	0
2	Tanaka	27	Japan	0
3	Raul	19	NULL	0
4		0	Japan	1

Show which entries in a frovedis DataFrame are NA:

For example,

```
# isna() demo to display fields which are only NA
fd_df.isna().show()
```

Output

index	Ename	Age	Country	isMale
0	0	0	0	0
1	1	0	0	0
2	0	0	0	0
3	0	0	1	0
4	0	0	0	0

Creating frovedis DataFrame from pandas Series:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# frovedis dataframe from a pandas Series object.
ser = pd.Series([5, 6, np.NaN])

# creating a frovedis dataframe
fd_df = fdf.DataFrame(ser)

# display frovedis dataframe
fd_df.show()
```

Output

index	0
0	5
1	6
2	NULL

Show which entries in corresponding frovedis DataFrame are NA:

For example,

```
# isna() demo to display na values mapped to corresponding dataframe
fd_df.isna().show()
```

Output

index	0
0	0
1	0
2	1

Return Value

It returns a new Frovedis DataFrame having all boolean values (0, 1) corresponding to each of the Frovedis DataFrame values depending on whether it is a valid NaN (True i.e. 1) value or not (False i.e. 0).

52.1.3.17 17. DataFrame.isnull()

Purpose

This method is used to detect missing values in the frovedis dataframe. It is an alias of isna().

It returns a boolean same-sized object indicating if the values are NA. NA values, such as None or numpy.NaN, gets mapped to True values. Everything else gets mapped to False values. Characters such as empty strings " or numpy.inf are not considered NA values.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# show which entries in a DataFrame are NA.
peopleDF = {
    'Ename' : ['Michael', None, 'Tanaka', 'Raul', ''],
    'Age' : [29, 30, 27, 19, 0],
    'Country' : ['USA', 'England', 'Japan', np.nan, 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe
pd_df = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
fd_df.show()
```

Output

index	Ename	Age	Country	isMale
0	Michael	29	USA	0
1	NULL	30	England	0
2	Tanaka	27	Japan	0
3	Raul	19	NULL	0
4		0	Japan	1

Show which entries in a frovedis DataFrame are NULL:

For example,

```
# isnull() demo
fd_df.isnull().show()
```

Output

index	Ename	Age	Country	isMale
0	0	0	0	0
1	1	0	0	0
2	0	0	0	0
3	0	0	1	0
4	0	0	0	0

Creating frovedis DataFrame from pandas Series:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# frovedis dataframe from a Series object.
ser = pd.Series([5, 6, np.NaN])

# creating a frovedis dataframe
fd_df = fdf.DataFrame(ser)
```

```
# display frovedis dataframe
fd_df.show()
```

Output

```
index    0
0         5
1         6
2        NULL
```

Show which entries in corresponding frovedis DataFrame are NULL:

For example,

```
# isnull() demo on a frovedis dataframe converted from Series object
fd_df.isnull().show()
```

Output

```
index    0
0         0
1         0
2         1
```

Return Value

It returns a Frovedis DataFrame having boolean values (0, 1) corresponding to each of the Frovedis DataFrame value depending of whether it is a valid NaN (True i.e. 1) value or not (False i.e. 0).

52.1.3.18 18. DataFrame.join(right, on, how = 'inner', lsuffix = '_left', rsuffix = '_right', sort = False, join_type = 'bcast')

Parameters

right: It accepts a Frovedis DataFrame instance or a pandas DataFrame instance or a list of DataFrame instances as parameter. Index should be similar to one of the columns in this one. If a pandas Series instance is passed, its name attribute must be set, and that will be used as the column name in the resulting joined dataframe.

on: It accepts a string object or a list of strings as parameter. It is the column or index name(s) in the caller to join on the index in other, otherwise joins index-on-index. **This parameter must be provided. It can not be None.**

how: It accepts a string object as parameter that specifies how to handle the operation of the two dataframes. (Default: 'inner')

1. **'left'**: form union of calling dataframe's index (or column if 'on' is specified) with other dataframe's index, and sort it lexicographically.
2. **'inner'**: form intersection of calling dataframe's index (or column if 'on' is specified) with other dataframe's index, preserving the order of the calling's one.

lsuffix: It accepts a string object as parameter. It adds the suffix to left DataFrame's overlapping columns. (Default: '_left')

rsuffix: It accepts a string object as parameter. It adds the suffix to right DataFrame's overlapping columns. (Default: '_right')

sort: It accepts a boolean type value. It orders resultant dataframe lexicographically by the join key. If False, the order of the join key depends on the join type ('how' keyword). (Default: False)

join_type: It accepts a string type object as parameter. It represents the type of join to be used internally. It can be specified as 'bcast' for broadcast join or 'hash' for hash join. (Default: 'bcast')

Purpose

It joins columns of another dataframe.

It joins columns with other dataframe either on index or on a key column. Efficiently join multiple DataFrame instances by index at once by passing a list.

Note:- Parameters 'on', 'lsuffix', and 'rsuffix' are not supported when passing a list of DataFrame instances.

Creating two frovedis dataframes:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame({'key': ['K0', 'K1', 'K2', 'K3', 'K4', 'K5'],
                        'A': ['A0', 'A1', 'A2', 'A3', 'A4', 'A5']})

# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display frovedis dataframe
fd_df1.show()

# creating another pandas dataframe
pd_df2 = pd.DataFrame({'key': ['K0', 'K1', 'K2'],
                        'B': ['B0', 'B1', 'B2']})

# creating another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)

# display other frovedis dataframe
fd_df2.show()
```

Output

index	key	A
0	K0	A0
1	K1	A1
2	K2	A2
3	K3	A3
4	K4	A4
5	K5	A5

index	key	B
0	K0	B0
1	K1	B1
2	K2	B2

Join dataframes using their indexes:

For example,

```
# join() demo with lsuffix, rsuffix and index parameters
fd_df1.join(fd_df2, 'index', lsuffix = '_caller', rsuffix = '_other').show()
```

Output

index	key_caller	A	key_other	B
0	K0	A0	K0	B0

1	K1	A1	K1	B1
2	K2	A2	K2	B2

Using the on parameter:

For example,

```
# join() demo using the 'key' columns,
# 'key' to be available in both fd_df1 and fd_df2
fd_df1.join(fd_df2, on = 'key').show()
```

Output

index	key	A	B
0	K0	A0	B0
1	K1	A1	B1
2	K2	A2	B2

Using the how parameter:

For example,

```
# join() demo with parameter how = 'left'
fd_df1.join(fd_df2, 'key', how = 'left').show()
```

Output,

index	key	A	B
0	K0	A0	B0
1	K1	A1	B1
2	K2	A2	B2
3	K3	A3	NULL
4	K4	A4	NULL
5	K5	A5	NULL

Return Value

It returns a new Frovedis DataFrame containing columns from both the DataFrame instances.

52.1.3.19 19. DataFrame.merge(right, on = None, how = 'inner', left_on = None, right_on = None, left_index = False, right_index = False, sort = False, suffixes = ('_x', '_y'), copy = True, indicator = False, join_type = 'bcast')

Parameters

right: It accepts a Frovedis DataFrame instance or a pandas DataFrame instance or a list of Frovedis DataFrame instances as parameter. Index should be similar to one of the columns in this one. If a pandas Series instance is passed, its name attribute must be set, and that will be used as the column name in the resulting joined dataframe.

on: It accepts a string object or a list of strings as parameter. It is the column or index level names to join on. These must be present in both dataframes. (Default: None)

When it is None and not merging on indexes then this defaults to the intersection of the columns in both dataframes.

how: It accepts a string object as parameter. It informs the type of merge operation on the two objects. (Default: 'inner')

1. **'left':** form union of calling dataframe's index (or column if 'on' is specified) with other's index and sort it lexicographically.

2. **'inner':** form intersection of calling dataframe's index (or column if 'on' is specified) with other's index, preserving the order of the calling's one.

left_on: It accepts a string object or a list of strings as parameter. It represents column names to join on in the left dataframe. It can also be an array or list of arrays of the length of the left dataframe. These arrays are treated as if they are columns. (Default: None)

right_on: It accepts a string object or a list of strings as parameter. It represents column names to join on in the right dataframe. It can also be an array or list of arrays of the length of the right dataframe. These arrays are treated as if they are columns. (Default: None)

left_index: It accepts a boolean value as parameter. It is used to specify whether to use the index from the left dataframe as the join key. **Either parameter 'left_on' or 'left_index' can be used, but not combination of both.** (Default: False)

right_index: It accepts a boolean value as parameter. It is used to specify whether to use the index from the right dataframe as the join key. **Either parameter 'right_on' or 'right_index' can be used, but not combination of both.** (Default: False)

sort: It accepts a boolean value. When this is explicitly set to True, it sorts the join keys lexicographically in the resultant dataframe. When it is False, the order of the join keys depends on the join type ('how' parameter). (Default: False)

suffixes: It accepts a list like (list or tuple) object of strings of length two as parameter. It indicates the suffix to be added to the overlapping column names in left and right respectively. Need to explicitly pass a value of None instead of a string to indicate that the column name from left or right should be left as-it is, with no suffix. At least one of the values must not be None. (Default: ("_x", "_y"))

Note:- During merging two DataFrames, the overlapping column names should be different. For example: `suffixes = (False, False)`, then the overlapping columns would have the same name so merging operation will fail. Also when there is no overlapping column, then this parameter is ignored automatically.

copy: It is an unused parameter. (Default: True)

indicator: It is an unused parameter. (Default: False)

join_type: It accepts a string type object as parameter. It represents the type of join to be used internally. It can be specified as 'bcast' for broadcast join or 'hash' for hash join. (Default: 'bcast')

Purpose

It is a utility to merge dataframe objects with a database-style join.

The join is done on columns or indexes. If joining columns on columns, the DataFrame indexes will be ignored. Otherwise if joining indexes on indexes or indexes on a column or columns, the index will be passed on.

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame({'lkey': ['foo', 'bar', 'baz', 'foo'],
                       'value': [1, 2, 3, 5]})

# creating another pandas dataframe
pd_df2 = pd.DataFrame({'rkey': ['foo', 'bar', 'baz', 'foo'],
                       'value': [5, 6, 7, 8]})

# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display frovedis dataframe
fd_df1.show()

# creating another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)
```

```
# display other frovedis dataframe
fd_df2.show()
```

Output

index	lkey	value
0	foo	1
1	bar	2
2	baz	3
3	foo	5

index	rkey	value
0	foo	5
1	bar	6
2	baz	7
3	foo	8

For example,

```
# merge() demo with parameters left_on and right_on
fd_df1.merge(fd_df2, left_on = 'lkey', right_on = 'rkey').show()
```

Output

index	lkey	value_x	rkey	value_y
0	foo	1	foo	8
1	foo	1	foo	5
2	bar	2	bar	6
3	baz	3	baz	7
4	foo	5	foo	8
5	foo	5	foo	5

For example,

```
# merge() demo with parameters left_on , right_on and suffixes
fd_df1.merge(fd_df2, left_on = 'lkey', right_on = 'rkey', suffixes = ('_left', '_right')).show()
```

Output

index	lkey	value_left	rkey	value_right
0	foo	1	foo	8
1	foo	1	foo	5
2	bar	2	bar	6
3	baz	3	baz	7
4	foo	5	foo	8
5	foo	5	foo	5

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df1 = pd.DataFrame({'lkey': ['foo', 'bar', 'baz', 'foo'],
                       'left_value': [1, 2, 3, 5]})

# creating another pandas dataframe
pd_df2 = pd.DataFrame({'rkey': ['foo', 'bar', 'baz', 'foo'],
                       'right_value': [5, 6, 7, 8]})
```

```
# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)

# display a frovedis dataframe
fd_df1.show()

# creating another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)

# display other frovedis dataframe
fd_df2.show()
```

Output

index	lkey	left_value
0	foo	1
1	bar	2
2	baz	3
3	foo	5

index	rkey	right_value
0	foo	5
1	bar	6
2	baz	7
3	foo	8

For example,

```
# merge() demo with same suffixes on 'lkey' and 'rkey'
fd_df1.merge(fd_df2, left_on = 'lkey', right_on = 'rkey', suffixes = ('_test', '_test')).show()
```

Output

index	lkey	left_value	rkey	right_value
0	foo	1	foo	8
1	foo	1	foo	5
2	bar	2	bar	6
3	baz	3	baz	7
4	foo	5	foo	8
5	foo	5	foo	5

Note:- In above example, suffix is ignored as merging column labels are different. To have suffix, the column name must be same in both dataframes but then the suffixes must be different.

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating two pandas datframes
pd_df1 = pd.DataFrame({'lkey': ['foo', 'bar', 'baz', 'foo'],
                        'value': [1, 2, 3, 5]})
pd_df2 = pd.DataFrame({'rkey': ['foo', 'bar', 'baz', 'foo'],
                        'value': [5, 6, 7, 8]})

# creating a frovedis dataframe
fd_df1 = fdf.DataFrame(pd_df1)
```

```

# update index values of fd_df1
fd_df1.update_index(['a','b', 'c', 'd'], key = 'index', inplace = True)

# display frovedis dataframe
fd_df1.show()

# creating another frovedis dataframe
fd_df2 = fdf.DataFrame(pd_df2)

# update index values of fd_df2
fd_df2.update_index(['a','b', 'c', 'd'], key = 'index', inplace = True)

# display frovedis dataframe
fd_df2.show()

```

Output

index	lkey	value
a	foo	1
b	bar	2
c	baz	3
d	foo	5

index	rkey	value
a	foo	5
b	bar	6
c	baz	7
d	foo	8

For example,

```

# merge() demo with left_index = True and right_index = True
fd_df1.merge(fd_df2, left_index = True, right_index = True).show()

```

Output

index	lkey	value_x	rkey	value_y
a	foo	1	foo	5
b	bar	2	bar	6
c	baz	3	baz	7
d	foo	5	foo	8

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

# creating two pandas datframes
pd_df1 = pd.DataFrame({'a': ['foo', 'bar'], 'b': [1, 2]})
pd_df2 = pd.DataFrame({'a': ['foo', 'baz'], 'c': [3, 4]})

# creating two frovedis datframes
fd_df1 = fdf.DataFrame(pd_df1)
fd_df2 = fdf.DataFrame(pd_df2)

# display frovedis dataframes
fd_df1.show()

```

```
fd_df2.show()
```

Output

```
index  a      b
0      foo    1
1      bar    2
```

```
index  a      c
0      foo    3
1      baz    4
```

For example,

```
# merge() demo with how = 'inner' and 'on' parameter
fd_df1.merge(fd_df2, how = 'inner', on = 'a').show()
```

Output

```
index  a      b      c
0      foo    1      3
```

For example,

```
# merge using 'how' = left and 'on' parameter
fd_df1.merge(fd_df2, how = 'left', on = 'a').show()
```

Output,

```
index  a      b      c
0      foo    1      3
1      bar    2     NULL
```

Return Value

It returns a new Frovedis DataFrame instance with the merged entries of the two DataFrame instances.

52.1.3.20 20. DataFrame.rename(columns, inplace = False)

Parameters

columns: It accepts a dictionary object as parameter. It contains the key as the name of the labels to be renamed and values as the final names.

inplace.: It accepts a boolean object as parameter which specify whether to modify the original DataFrame instance or to return a copy. When it is set to True then the original DataFrame instance is modified. (Default: False)

Purpose

It is used to set the name of the columns.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame({"A": [1, 2, 3], "B": [4, 5, 6]})

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)
```

```
# display frovedis dataframe
fd_df.show()
```

Output

index	A	B
0	1	4
1	2	5
2	3	6

Rename columns using a mapping:

For example,

```
# rename columns 'A' and 'B' to 'a' and 'c' respectively
fd_df.rename(columns={"A": "a", "B": "c"}).show()
```

Output

index	a	c
0	1	4
1	2	5
2	3	6

Rename columns using a mapping and keep the dataframe in the same variable:

For example,

```
# rename() demo with inplace = True
fd_df.rename(columns={"A": "X", "B": "Y"}, inplace = True)
```

Output,

index	X	Y
0	1	4
1	2	5
2	3	6

Return Value

- It returns a new Frovedis DataFrame with the updated label name for the specified columns.
- It returns None when 'inplace' parameter is set to True.

52.1.3.21 21. DataFrame.reset_index(drop = False, inplace = False)

Parameters

drop: It accepts a boolean value as parameter. Do not try to insert index into dataframe columns. This resets the index to the default integer index. (Default: False)

inplace: It accepts a boolean values as parameter. When it is explicitly set to True, it modifies the original object directly instead of creating a copy of DataFrame object. (Default: False)

Purpose

It is used to reset the Index label of the DataFrame. A new Index label is inserted with default integer values from 0 to n-1 where n is the number of rows.

Note:- MultiIndex is not supported by Frovedis DataFrame.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe
pd_df = pd.DataFrame(peopleDF)

# updating the index values of pandas dataframe
pd_df.index = ['a', 'b', 'c', 'd', 'a']

# creating frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display the frovedis dataframe
fd_df.show()

```

Output

index	Ename	Age	Country	isMale
a	Michael	29	USA	0
b	Andy	30	England	0
c	Tanaka	27	Japan	0
d	Raul	19	France	0
a	Yuta	31	Japan	1

Resetting the index and keeping the old index is as a column:

For example,

```

# reset_index() demo
fd_df.reset_index().show

```

Output

label_0	index_col	Ename	Age	Country	isMale
0	a	Michael	29	USA	0
1	b	Andy	30	England	0
2	c	Tanaka	27	Japan	0
3	d	Raul	19	France	0
4	a	Yuta	31	Japan	1

When we reset the index, the old index is added as a column, and a new sequential index is used.

Using drop parameter:

For example,

```

# reset_index() demo with drop parameter
fd_df.reset_index(drop = True).show()

```

Output

index	Ename	Age	Country	isMale
0	Michael	29	USA	0

1	Andy	30	England	0
2	Tanaka	27	Japan	0
3	Raul	19	France	0
4	Yuta	31	Japan	1

We can use the drop parameter to avoid the old index being added as a column.

Return Value

- It returns a new Frovedis DataFrame with the default sequence in index label.
- It returns None if 'inplace' parameter is set to True.

52.1.3.22 22. DataFrame.set_index(keys, drop = True, append = False, inplace = False, verify_integrity = False)

Parameters

keys: It accepts a string object as parameter. This parameter can be a single column key.

drop: It accepts a boolean value as parameter. When it is set to True, it will remove the column which is selected as new index. **Currently, Frovedis doesn't support drop = False.** (Default: True)

append: It accepts a boolean value as parameter. It will decide whether to append columns to existing index. **Currently, Frovedis doesn't support append = True.** (Default: False)

inplace: It accepts a boolean values as parameter which is when explicitly set to True, it modifies the original object directly instead of creating a copy of DataFrame instance. (Default: False)

verify_integrity: It accepts a boolean value as parameter. When it is set to True, it checks the new index for duplicates. Performance of this method will be better when it is set to False. (Default: False)

Purpose

It is used to set the frovedis dataframe index using existing columns. The index will replace the existing index.

Note:- Frovedis DataFrame does not support Multi Index.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame({'month': [1, 4, 1, 10],
                      'year': [2012, 2014, 2013, 2014],
                      'sale': [55, 40, 84, 31]})

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
fd_df.show()
```

Output

index	month	year	sale
0	1	2012	55
1	4	2014	40
2	1	2013	84
3	10	2014	31

Set the ‘month’ column to be the new index:

For example,

```
fd_df.set_index('month').show()
```

Output

month	year	sale
1	2012	55
4	2014	40
1	2013	84
10	2014	31

Set the ‘month’ column to be the new index. Also, it checks for duplicates before setting:

For example,

```
# set_index() demo with verify_integrity = True
fd_df.set_index('month', verify_integrity = True).show()
```

Output

month	year	sale
1	2012	55
4	2014	40
7	2013	84
10	2014	31

Note:- In above example, the column which is being selected as index must have unique values.

Return Value

- It returns a new Frovedis DataFrame where the Index column label is replace with specified column label.
- It returns None when ‘inplace’ parameter is set to True.

52.1.3.23 23. DataFrame.show()

Purpose

This method is used to display the Frovedis DataFrame on the console. It can display full dataframe or some selected columns of the DataFrame (single or multi-column).

It can be used either with any method which returns a Frovedis DataFrame instance type compatible with string type.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating the dataframe
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}
```

```
pdf = pd.DataFrame(peopleDF)

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pdf)

# display frovedis dataframe
print("Displaying complete frovedis dataframe")
fd_df.show()
```

Output

```
Displaying complete frovedis dataframe
index  Ename  Age   Country isMale
0      Michael 29    USA      0
1      Andy   30   England  0
2      Tanaka 27    Japan   0
3      Raul   19   France   0
4      Yuta   31    Japan   1
```

Display dataframe by selecting single column:

For example,

```
print("Displaying frovedis dataframe with just Ename column")
fd_df["Ename"].show()      # single column
```

Output

```
Displaying frovedis dataframe with just Ename column
index  Ename
0      Michael
1      Andy
2      Tanaka
3      Raul
4      Yuta
```

Displaying multiple columns:

For example,

```
print("Displaying frovedis dataframe with Ename and Age columns")
fd_df[["Ename","Age"]].show() # multiple column
```

Output

```
Displaying frovedis dataframe with Ename and Age columns
index  Ename  Age
0      Michael 29
1      Andy   30
2      Tanaka 27
3      Raul   19
4      Yuta   31
```

Displaying frovedis dataframe using condition based indexing operator:

For example,

```
print("Displaying frovedis dataframe using condition based slicing operator")
fd_df[fd_df.Age > 19].show()
```

Output

Displaying frovedis dataframe using condition based slicing operator

```
index  Ename  Age   Country isMale
0      Michael 29    USA      0
1      Andy   30   England 0
2      Tanaka 27    Japan   0
4      Yuta   31    Japan   1
```

Displaying frovedis dataframe using chaining of methods:

For example,

```
print("Displaying frovedis dataframe using chaining of methods")
fd_df[fd_df.Country.str.contains("a")].show()
```

Output

Displaying frovedis dataframe using chaining of methods

```
index  Ename  Age   Country isMale
1      Andy   30   England 0
2      Tanaka 27    Japan   0
3      Raul   19   France   0
4      Yuta   31    Japan   1
```

Return Value

It return nothing.

52.1.3.24 24. DataFrame.tail(n = 5)

Parameters

n: It accepts an integer as parameter. It represents the number of rows which is to be selected. (Default: 5)

Purpose

This utility is used to return the last **n** rows from the DataFrame.

It is useful for quickly verifying data, for example, after sorting or appending rows.

For negative values of **n**, this function returns all rows except the first **n** rows, equivalent to **fdf[n:]**.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame({'animal': ['alligator', 'bee', 'falcon', 'lion',
                                'monkey', 'parrot', 'shark', 'whale', 'zebra']})

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

print("Viewing all entries of dataframe")
fd_df.show()
```

Output

Viewing all entries of dataframe

```
index  animal
0      alligator
```

```
1      bee
2      falcon
3      lion
4      monkey
5      parrot
6      shark
7      whale
8      zebra
```

Viewing the last 5 lines (by default n=5):

For example,

```
# tail() demo
print("Viewing the last 5 lines")
fd_df.tail().show()
```

Output

```
Viewing the last 5 lines
index  animal
4      monkey
5      parrot
6      shark
7      whale
8      zebra
```

Viewing the last 3 lines:

For example,

```
# tail() demo with n = 3
print("Viewing the last 3 lines")
fd_df.tail(3).show()
```

Output

```
Viewing the last 3 lines
index  animal
6      shark
7      whale
8      zebra
```

For negative values of n:

For example,

```
# tail() demo with n = -2
print("For negative values of n")
fd_df.tail(-2).show()
```

Output

```
For negative values of n
index  animal
2      falcon
3      lion
4      monkey
5      parrot
6      shark
7      whale
```

8 zebra

Return Value

It returns a new Frowedis DataFrame instance with last **n** rows.

52.2 SEE ALSO

- **DataFrame - Introduction**
- **DataFrame - Conversion Functions**
- **DataFrame - Sorting Functions**
- **DataFrame - Math Functions**
- **DataFrame - Aggregate Functions**

Chapter 53

DataFrame Conversion Functions

53.1 NAME

DataFrame Conversion Functions - this manual contains all functions related to conversion of datatypes with respect to frovedis dataframe.

53.1.1 DESCRIPTION

In frovedis during data pre-processing and manipulation, there might be the need to change the data type of the variable to a particular type for better cleaning and understanding of the data.

For this inter-conversion within the variables, frovedis dataframe offer various conversion functions like `asDF()`, `to_dict()`, `to_numpy()`, etc.

53.1.2 Public Member Functions

1. `asDF(df)`
2. `to_dict(orient = "dict", into = dict)`
3. `to_numpy(dtype = None, copy = False, na_value = None)`
4. `to_pandas()`
5. `to_frovedis_rowmajor_matrix(t_cols, dtype = np.float32)`
6. `to_frovedis_colmajor_matrix(t_cols, dtype = np.float32)`
7. `to_frovedis_crs_matrix(t_cols, cat_cols, dtype = np.float32, need_info = False)`
8. `to_frovedis_crs_matrix_using_info(info, dtype = np.float32)`

53.1.3 Detailed Description

53.1.3.1 1. DataFrame.asDF(df)

Parameters

df: It takes either a Frovedis DataFrame or a Pandas DataFrame or a Series instance.

Purpose

It creates a new Frovedis DataFrame after suitable conversion.

Using `asDF()` to create frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# convert to pandas dataframe
pd_df = pd.DataFrame(peopleDF)

print("display the pandas dataframe type")
print(type(pd_df))

# convert pandas DataFrame to frovedis DataFrame object
print("display type after conversion to frovedis dataframe")
print(type(fdf.DataFrame.asDF(pd_df)))
```

Output

```
display the pandas dataframe type
<class 'pandas.core.frame.DataFrame'>

display type after conversion to frovedis dataframe
<class 'frovedis.dataframe.df.DataFrame'>
```

Using asDF() to create frovedis DataFrame from pandas Series:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# create a Series object
sdf1 = pd.Series([1, 2], dtype='int32')
print("display series type")
print(type(sdf1))

# convert a Series object to Frovedis DataFrame object
print("display type after conversion to frovedis dataframe")
print(type(fdf.DataFrame.asDF(sdf1)))
```

Output

```
display series type
<class 'pandas.core.series.Series'>

display type after conversion to frovedis dataframe
<class 'frovedis.dataframe.df.DataFrame'>
```

Return Value

It returns a Frovedis DataFrame instance after suitable conversion

53.1.3.2 2. DataFrame.to_dict(orient = “dict”, into = dict)**Parameters**

orient: It accepts a string object as parameter. It is used to determine the type of the values of the dictionary. (Default: ‘dict’)

1. ‘dict’: dict like {column -> {index -> value}}
2. ‘list’: dict like {column -> [values]}

into: This parameter is used for mapping in the return value. Currently it only supports OrderedDict as return type. (Default: dict)

Purpose

It is used to convert the Frovedis DataFrame to a dictionary.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a pandas dataframe
pd_df = pd.DataFrame({'col1': [1, 2],
                      'col2': [0.5, 0.75]},
                     index=['row1', 'row2'])

# create a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# to display frovedis dataframe
fd_df.show()
```

Output

```
index  col1  col2
row1    1    0.5
row2    2    0.75
```

Converting frovedis DataFrame to dictionary:

For example,

```
print(fd_df.to_dict())
```

Output

```
OrderedDict([('col1', {'row1': 1, 'row2': 2}), ('col2', {'row1': 0.5, 'row2': 0.75})])
```

Converting frovedis DataFrame to dictionary and using orient parameter:

For example,

```
# to_dict() demo with orient = 'list'
print(fd_df.to_dict(orient = 'list'))
```

Output

```
OrderedDict([('index', array(['row1', 'row2'], dtype='<U4')),
             ('col1', array([1, 2])), ('col2', array([0.5 , 0.75]))])
```

Return Value

It returns a dictionary representing the Frovedis DataFrame instance. The resulting transformation depends on the ‘orient’ parameter.

53.1.3.3 3. DataFrame.to_numpy(dtype = None, copy = False, na_value = None)**Parameters**

dtype : It accepts the dtype parameter which decides the datatype of numpy ndarray. (Default: None)
When it is None (not specified explicitly), it will be set as double (float64).

copy : This is an unused parameter. (Default: False)

na_value : This is an unused parameter. (Default: None)

Purpose

This method is used to convert a frovedis dataframe into a numpy array.

The parameters: “copy” and “na_value” are simply kept in to make the interface uniform to the pandas DataFrame.to_numpy().

This is not used in the frovedis implementation.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a python dict
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# a pandas dataframe object from dict
pd_df = pd.DataFrame(peopleDF)

# to create a frovedis dataframe object
fd_df = fdf.DataFrame(pd_df)

# display frovedis dataframe
fd_df.show()

# below will display a Frovedis dataframe type
print(type(fd_df))
```

Output

index	Ename	Age	Country	isMale
0	Michael	29	USA	0
1	Andy	30	England	0
2	Tanaka	27	Japan	0
3	Raul	19	France	0
4	Yuta	31	Japan	1

```
<class 'frovedis.dataframe.df.DataFrame'>
```

Convert frovedis DataFrame into numpy array:

For example,

```
# to_numpy() demo to convert a frovedis dataframe into numpy array
# below will display a numpy array type
```

```
print(fd_df.to_numpy())

# below will display a numpy array type
print(type(fd_df.to_numpy()))
```

Output

```
[['Michael' '29' 'USA' '0']
 ['Andy' '30' 'England' '0']
 ['Tanaka' '27' 'Japan' '0']
 ['Raul' '19' 'France' '0']
 ['Yuta' '31' 'Japan' '1']]
```

```
<<class 'numpy.ndarray'>
```

Return Value

It returns a numpy ndarray that represents a Frovedis DataFrame instance. It has shape (**nRows** , **nCols**).

53.1.3.4 4. DataFrame.to_pandas()

Purpose

It converts a Frovedis DataFrame instance into a pandas DataFrame instance.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a python dictionary
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}

# creating a pandas dataframe from python dict
pd_df = pd.DataFrame(peopleDF)

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# below will display a Frovedis dataframe
fd_df.show()

# below will display a Frovedis dataframe type
print(type(fd_df))
```

Output

index	Ename	Age	Country	isMale
0	Michael	29	USA	0
1	Andy	30	England	0
2	Tanaka	27	Japan	0
3	Raul	19	France	0

```
4      Yuta      31      Japan      1
```

```
<class 'frovedis.dataframe.df.DataFrame'>
```

Convert frovedis DataFrame to pandas DataFrame:

For example,

```
# to_pandas() demo to convert frovedis dataframe to pandas dataframe
print(fd_df.to_pandas())
```

```
# below will display a pandas dataframe type after conversion
print(type(fd_df.to_pandas()))
```

Output

```
index  Ename  Age  Country  isMale
0      Michael  29      USA    False
1         Andy  30  England    False
2       Tanaka  27   Japan    False
3        Raul  19   France    False
4         Yuta  31   Japan     True
```

```
<class 'pandas.core.frame.DataFrame'>
```

Return Value

It returns a pandas DataFrame instance after suitable conversion.

53.1.3.5 5. DataFrame.to_frovedis_rowmajor_matrix(t_cols, dtype = np.float32)

Parameters

t_cols: It accepts a list of string type argument where each of the member of the list is the name of the column labels.

dtype: It accepts a dtype type argument which is the type of the resultant values. Currently, only float (float32) and double (float64) types are supported. (Default: np.float32)

Purpose

This method is used to convert a Frovedis DataFrame instance into FrovedisRowmajorMatrix instance.

Loading input file data into frovedis DataFrame:

For example,

```
import frovedis.dataframe as fdf
```

```
# read_csv demo to get values
```

```
df = fdf.read_csv("./input/numbers.csv", names = ['one', 'two', 'three', 'four'])
```

```
# display frovedis dataframe
```

```
df.show()
```

Output

```
index  one    two    three  four
0      10    10.23    F      0
1      12    12.2     F      0
2      13    34.8999  D      1
3      15    100.12   A      2
```

Convert frovedis DataFrame into FrovedisRowmajorMatrix:

For example,

```
# to convert frovedis dataframe into frovedis rowmajor matrix
row_mat = df.to_frovedis_rowmajor_matrix(['one', 'two'], dtype = np.float64)

# display row_mat (row major matrix)
row_mat.debug_print()
```

Output

```
matrix:
num_row = 4, num_col = 2
node 0
node = 0, local_num_row = 4, local_num_col = 2, val = 10 10.23 12 12.2 13 34.9 15 100.12
```

Return Value

It returns a FrovedisRowmajorMatrix instance after suitable conversion.

53.1.3.6 6. DataFrame.to_frovedis_colmajor_matrix(t_cols, dtype = np.float32)**Parameters**

t_cols: It accepts a list of string type argument where each of the member of the list is the name of the column labels.

dtype: It accepts a dtype type argument which is the type of the resultant values. Currently only float (float32) and double (float64) types are supported. (Default: np.float32)

Purpose

This method converts a Frovedis DataFrame instance to FrovedisColmajorMatrix instance.

Loading input file data into frovedis DataFrame:

For example,

```
import frovedis.dataframe as fdf

# read_csv demo
df = fdf.read_csv("./input/numbers.csv", names=['one', 'two', 'three', 'four'])

# display frovedis dataframe
df.show()
```

Output

index	one	two	three	four
0	10	10.23	F	0
1	12	12.2	F	0
2	13	34.8999	D	1
3	15	100.12	A	2

Convert frovedis DataFrame into FrovedisColmajorMatrix:

For example,

```
# to convert frovedis dataframe into frovedis colmajor matrix
col_mat = df.to_frovedis_colmajor_matrix(['one', 'two']) # default dtype = np.float32

# display col_mat
col_mat.debug_print()
```

Output

```
matrix:
num_row = 4, num_col = 2
node 0
node = 0, local_num_row = 4, local_num_col = 2, val = 10 12 13 15 10.23 12.2 34.9 100.12
```

Return Value

It returns a FrovedisColmajorMatrix instance after converting the original Frovedis DataFrame instance.

53.1.3.7 7. DataFrame.to_frovedis_crs_matrix(t_cols, cat_cols, dtype = np.float32, need_info = False)

Parameters

t_cols: It accepts a list of string type argument where each of the member of the list is the name of the column labels.

cat_cols: It accepts a list of strings as parameter where strings are the column names. It stands for categorical columns. It represents how the repetitive values are distributed in the specified column. It adds number of columns corresponding to the number of distinct values in the specified column to the matrix. And values corresponding to the value of the specified column is set to 1 and remaining values are set to 0.

dtype: It accepts a datatype as parameter which is the type of the resultant values. Currently only float (float32) and double (float64) types are supported. (Default: np.float32)

need_info: It accepts a boolean value as parameter. When this is explicitly set to True, it returns an additional value of type 'df_to_sparse_info'. (Default: False)

Purpose

It converts a Frovedis DataFrame instance to FrovedisCRSMatrix instance.

Loading input file data into frovedis DataFrame:

For example,

```
import frovedis.dataframe as fdf
import numpy as np

# read_csv demo
df = fdf.read_csv("./input/numbers.csv", names = ['one', 'two', 'three', 'four'])

# display frovedis dataframe
df.show()
```

Output

index	one	two	three	four
0	10	10.23	F	0
1	12	12.2	F	0
2	13	34.8999	D	1
3	15	100.12	A	2

Convert frovedis DataFrame to FrovedisCRSMatrix:

For example,

```
# to convert frovedis dataframe to frovedis crs matrix
crs_mat,info = df.to_frovedis_crs_matrix(['one', 'two', 'four'],
                                         ['three'],
                                         need_info = True) # default dtype = np.float32

# display crs_matrix
```

```
crs_mat.debug_print()

# converting crs_matrix into display it in better form
mat_t = crs_mat.to_scipy_matrix().todense()
print(mat_t)
```

Output

```
Active Elements: 12
matrix:
num_row = 4, num_col = 5
node 0
local_num_row = 4, local_num_col = 5
val : 10 10.23 1 12 12.2 1 13 34.9 1 15 100.12 1
idx : 0 1 2 0 1 2 0 1 3 0 1 4
off : 0 3 6 9 12
```

```
[[ 10.    10.23   0.     0.     1.  ]
 [ 12.    12.2    0.     0.     1.  ]
 [ 13.    34.9    0.     1.     0.  ]
 [ 15.   100.12   1.     0.     0.  ]]
```

Return Value

It returns a FrovedisCRSMatrix instance after converting the original Frovedis DataFrame instance.

53.1.3.8 8. DataFrame.to_frovedis_crs_matrix_using_info(info, dtype = np.float32)

Parameters

info: It accepts an instance of 'df_to_sparse_info' type.

dtype: It accepts a dtype type argument which is the type of the resultant values. Currently only float (float32) and double (float64) types are supported. (Default: np.float32)

Purpose

It converts a Frovedis DataFrame instance to FrovedisCRSMatrix instance and provided an info object of 'df_to_sparse_info' class.

Loading input file data into frovedis DataFrame:

For example,

```
import frovedis.dataframe as fdf

# read_csv demo
df = fdf.read_csv("./input/numbers.csv", names=['one', 'two', 'three', 'four'])

# display frovedis dataframe
df.show()
```

Output

index	one	two	three	four
0	10	10.23	F	0
1	12	12.2	F	0
2	13	34.8999	D	1
3	15	100.12	A	2

Convert frovedis DataFrame to FrovedisCRSMatrix:

For example,

```
# to use to_frovedis_crs_matrix_using_info
crs_mat,info = df.to_frovedis_crs_matrix(['one', 'two', 'four'],
                                         ['four'],
                                         need_info=True) # default dtype = np.float32

print(type(info))

crs_mat2 = df.to_frovedis_crs_matrix_using_info(info)
print(type(crs_mat2))

# display crs_matrix
crs_mat2.debug_print()
```

Output

```
<class 'frovedis.dataframe.info.df_to_sparse_info'>

<class 'frovedis.matrix.crs.FrovedisCRSMatrix'>

Active Elements: 12
matrix:
num_row = 4, num_col = 5
node 0
local_num_row = 4, local_num_col = 5
val : 10 10.23 1 12 12.2 1 13 34.9 1 15 100.12 1
idx : 0 1 2 0 1 2 0 1 3 0 1 4
off : 0 3 6 9 12
```

Return Value

It returns a FrovedisCRSMatrix instance after converting Frovedis DataFrame instance.

53.2 SEE ALSO

- **DataFrame - Introduction**
- **DataFrame - Generic Functions**
- **DataFrame - Sorting Functions**
- **DataFrame - Math Functions**
- **DataFrame - Aggregate Functions**

Chapter 54

DataFrame Sorting Functions

54.1 NAME

DataFrame Sorting Functions - this manual contains list of all functions related to sorting operations performed on frovedis dataframe.

54.1.1 DESCRIPTION

Frovedis dataframe provides various sorting functions such as `sort_index()`, `sort_values()`, etc.

These can be used to sort data along rows or columns of frovedis dataframe, either in ascending or descending order.

Currently, sorting operations will be performed on a copy of frovedis dataframe. Inplace sorting is not supported yet.

54.1.2 Public Member Functions

1. `nlargest(n, columns, keep = 'first')`
2. `nsmallest(n, columns, keep = 'first')`
3. `sort(columns = None, axis = 0, ascending = True, inplace = False, kind = 'radixsort', na_position = 'last', **kwargs)`
4. `sort_index(axis = 0, ascending = True, inplace = False, kind = 'quicksort', na_position = 'last')`
5. `sort_values(by, axis = 0, ascending = True, inplace = False, kind = 'radixsort', na_position = 'last')`

54.1.3 Detailed Description

54.1.3.1 1. DataFrame.nlargest(n, columns, keep = 'first')

Parameters

n: It accepts an integer parameter which represents the number of rows to return.

columns: It accepts a string type or list of strings type parameter which represents the column label(s) to order by.

keep: It accepts a string object parameter. This parameter is used when there are duplicate values. (Default: 'first')

- 'first' : it prioritizes the first occurrence(s).

- **'last'** : it prioritizes the last occurrence(s).
- **'all'** : it is used to not drop any duplicates, even when it means selecting more than **n** items.

Purpose

It is used to return the **top n rows** ordered by the specified columns in descending order. The columns that are not specified are returned as well, but not used for ordering.

This method is equivalent to `DataFrame.sort_values(columns, ascending = False).head(n)`, but it is more efficient.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

pd_df = pd.DataFrame({'population': [59000000, 65000000, 434000, 434000,
                                     434000, 337000, 11300, 11300, 11300],
                      'GDP': [1937894, 2583560, 12011, 4520, 12128,
                              17036, 182, 38, 311],
                      'alpha-2': ["IT", "FR", "MT", "MV", "BN",
                                  "IS", "NR", "TV", "AI"]},
                      index=["Italy", "France", "Malta",
                              "Maldives", "Brunei", "Iceland",
                              "Nauru", "Tuvalu", "Anguilla"])

# create a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# to display frovedis dataframe
fd_df.show()
```

Output

index	population	GDP	alpha-2
Italy	59000000	1937894	IT
France	65000000	2583560	FR
Malta	434000	12011	MT
Maldives	434000	4520	MV
Brunei	434000	12128	BN
Iceland	337000	17036	IS
Nauru	11300	182	NR
Tuvalu	11300	38	TV
Anguilla	11300	311	AI

In order to use `nlargest()` to select the 3 rows having the largest values in 'population' column:

For example,

```
fd_df.nlargest(3, 'population').show()
```

Output

index	population	GDP	alpha-2
France	65000000	2583560	FR
Italy	59000000	1937894	IT
Malta	434000	12011	MT

When using `keep='last'`, last occurrences are prioritized. Ties are resolved in reverse order:

For example,

```
# nlargest() demo with keep = 'last'
fd_df.nlargest(3, 'population', keep='last').show()
```

Output

index	population	GDP	alpha-2
France	65000000	2583560	FR
Italy	59000000	1937894	IT
Brunei	434000	12128	BN

When using keep='all', all duplicate items are maintained (not dropped):

For example,

```
# nlargest() demo with keep = 'all'
fdf.nlargest(3, 'population', keep='all').show()
```

Output

index	population	GDP	alpha-2
France	65000000	2583560	FR
Italy	59000000	1937894	IT
Malta	434000	12011	MT
Maldives	434000	4520	MV
Brunei	434000	12128	BN

To order by the largest values in 'population' and then 'GDP' column, multiple columns may be specified:

For example,

```
# nlargest() demo with use of list of columns
fd_df.nlargest(3, ['population', 'GDP']).show()
```

Output

index	population	GDP	alpha-2
France	65000000	2583560	FR
Italy	59000000	1937894	IT
Brunei	434000	12128	BN

Return Value

It returns a Frovedis DataFrame instance with **n** rows ordered by the specified columns in descending order.

54.1.3.2 2. DataFrame.nsmallest(n, columns, keep = 'first')

Parameters

n: It accepts an integer type argument that represents the number of rows to return.

columns: It accepts a string type or list of strings type parameter which represents the column label(s) to order by.

keep: It accepts a string object parameter. This parameter is used when there are duplicate values. (Default: 'first')

1. **'first'**: it prioritizes the first occurrence(s).
2. **'last'**: it prioritizes the last last occurrence(s).
3. **'all'**: it is used to not drop any duplicates, even it means selecting more than **n** items.

Purpose

It is used to return the **top n rows** ordered by the specified columns in ascending order.

It returns the first **n** rows with the smallest values in columns, in ascending order. The columns that are not specified are returned as well, but not used for ordering.

This method is equivalent to `DataFrame.sort_values(columns, ascending=True).head(n)`, but it is more efficient.

Creating frovedis DataFrame from pandas DataFrame:

For example

```
import pandas as pd
import frovedis.dataframe as fdf

pd_df = pd.DataFrame({'population': [59000000, 65000000, 434000,
                                     434000, 434000, 337000, 11300,
                                     11300, 11300],
                      'GDP': [1937894, 2583560, 12011, 4520, 12128,
                              17036, 182, 38, 311],
                      'alpha-2': ["IT", "FR", "MT", "MV", "BN",
                                  "IS", "NR", "TV", "AI"]},
                      index=["Italy", "France", "Malta",
                              "Maldives", "Brunei", "Iceland",
                              "Nauru", "Tuvalu", "Anguilla"])

# create a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# to display frovedis dataframe
fd_df.show()
```

Output

index	population	GDP	alpha-2
Italy	59000000	1937894	IT
France	65000000	2583560	FR
Malta	434000	12011	MT
Maldives	434000	4520	MV
Brunei	434000	12128	BN
Iceland	337000	17036	IS
Nauru	11300	182	NR
Tuvalu	11300	38	TV
Anguilla	11300	311	AI

In order to use `nlargest()` to select the 3 rows having the largest values in ‘population’ column:

For example,

```
fd_df.nsmallest(3, 'population').show()
```

Output

index	population	GDP	alpha-2
Nauru	11300	182	NR
Tuvalu	11300	38	TV
Anguilla	11300	311	AI

When using `keep='last'`, last occurrences are prioritized. Ties are resolved in reverse order:

For example,

```
# nsmallest() demo with keep = 'last' parameter
fd_df.nsmallest(3, 'population', keep='last').show()
```

Output

index	population	GDP	alpha-2
Anguilla	11300	311	AI
Tuvalu	11300	38	TV
Nauru	11300	182	NR

When using `keep='all'`, all duplicate items are maintained (not dropped):

For example

```
# example to use nsmallest with keep = 'all' parameter
fd_df.nsmallest(3, 'population', keep='all').show()
```

Output

index	population	GDP	alpha-2
Nauru	11300	182	NR
Tuvalu	11300	38	TV
Anguilla	11300	311	AI

To order by the largest values in ‘population’ and then ‘GDP’ column, multiple columns may be specified:

For example,

```
# nsmallest() demo with using a list of columns names
fd_df.nsmallest(3, ['population', 'GDP'])
```

Output

index	population	GDP	alpha-2
Tuvalu	11300	38	TV
Nauru	11300	182	NR
Anguilla	11300	311	AI

Return Value

It returns a Frovedis DataFrame object with **n** rows ordered by the specified columns in ascending order.

54.1.3.3 3. DataFrame.sort(columns = None, axis = 0, ascending = True, inplace = False, kind = ‘radixsort’, na_position = ‘last’, **kwargs)

Parameters

columns: It accepts the name or list of names on which sorting will be applied. (Default: None)

If axis is 0 or ‘index’ then this parameter should contain index levels and/or column labels. Currently axis = 0 or ‘index’ is only supported in Frovedis DataFrame.

When it is None (not specified explicitly), it will not perform sorting and it will raise an exception.

axis: It accepts an integer or a string object as parameter. To perform sorting along rows or columns, it is selected by this parameter. (Default: 0)

If axis is 0 or ‘index’, sorting is performed along the row. Currently axis = 0 or ‘index’ is only supported in Frovedis DataFrame.

ascending: It accepts a boolean value or a list of booleans as parameter. The order of sorting is decided by this parameter. Need to specify a list of booleans for multiple sort orders. If this is a list of booleans, then it must match the length of the ‘columns’ parameter. By default, the order of sorting will be ascending and to change the order to descending, explicitly pass it as False. (Default: True)

inplace: It accepts a boolean value as parameter. When it is explicitly set to True, it modifies the original object directly instead of creating a copy of DataFrame object. **Currently, ‘inplace’ = True is not supported by this Frovedis DataFrame method.** (Default: False)

kind: It accepts a string object as parameter to select the type of sorting algorithm. **Currently, Frovedis supports only ‘radixsort’ and other values for ‘kind’ parameter will be ignored internally with a warning.** (Default: ‘radixsort’)

na_position: It accepts a string object as parameter. It decides the position of NaNs after sorting. When it is set to 'last', it puts NaNs at the end. **Currently, Frovedis only supports 'na_position' = 'last'.** (Default: 'last')

****kwargs:** It accepts a dictionary object as parameter. It is used to pass all the other parameters at once in the form of a dictionary object. Currently this is not supported.

Purpose

It is used to sort the values in the specified column(s) along axis = 0 or axis = 'index' in the Frovedis DataFrame.

This method is present only in frovedis. It internally uses sort_values().

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

pd_df = pd.DataFrame({
    'col1': ['A', 'A', 'B', np.nan, 'D', 'C'],
    'col2': [2, 1, 9, 8, 7, 4],
    'col3': [0, 1, 9, 4, 2, 3],
    'col4': ['a', 'B', 'c', 'D', 'e', 'F'] })

# create a frovedis dataframe from pandas dataframe
fd_df = fdf.DataFrame(pd_df)
```

```
# display frovedis dataframe
fd_df.show()
```

Output

index	col1	col2	col3	col4
0	A	2	0	a
1	A	1	1	B
2	B	9	9	c
3	NULL	8	4	D
4	D	7	2	e
5	C	4	3	F

Sort by 'col1' on frovedis dataframe:

For example,

```
# Sort dataframe by 'col1'
fd_df.sort('col1').show()
```

Output

index	col1	col2	col3	col4
0	A	2	0	a
1	A	1	1	B
2	B	9	9	c
5	C	4	3	F
4	D	7	2	e
3	NULL	8	4	D

Sorting frovedis dataframe in ascending order (by default) by using multiple columns:

For example,

```
# Sort dataframe by multiple columns
fd_df.sort(['col1', 'col2']).show()
```

Output

index	col1	col2	col3	col4
1	A	1	1	B
0	A	2	0	a
2	B	9	9	c
5	C	4	3	F
4	D	7	2	e
3	NULL	8	4	D

NOTE: In the above example, in case of multiple columns sorting, ‘col1’ will be sorted first and ‘col2’ will only be considered for sorting in case of duplicate entries present in ‘col1’.

Sorting frovedis dataframe in descending orde by using ascending=False:

For example,

```
# Sort datafrme in descending order
fd_df.sort('col1', ascending = False).show()
```

Output

index	col1	col2	col3	col4
3	NULL	8	4	D
4	D	7	2	e
5	C	4	3	F
2	B	9	9	c
0	A	2	0	a
1	A	1	1	B

Return Value

It returns a new Frovedis DataFrame with sorted values.

54.1.3.4 4. DataFrame.sort_index(axis = 0, ascending = True, inplace = False, kind = ‘quicksort’, na_position = ‘last’)

Parameters

axis: It accepts an interger or a string object as parameter. It is the axis along which the sorting will be performed. (Default: 0)

When axis = 0 or axis = ‘index’, operation will be performed on rows. **Currently only axis = 0 or axis = ‘index’ is supported.**

ascending: It accepts a boolean value as parameter. This parameter decides the order to sort the data. (Default: True)

When this parameter is explicitly passed as False, it sorts the data into descending order.

inplace: It accepts a boolean value as paramter. To modify the original DataFrame object, argument is explicitly passed as True. Otherwise operation is performed on a copy of Frovedis DataFrame object. **Currently ‘inplace’ = True is not supported by this method.** (Default: False)

kind: It accepts a string object as parameter. This parameter is used to select the sorting algorithms. (Default: ‘quicksort’)

na_position: It accepts a string object as parameter. It is used to decide where to puts NaNs i.e at the beginning or at the end. (Default: ‘last’)

When na_position = ‘last’, it puts NaNs at the end. **Currently, na_position = ‘last’ is only supported for this method.**

Purpose

It is used to sort Frovedis DataFrame according to index values. It creates a new sorted DataFrame by the specified label.

Currently it only supports ‘radixsort’ and other values for ‘kind’ parameter are ignored internally along with a warning. Also, this method does not support MultiIndex yet.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame([1, 2, 3, 4, 5],
                      index=[100, 29, 234, 1, 150],
                      columns=['A'])

# creating frovedis dataframe from pandas dataframe
fd_df = fdf.DataFrame(pd_df)

# to display frovedis dataframe
fd_df.show()
```

Output

index	A
100	1
29	2
234	3
1	4
150	5

Sort values in dataframe, by default in ascending order (ascending=True):

For example,

```
# to display sorted dataframe by index
fd_df.sort_index().show()
```

Output

index	A
1	4
29	2
100	1
150	5
234	3

In order to sort in descending order, use ascending=False:

For example,

```
fd_df.sort_index(ascending=False)
```

Output

index	A
234	3
150	5
100	1


```
29      2
1      4
```

Return Value

It returns a new Frovedis DataFrame instance sorted by the labels.

54.1.3.5 5. DataFrame.sort_values(by, axis = 0, ascending = True, inplace = False, kind = 'radixsort', na_position = 'last')

Parameters

by: It accepts the name or list of names on which sorting will be applied.

If axis is 0 or 'index', then 'by' should contain index levels and/or column labels. **Currently axis = 1 or axis = 'columns' is not supported.**

axis: It accepts an integer or a string object as parameter. To perform sorting along rows or columns, it is selected by this parameter. (Default: 0)

If axis is 0 or 'index', sorting is performed along the row. **Currently axis = 1 or axis = 'columns' is not supported.**

ascending: It accepts a boolean value or a list of boolean values as parameter. The order of sorting is decided by this parameter. Need to specify a list of booleans for multiple sorting orders. If this is a list of booleans, the length of the list must match the length of the 'by' parameter list. By default, the order of sorting will be ascending and to change the order to descending, explicitly pass it as False. (Default: True)

inplace: It accepts a boolean value as parameter. To modify the original DataFrame object, argument is explicitly passed as True. Otherwise operation is performed on a copy of Frovedis DataFrame object. **Currently 'inplace' = True is not supported by this method.** (Default: False)

kind: It accepts a string object as parameter. The type of sorting algorithm is decided from this parameter. **Currently it only supports 'radixsort' and other values for 'kind' parameter are ignored internally along with a warning.** (Default: 'radixsort')

na_position: It accepts a string object as parameter. It is used to decide where to puts NaNs i.e at the beginning or at the end. (Default: 'last')

When na_position = 'last', it puts NaNs at the end. **Currently, it only supports na_position = 'last'.**

Purpose

To sort the DataFrame by the values along axis = 0 or 'index'.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# creating a pandas dataframe
pd_df = pd.DataFrame({
    'col1': ['A', 'A', 'B', np.nan, 'D', 'C'],
    'col2': [2, 1, 9, 8, 7, 4],
    'col3': [0, 1, 9, 4, 2, 3],
    'col4': ['a', 'B', 'c', 'D', 'e', 'F']
})

# creating a frovedis dataframe
fd_df = fdf.DataFrame(pd_df)

# displaying the frovedis dataframe
fd_df.show()
```

Output

index	col1	col2	col3	col4
0	A	2	0	a
1	A	1	1	B
2	B	9	9	c
3	NULL	8	4	D
4	D	7	2	e
5	C	4	3	F

Sort by 'col1' on frovedis dataframe:

For example,

```
fd_df.sort_values(by=['col1']).show() #Sort by col1
```

Output

index	col1	col2	col3	col4
0	A	2	0	a
1	A	1	1	B
2	B	9	9	c
5	C	4	3	F
4	D	7	2	e
3	NULL	8	4	D

Sorting frovedis dataframe in ascending order (by default) by using multiple columns:

For example,

```
fd_df.sort_values(by=['col1', 'col2'], ascending = [True, True]).show()
```

Output

index	col1	col2	col3	col4
1	A	1	1	B
0	A	2	0	a
2	B	9	9	c
5	C	4	3	F
4	D	7	2	e
3	NULL	8	4	D

Sorting frovedis dataframe in descending orde by using ascending=False:

For example,

```
# Sort in descending order
fd_df.sort_values(by='col1', ascending=False).show()
```

Output

index	col1	col2	col3	col4
3	NULL	8	4	D
4	D	7	2	e
5	C	4	3	F
2	B	9	9	c
0	A	2	0	a
1	A	1	1	B

Return Value

It returns a new Frovedis DataFrame with sorted values.

54.2 SEE ALSO

- **DataFrame - Introduction**
- **DataFrame - Generic Functions**
- **DataFrame - Conversion Functions**
- **DataFrame - Math Functions**
- **DataFrame - Aggregate Functions**

Chapter 55

DataFrame Aggregate Functions

55.1 NAME

DataFrame Aggregate Functions - list of all functions related to aggregate operations on frovedis dataframe are illustrated here.

55.1.1 DESCRIPTION

An essential piece of analysis of large data is efficient summarization: computing aggregations like `sum()`, `mean()`, `median()`, `min()`, and `max()`, which gives insight into the nature of a potentially large dataset. In this section, we'll explore list of all such aggregation operations done on frovedis dataframe.

Aggregation can be performed in **two ways** on frovedis dataframe:

- Either using `agg()`.
- Or using aggregation functions such as `min()`, `max()`, `median()`, `mode()`, etc. on frovedis dataframe.

55.1.2 Public Member Functions

1. `agg(func)`
2. `cov((min_periods = None, ddof = 1.0, low_memory = True, other = None)`
3. `mad(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)`
4. `max(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)`
5. `mean(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)`
6. `median(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)`
7. `min(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)`
8. `mode(axis = 0, numeric_only = False, dropna = True)`
9. `sem(axis = None, skipna = None, level = None, ddof = 1, numeric_only = None, **kwargs)`
10. `std(axis = None, skipna = None, level = None, ddof = 1, numeric_only = None, **kwargs)`
11. `sum(axis = None, skipna = None, level = None, numeric_only = None, min_count = 0, **kwargs)`
12. `var(axis = None, skipna = None, level = None, ddof = 1, numeric_only = None, **kwargs)`

55.1.3 Detailed Description

55.1.3.1 1. DataFrame.agg(func)

Parameters

func: Names of functions to use for aggregating the data. The input to be used with the function must be a frovedis DataFrame instance having atleast one numeric column.

Accepted combinations for this parameter are:

- A string function name such as 'max', 'min', etc.
- list of functions and/or function names, For example, ['max', 'mean'].
- dictionary with keys as column labels and values as function name or list of such functions.
For Example, {'Age': ['max', 'min', 'mean'], 'Ename': ['count']}

Purpose

It computes an aggregate operation based on the condition specified in 'func'.

Currently, this method will perform aggregation operation along the rows.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52

```
7      Abhi      32      Kanpur      B.Tech      NULL
```

Aggregate these functions over the rows, where func is a function string name:

For example,

```
print(fdf1.agg('max'))
```

Output

```
Name          nan
Age           36
City          nan
Qualification  nan
Score         52
Name: max, dtype: object
```

It displays a pandas dataframe containing numeric column(s) with newly computed aggregates of each groups.

Performing aggregation along the rows where func is a dictionary:

For example,

```
print(fdf1.agg({"Age": ["std", "mean"]}))
```

Output

```
          Age
mean  29.125000
std   4.853202
```

Aggregation along the rows where func is a list of functions:

For example,

```
print(fdf1['Age'].agg(['max', 'min', 'mean']))
```

Output

```
          Age
max   36.000
min   22.000
mean  29.125
```

Return Value

1. If one ‘func’ provided and ‘func’ is a string:

- It returns a pandas Series instance with numeric column(s) only, after aggregation function is completed.

2. If one or more ‘func’ provided and ‘func’ is list/dict of string:

- It returns a pandas DataFrame instance with numeric column(s) only, after aggregation function is completed.

55.1.3.2 2. DataFrame.cov(min_periods = None, ddof = 1.0, low_memory = True, other = None)

Parameters

min_periods: It accepts an integer as parameter. It specifies the minimum number of observations required per pair of columns to have a valid result. (Default: None)

When it is None (not specified explicitly), then **min_periods = 1**.

ddof: It accepts a float parameter that specifies the delta degrees of freedom. (Default: 1.0)

low_memory: It accepts boolean parameter that specifies whethet to enable memory optimised computation or time optimised computation. (Default: True)

other: It accepts frovedis dataframe as parameter, where it must be expressed in “df[col_name]” form. Also, it can be expressed in “df.col_name” form as well. (Default: None)

- **When it is not None (specified explicitly)**, it performs covariance operation between both the given frovedis dataframes. Although, the input dataframe must be used as expressions mentioned above.
- **When it is None (not specified explicitly)**, it will perform covarince on input dataframe to give covraince matrix represented as a dataframe.

Purpose

It computes pairwise covariance of columns, excluding missing values.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'Score': [23, 34, 35, 45, 23, 50, 52, 34]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Age	Score
0	27	23
1	24	34
2	22	35
3	32	45
4	33	23
5	36	50
6	27	52
7	32	34

Compute covariance on frovedis dataframe:

For example,

```
# cov() demo
fdf1.cov().show()
```

Output

index	Age	Score
-------	-----	-------


```
Age      23.5535  11
Score    11      124.571
```

It displays a covariance matrix as the frovedis DataFrame instance.

Using min_periods parameter to calculate covariance:

For example,

```
# cov() demo using min_periods = 8
fdf1.cov(min_periods = 8).show()
```

Output

```
index  Age      Score
Age     NULL     NULL
Score  NULL     NULL
```

Using ddof parameter to calculate covariance:

For example,

```
# cov() demo using ddof = 2
fdf1.cov(ddof = 2).show()
```

Output

```
index  Age      Score
Age     27.4791  12.8333
Score   12.8333 145.333
```

In the below example, while using ‘other’ parameter, both inputs must be frovedis series. Also, the output returned by this method will be a float value.

For example,

```
# create another dataframe
pdf2 = pd.DataFrame({'Score': [51, 34, 33, 45, 12, 82, 67, 91]})
fdf2 = fdf.DataFrame(pdf2)

# cov() demo using 'other' parameter
print(fdf1['Score'].cov(other = fdf2['Score']))
```

Output

```
158.28571428571428
```

Here, it could also be expressed as “`fdf1['Score'].cov(other = fdf2['Score'])`”.

Note:- While using input dataframe in the form ‘`fdf1['Score']`’ or ‘`fdf1.Score`’, ‘other’ parameter must be provided.

Return Value

- **If other = None:**
It returns a covariance matrix represented as frovedis DataFrame instance.
- **If other != None:**
It returns covariance as scalar value.

55.1.3.3 3. DataFrame.mad(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform mean absolute deviation along the columns or rows. (Default: None)

- **0 or 'index'**: perform mean absolute deviation along the indices.
- **1 or 'columns'**: perform mean absolute deviation along the columns.

When it is None (not specified explicitly), it performs mean absolute deviation along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during mean absolute deviation computation.

level: This is an unused parameter. (Default: None)

numeric_only: This is an unused parameter. (Default: None)

****kwargs**: Additional keyword arguments to be passed to the function.

Purpose

It computes the mean absolute deviation of the values over the requested axis.

Currently, mean absolute deviation will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.mad(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Mean absolute deviation along the rows (by default):

For example,

```
# mad() demo, axis = 0 by default
fdf1.mad().show()
```

Output

```
index  mad
Age    4.125
Score  9.16666
```

It displays a frovedis dataframe with numeric column(s) containing the newly computed mean absolute deviation for each column.

Also, it excludes the missing value in ‘**Score**’ column while computing the mean absolute deviation.

Mean absolute deviation along the rows and using skipna parameter:

For example,

```
# mad() demo using skipna = False
fdf1.mad(skipna = False).show()
```

Output

```
index  mad
Age    4.125
Score  NULL
```

Here, it includes the missing value in ‘**Score**’ column while computing the mean absolute deviation.

Mean absolute deviation along the columns:

For example,

```
# mad() demo using axis = 1
fdf1.mad(axis = 1).show()
```

Output

```
index  mad
0      2
1      5
2     6.5
3     6.5
4      0
5      7
6    12.5
7      0
```

Mean absolute deviation along the columns and using skipna parameter:

For example,

```
# mad() demo using axis = 1 and skipna = False
fdf1.mad(axis = 1, skipna = False).show()
```

Output

```
index    mad
0         2
1         5
2        6.5
3        6.5
4        NULL
5         7
6       12.5
7        NULL
```

Return Value

It returns a frovedis DataFrame instance with the result of the specified aggregate operation.

55.1.3.4 4. DataFrame.max(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform maximum operation along the columns or rows. (Default: None)

- **0 or 'index':** perform maximum operation along the indices to get the maximum value.
- **1 or 'columns':** perform maximum operation along the columns to get the maximum value.

When it is None (not specified explicitly), it performs maximum operation along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during maximum value computation.

level: This is an unused parameter. (Default: None)

numeric_only: This is an unused parameter. (Default: None)

****kwargs:** Additional keyword arguments to be passed to the function.

Purpose

It computes the maximum of the values over the requested axis.

Currently, maximum operation will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.max(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
```

```

peopleDF = {
    'Name':['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age':[27, 24, 22, 32, 33, 36, 27, 32],
    'City':['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification':['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Max operation along the rows (by default):

For example,

```

# max() demo
fdf1.max().show()

```

Output

index	max
Age	36
Score	52

It displays a frovedis dataframe with numeric column(s) containing the newly computed maximum value for each column.

Also, it excludes the missing value in ‘Score’ column while computing the maximum value.

Max operation along the rows and using skipna parameter:

For example,

```

# max() demo using skipna = False
fdf1.max(skipna = False).show()

```

Output

index	max
Age	36
Score	NULL

Here, it includes the missing value in ‘**Score**’ column while computing the maximum value.

Max operation along the columns:

For example,

```
# max() demo using axis = 1
fdf1.max(axis = 1).show()
```

Output

index	max
0	27
1	34
2	35
3	45
4	33
5	50
6	52
7	32

Max operation along the columns and using skipna parameter:

For example,

```
# max() demo using axis = 1 and skipna = False
fdf1.max(axis = 1, skipna = False).show()
```

Output

index	max
0	27
1	34
2	35
3	45
4	NULL
5	50
6	52
7	NULL

Return Value

It returns a frovedis DataFrame instance with the result of the specified aggregate operation.

55.1.3.5 5. DataFrame.mean(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform mean along the columns or rows. (Default: None)

- **0** or ‘**index**’: perform mean along the indices.

- **1** or ‘**columns**’: perform mean along the columns.

When it is None (not specified explicitly), it performs mean operation along the rows.

skipna: It is a boolean parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during mean computation.

level: This is an unused parameter. (Default: None)

numeric_only: This is an unused parameter. (Default: None)

****kwargs**: Additional keyword arguments to be passed to the function.

Purpose

It computes mean of the values over the requested axis.

Currently, mean will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.mean(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Mean computation along the rows (by default):

For example,

```
# mean() demo
fdf1.mean(),show()
```

Output

index	mean
Age	29.125
Score	39.8333

It displays a `frovedis` dataframe with numeric column(s) containing the newly computed mean for each column.

Also, it excludes the missing value in ‘**Score**’ column while computing the mean.

Mean computation along the rows and using `skipna` parameter:

For example,

```
# mean() demo using skipna = False
fdf1.mean(skipna = False).show()
```

Output

index	mean
Age	29.125
Score	NULL

Here, it includes the missing value in ‘**Score**’ column while computing the mean.

Max operation along the columns:

For example,

```
# mean() demo using axis = 1
fdf1.mean(axis = 1).show()
```

Output

index	mean
0	25
1	29
2	28.5
3	38.5
4	33
5	43
6	39.5
7	32

Max operation along the columns and using `skipna` parameter:

For example,

```
# mean() demo using axis = 1 and skipna = False
fdf1.mean(axis = 1, skipna = False).show()
```

Output

index	mean
0	25
1	29
2	28.5
3	38.5
4	NULL
5	43
6	39.5
7	NULL

Return Value

It returns a `frovedis DataFrame` instance.

55.1.3.6 6. DataFrame.median(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform median operation along the columns or rows. (Default: None)

- **0 or 'index':** perform median operation along the indices.
- **1 or 'columns':** perform median operation along the columns.

When it is None (not specified explicitly), it performs median operation along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during median computation.

level: This is an unused parameter. (Default: None)

numeric_only: This is an unused parameter. (Default: None)

****kwargs:** Additional keyword arguments to be passed to the function.

Purpose

It computes median of the values over the requested axis.

Currently, median will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.median(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23

1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Median computation along the rows (by default):

For example,

```
# median() demo
fdf1.median().show()
```

Output

```
index  median
Age    29.5
Score  40
```

It displays a frovedis dataframe with numeric column(s) containing the newly computed median for each column.

Also, it excludes the missing value in ‘**Score**’ column while computing the median.

Median computation along the rows and using skipna parameter:

For example,

```
# median() demo using skipna = False
fdf1.median(skipna = False).show()
```

Output

```
index  median
Age    29.5
Score  NULL
```

Here, it includes the missing value in ‘**Score**’ column while computing the median.

Median computation along the columns:

For example,

```
# median() demo using axis = 1
fdf1.median(axis = 1).show()
```

Output

```
index  median
0      25
1      29
2     28.5
3     38.5
4      33
5      43
6     39.5
7      32
```

Median computation along the columns and using skipna parameter:

For example,

```
# median() demo using axis = 1 and skipna = False
fdf1.median(axis = 1, skipna = False).show()
```

Output

```
index    median
0         25
1         29
2        28.5
3        38.5
4         NULL
5         43
6        39.5
7         NULL
```

Return Value

It returns a frovedis DataFrame instance.

55.1.3.7 7. DataFrame.min(axis = None, skipna = None, level = None, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform minimum operation along the columns or rows. (Default: None)

- **0 or 'index'**: perform minimum operation along the indices to get the minimum value.
- **1 or 'columns'**: perform minimum operation along the columns to get the minimum value.

When it is None (not specified explicitly), it performs minimum operation along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during minimum value computation.

level: This is an unused parameter. (Default: None)

numeric_only: This is an unused parameter. (Default: None)

****kwargs**: Additional keyword arguments to be passed to the function.

Purpose

It computes the minimum of the values over the requested axis.

Currently, minimum operation will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.min(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name':['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age':[27, 24, 22, 32, 33, 36, 27, 32],
```

```

    'City':['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification':['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Min operation along the rows (by default):

For example,

```

# min() demo
fdf1.min().show()

```

Output

index	min
Age	22
Score	23

It displays a frovedis dataframe with numeric column(s) containing the newly computed minimum value for each column.

Also, it excludes the missing value in ‘**Score**’ column while computing the minimum value.

Min operation along the rows and using skipna parameter:

For example,

```

# min() demo using skipna = False
fdf1.min(skipna = False).show()

```

Output

index	min
Age	22
Score	NULL

Here, it includes the missing value in ‘**Score**’ column while computing the median.

Min operation along the columns:

For example,

```
# min() demo using axis = 1
fdf1.min(axis = 1).show()
```

Output

index	min
0	23
1	24
2	22
3	32
4	33
5	36
6	27
7	32

Min operation along the columns and using skipna parameter:

For example,

```
# min() demo using axis = 1 and skipna = False
fdf1.min(axis = 1, skipna = False).show()
```

Output

index	min
0	23
1	24
2	22
3	32
4	NULL
5	36
6	27
7	NULL

Return Value

It returns a `provdis DataFrame` instance with the result of the specified aggregate operation.

55.1.3.8 8. `DataFrame.mode(axis = 0, numeric_only = False, dropna = True)`

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform standard error of the mean along the columns or rows. (Default: 0)

- **0 or 'index':** perform mode along the indices.
- **1 or 'columns':** perform mode along the columns.

When it is `None` (not specified explicitly), it performs standard error of the mean along the rows.

numeric_only: It accepts string object as parameter. If `True`, mode operation will result in a dataframe having only numeric columns. Otherwise, it will result in a dataframe having both numeric and non-numeric columns. (Default: `False`)

dropna: It accepts boolean as parameter. When set to `True`, it will exclude missing values while computing the result of mode operation. (Default: `True`)

When it is `None` (not specified explicitly), it excludes missing values during mode computation.

Purpose

This method gets the mode(s) of each element along the selected axis.

The mode of a set of values is the value that appears most often. It can be multiple values.

When input with non-numeric columns is used with `mode()`, then it ignores the non-numeric columns for mode computation.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Mode computation along the rows (by default):

For example,

```
# mode() demo
fdf1.mode().show()
```

Output

index	Score	Name	Age	City	Qualification
0	23	Anuj	27	Kanpur	B.Tech
1	34	Jai	32	NULL	Phd
2	35	Princi	NULL	NULL	NULL
3	45	NULL	NULL	NULL	NULL
4	50	NULL	NULL	NULL	NULL

```
5      52      NULL      NULL      NULL      NULL
```

Mode will be calculated for dataframe having string and numeric columns when `axis = 0` or `'index'`.

Also, resultant dataframe has both numeric and non-numeric columns.

Mode computation along the rows and using `numeric_only` parameter:

For example,

```
# mode() demo using numeric_only = True
fdf1.mode(numeric_only = True).show()
```

Output

index	Score	Age
0	23	27
1	34	32
2	35	NULL
3	45	NULL
4	50	NULL
5	52	NULL

Here, resultant dataframe has only numeric columns.

Mode computation along the rows and using `dropna` parameter:

For example,

```
# mode() demo using dropna = False
fdf1.mode(dropna = False).show()
```

Output

index	Name	Age	City	Qualification	Score
0	Anuj	27	Kanpur	B.Tech	NULL
1	Jai	32	NULL	Phd	NULL
2	Princi	NULL	NULL	NULL	NULL

For `axis = 1` or `'columns'`, mode will be calculated for dataframe having only numeric columns.

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
bmiDF = {
    'height': [157, 124, 162, np.nan, 133, 176, np.nan, 152],
    'weight': [53, 64, np.nan, 65, 63, 80, np.nan, 84]
}

# create pandas dataframe with only numeric columns
pdf1 = pd.DataFrame(bmiDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	height	weight
0	157	53
1	124	64
2	162	NULL
3	NULL	65
4	133	63
5	176	80
6	NULL	NULL
7	152	84

Mode computation along the columns:

For example,

```
# mode() demo using axis = 1
fdf1.mode(axis = 1).show()
```

Output

index	0
0	157
1	124
2	162
3	65
4	133
5	176
6	0
7	152

Mode computation along the columns and using dropna parameter:

For example,

```
# mode() demo using axis = 1 and dropna = False
fdf1.mode(axis = 1, dropna = False).show()
```

Output

index	0
0	157
1	124
2	162
3	NULL
4	133
5	176
6	NULL
7	152

Return Value

- **If `numeric_only = False`:**
It returns a `frovedis DataFrame` instance having both numeric and non-numeric columns (if any).
- **If `numeric_only = True`:**
It returns a `frovedis DataFrame` instance having only numeric columns.

55.1.3.9 9. DataFrame.sem(axis = None, skipna = None, level = None, ddof = 1, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform standard error of the mean along the columns or rows. (Default: None)

- **0 or 'index':** perform standard error of the mean along the indices.
- **1 or 'columns':** perform standard error of the mean along the columns.

When it is None (not specified explicitly), it performs standard error of the mean along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during standard error of the mean computation.

level: This is an unused parameter. (Default: None)

ddof: It accepts an integer parameter that specifies the delta degrees of freedom. (Default: 1)

numeric_only: This is an unused parameter. (Default: None)

****kwargs:** Additional keyword arguments to be passed to the function.

Purpose

It computes standard error of the mean over requested axis.

Currently, standard error of the mean will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.sem(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Standard error of the mean operation along the rows (by default):

For example,

```
# sem() demo
fdf1.sem().show()
```

Output

```
index    sem
Age      1.71586
Score    4.54178
```

It displays a frovedis dataframe with numeric column(s) containing the newly computed standard error of the mean for each column.

Standard error of the mean operation along the rows and using ddof parameter:

For example,

```
# sem() demo using ddof = 2
fdf1.sem(ddof = 2).show()
```

Output

```
index    sem
Age      1.85334
Score    5.07786
```

Standard error of the mean operation along the rows and using skipna parameter:

For example,

```
# sem() demo using skipna = False
fdf1.sem(skipna = False).show()
```

Output

```
index    sem
Age      1.71586
Score    NULL
```

Standard error of the mean operation along the columns:

For example,

```
# sem() demo using axis = 1
fdf1.sem(axis = 1).show()
```

Output

```
index    sem
0        2
1        5
```

```

2      6.49999
3      6.49999
4      NULL
5      6.99999
6      12.5
7      NULL

```

Standard error of the mean operation along the columns and using skipna parameter:

For example,

```

# sem() demo using axis = 1 and skipna = False
fdf1.sem(axis = 1, skipna = False).show()

```

Output

```

index    sem
0         2
1         5
2      6.49999
3      6.49999
4      NULL
5      6.99999
6      12.5
7      NULL

```

Return Value

It returns a `frovedis DataFrame` instance with the result of the specified aggregate operation.

55.1.3.10 10. `DataFrame.std(axis = None, skipna = None, level = None, ddof = 1, numeric_only = None, **kwargs)`

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform standard deviation along the columns or rows. (Default: None)

- **0 or 'index':** perform standard deviation along the indices.
- **1 or 'columns':** perform standard deviation along the columns.

When it is None (not specified explicitly), it performs standard deviation along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during standard deviation computation.

level: This is an unused parameter. (Default: None)

ddof: It accepts an integer parameter that specifies the delta degrees of freedom. (Default: 1)

numeric_only: This is an unused parameter. (Default: None)

****kwargs:** Additional keyword arguments to be passed to the function.

Purpose

It computes standard deviation over requested axis.

Currently, standard deviation will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas `DataFrame.std()`. These are not used internally in `frovedis`.

Creating `frovedis DataFrame` from pandas `DataFrame`:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Standard deviation computation along the rows (by default):

For example,

```
# std() demo
fdf1.std().show()
```

Output

index	std
Age	4.8532
Score	11.125

It displays a frovedis dataframe with numeric column(s) containing the newly computed standard deviation for each column.

Standard deviation computation along the rows and using skipna parameter:

For example,

```
# std() demo using skipna = False
fdf1.std(skipna = False).show()
```

Output

```
index    std
Age      4.8532
Score    NULL
```

Standard deviation computation along the rows and using ddof parameter:

For example,

```
# std() demo using ddof = 2
fdf1.std(ddof = 2).show()
```

Output

```
index    std
Age      5.24205
Score    12.4381
```

Standard deviation computation along the columns:

For example,

```
# std() demo using axis = 1
fdf1.std(axis = 1).show()
```

Output

```
index    std
0        2.82842
1        7.07106
2        9.19238
3        9.19238
4        NULL
5        9.89949
6        17.6776
7        NULL
```

Standard deviation computation along the columns and using skipna parameter:

For example,

```
# std() demo using axis = 1 and skipna = False
fdf1.std(axis = 1, skipna = False).show()
```

Output

```
index    std
0        2.82842
1        7.07106
2        9.19238
3        9.19238
4        NULL
5        9.89949
6        17.6776
7        NULL
```

Return Value

It returns a `frovedis DataFrame` instance with the result of the specified aggregate functions.

55.1.3.11 11. DataFrame.sum(axis = None, skipna = None, level = None, numeric_only = None, min_count = 0, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform summation operation along the columns or rows. (Default: None)

- **0 or 'index':** perform summation operation along the indices.
- **1 or 'columns':** perform summation operation along the columns.

When it is None (not specified explicitly), it performs summation operation along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during summation computation.

level: This is an unused parameter. (Default: None)

numeric_only: This is an unused parameter. (Default: None)

min_count: It is an integer, float or double (float64) parameter that specifies the minimum number of values that needs to be present to perform the action. (Default: 0)

****kwargs:** Additional keyword arguments to be passed to the function.

Purpose

It computes the sum of the values over the requested axis.

Currently, summation will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.sum(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
-------	------	-----	------	---------------	-------

0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Sum operation along the rows (by default):

For example,

```
# sum() demo
fdf1.sum().show()
```

Output

```
index    sum
Age      233
Score    239
```

It displays a frovedis dataframe with numeric column(s) containing the newly computed summation for each column.

Also, it excludes the missing value in ‘Score’ column while computing summation.

Sum operation along the rows and using skipna parameter:

For example,

```
# sum() demo using skipna = False
fdf1.sum(skipna = False).show()
```

Output

```
index    sum
Age      233
Score    NULL
```

Here, it includes the missing value in ‘Score’ column while computing the sum.

Sum operation along the columns:

For example,

```
# sum() demo using axis = 1
fdf1.sum(axis = 1).show()
```

Output

```
index    sum
0        50
1        58
2        57
3        77
4        33
5        86
6        79
7        32
```

Sum operation along the columns and using skipna parameter:

For example,

```
# sum() demo using axis = 1 and skipna = False
fdf1.sum(axis = 1, skipna = False).show()
```

Output

```
index    sum
0         50
1         58
2         57
3         77
4        NULL
5         86
6         79
7        NULL
```

Return Value

It returns a frovedis DataFrame instance with the result of the specified aggregate functions.

55.1.3.12 12. DataFrame.var(axis = None, skipna = None, level = None, ddof = 1, numeric_only = None, **kwargs)

Parameters

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform variance along the columns or rows. (Default: None)

- **0 or 'index':** perform variance along the indices.
- **1 or 'columns':** perform variance along the columns.

When it is None (not specified explicitly), it performs variance along the rows.

skipna: It accepts boolean as parameter. When set to True, it will exclude missing values while computing the result. (Default: None)

When it is None (not specified explicitly), it excludes missing values during variance computation.

level: This is an unused parameter. (Default: None)

ddof: It accepts an integer parameter that specifies the delta degrees of freedom. (Default: 1)

numeric_only: This is an unused parameter. (Default: None)

****kwargs:** Additional keyword arguments to be passed to the function.

Purpose

It computes variance over requested axis.

Currently, variance will be calculated for dataframe having atleast one numeric columns.

The parameters: “level”, “numeric_only”, “**kwargs” are simply kept in to make the interface uniform to the pandas DataFrame.var(). These are not used internally in frovedis.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name':['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age':[27, 24, 22, 32, 33, 36, 27, 32],
    'City':['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
```



```

        'Kanpur', 'Kanpur', 'Kanpur'],
        'Qualification':['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
        'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
    }

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Variance computation along the rows (by default):

For example,

```

# var() demo
fdf1.var().show()

```

Output

index	var
Age	23.5535
Score	123.766

It displays a frovedis dataframe with numeric column(s) containing the newly computed variance for each column.

Also, it excludes the missing value in ‘**Score**’ column while computing variance.

Variance computation along the rows and using skipna parameter:

For example,

```

# var() demo using skipna = False
fdf1.var(skipna = False).show()

```

Output

index	var
Age	23.5535
Score	NULL

Variance computation along the rows and using ddof parameter:

For example,

```

# var() demo using ddof = 2

```

```
fdf1.var(ddof = 2).show()
```

Output

```
index    var
Age      27.4791
Score    154.708
```

Variance computation along the columns:

For example,

```
# var() demo using axis = 1
fdf1.var(axis = 1).show()
```

Output

```
index    var
0         8
1        50
2       84.5
3       84.5
4       NULL
5        98
6      312.5
7       NULL
```

Variance computation along the columns and using skipna parameter:

For example,

```
# var() demo using axis = 1 and skipna = False
fdf1.var(axis = 1, skipna = False).show()
```

Output

```
index    var
0         8
1        50
2       84.5
3       84.5
4       NULL
5        98
6      312.5
7       NULL
```

Return Value

It returns a `frovedis DataFrame` instance with the result of the specified aggregate operation.

55.2 SEE ALSO

- **DataFrame - Introduction**
- **DataFrame - Generic Functions**
- **DataFrame - Conversion Functions**
- **DataFrame - Sorting Functions**

- **DataFrame - Math Functions**

Chapter 56

DataFrame Math Functions

56.1 NAME

DataFrame Math Functions - this manual contains all the methods for carrying out mathematical operations.

56.1.1 DESCRIPTION

Frovedis dataframe has several math functions defined for performing operations like `add()`, `sub()`, `mul()`, etc. between two dataframes or between scalar value and dataframe. These functions return a new frovedis DataFrame instance as a result.

Also, it contains reverse operations such as `radd()`, `rsub()`, `rmul()`, etc.

However, there are some special cases while using frovedis dataframe with mathematical operations as mentioned below:

Binary operation on two frovedis dataframes having same columns but different datatypes:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

pdf1 = pd.DataFrame(data1)
fdf1 = fdf.DataFrame(pdf1)

data2 = {
    "points": [2., 3., 8.],
    "total": [7, 9, 11]
}

pdf2 = pd.DataFrame(data2)
fdf2 = fdf.DataFrame(pdf2)

print(fdf1 + fdf2)
```

Output

index	points	total
0	7	17
1	9	20
2	12	23

Here, ‘points’ column in first dataframe is **int type** and in other dataframe is **float type**.

Type conversion occurs for ‘points’ column in resultant dataframe.

```
fdf1.points(int) + fdf2.points(float) -> res.points(float)
```

Binary operation on two frovedis dataframes having atleast one common column:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}
```

```
pdf1 = pd.DataFrame(data1)
fdf1 = fdf.DataFrame(pdf1)
```

```
data2 = {
    "score": [2, 3, 8],
    "total": [7, 9, 11]
}
```

```
pdf2 = pd.DataFrame(data2)
fdf2 = fdf.DataFrame(pdf2)
```

```
print(fdf1 + fdf2)
```

Output

index	points	score	total
0	NULL	NULL	17
1	NULL	NULL	20
2	NULL	NULL	23

Here, the resultant dataframe will contains all columns from input frovedis dataframes lexicographically.

Also, binary operation (i.e. addition) is performed on the common column only.

Binary operation on two frovedis dataframes having same columns. Also, same indices but in different order:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}
```

```
pdf1 = pd.DataFrame(data1, index = [1,2,3])
```

```
fdf1 = fdf.DataFrame(pdf1)

data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

pdf2 = pd.DataFrame(data2, index = [2,3,1])
fdf2 = fdf.DataFrame(pdf2)

print(fdf1 + fdf2)
```

Output

index	points	total
1	13	21
2	8	18
3	7	21

Binary operation (i.e. addition) will be performed between same indices for each column irrespective of the index order.

Currently, binary operation on two frovedis dataframes having same columns but different indices is not supported.

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

pdf1 = pd.DataFrame(data1, index = [0,1,2])
fdf1 = fdf.DataFrame(pdf1)

data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

pdf2 = pd.DataFrame(data2, index = [0,1,3])
fdf2 = fdf.DataFrame(pdf2)

print(fdf1 + fdf2)
```

In this case, dataframes have same column but indices are different. This will raise an exception in frovedis.

Binary operation between columns of frovedis dataframes. Also, having same indices:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}
```

```
pdf1 = pd.DataFrame(data1, index = [1,2,3])
fdf1 = fdf.DataFrame(pdf1)

data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

pdf2 = pd.DataFrame(data2, index = [1,2,3])
fdf2 = fdf.DataFrame(pdf2)

print(fdf1['points'] + fdf2['points'])
```

Output

```
index    (points+points)
1         7
2         9
3        12
```

Binary operation between two frovedis dataframes having single column (common) only. Also, having same indices but different order:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
pdf1 = pd.Series([27, 24, 22, 32, 33, 36, 27, 32],index = [0,1,2,3,4,5,6,7])
pdf2 = pd.Series([23, 34, 35, 45, 23, 50, 52, 34],index = [2,3,1,0,7,4,5,6])

fdf1 = fdf.DataFrame(pdf1)
fdf2 = fdf.DataFrame(pdf2)

print(fdf1 + fdf2)
```

Output

```
index    (0+0)
0         72
1         59
2         45
3         66
4         83
5         88
6         61
7         55
```

Binary operation (i.e. addition) will be performed between same indices irrespective of the index order in the frovedis dataframe.

Currently, binary operation on two frovedis dataframes single column only but different indices is not supported.

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
pdf1 = pd.Series([27, 24, 22, 32, 33, 36, 27, 32],index = [0,1,2,3,4,5,6,7])
```



```
pdf2 = pd.Series([23, 34, 35, 45, 23, 50, 52, 34],index = [2,3,1,0,8,4,5,9])

fdf1 = fdf.DataFrame(pdf1)
fdf2 = fdf.DataFrame(pdf2)

print(fdf1 + fdf2)
```

In this case, resultant dataframe has single column but indices are different. This will raise an exception in frovedis.

Binary operation between frovedis dataframe and scalar value (float type):

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

pdf1 = pd.DataFrame(data1)
fdf1 = fdf.DataFrame(pdf1)

print(fdf1 + 12.)
```

Output

index	points	total
0	17	22
1	18	23
2	16	24

Here, binary operation (i.e. addition) will be performed between the scalar value of float type on each column of the resultant frovedis dataframe.

Binary operation between frovedis dataframe and row vector:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

pdf1 = pd.DataFrame(data1)
fdf1 = fdf.DataFrame(pdf1)

print(fdf1 + [51, 34])
```

Output

index	points	total
0	56	44
1	57	45
2	55	46

Here, binary operation (i.e addition) will be performed between each element of row vector with each column on the given index of the frovedis dataframe.

Binary operation between frovedis dataframe and column vector:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

pdf1 = pd.DataFrame(data1)
fdf1 = fdf.DataFrame(pdf1)

print(fdf1['points'] + [21, 34, 45])
print(fdf1['total'] + [21, 34, 45])
```

Output

```
index  points
0      26
1      40
2      49

index  total
0      31
1      45
2      57
```

Here, binary operation (i.e addition) will be performed between the array and the given column of frovedis dataframe.

56.1.2 Public Member Functions

1. `abs()`
2. `add(other, axis = 'columns', level = None, fill_value = None)`
3. `div(other, axis = 'columns', level = None, fill_value = None)`
4. `floordiv(other, axis = 'columns', level = None, fill_value = None)`
5. `mod(other, axis = 'columns', level = None, fill_value = None)`
6. `mul(other, axis = 'columns', level = None, fill_value = None)`
7. `pow(other, axis = 'columns', level = None, fill_value = None)`
8. `sub(other, axis = 'columns', level = None, fill_value = None)`
9. `truediv(other, axis = 'columns', level = None, fill_value = None)`
10. `radd(other, axis = 'columns', level = None, fill_value = None)`
11. `rdiv(other, axis = 'columns', level = None, fill_value = None)`
12. `rfloordiv(other, axis = 'columns', level = None, fill_value = None)`
13. `rmod(other, axis = 'columns', level = None, fill_value = None)`
14. `rmul(other, axis = 'columns', level = None, fill_value = None)`
15. `rpow(other, axis = 'columns', level = None, fill_value = None)`
16. `rsub(other, axis = 'columns', level = None, fill_value = None)`
17. `rtruediv(other, axis = 'columns', level = None, fill_value = None)`

56.1.3 Detailed Description

56.1.3.1 1. DataFrame.abs()

Purpose

It computes absolute numeric value of each element.

This function only applies to elements that are all numeric.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
tempDF = {
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Temperature': [-2, 10, 18, 34, -8, -4, 36, 45]
}

# create pandas dataframe
pdf1 = pd.DataFrame(tempDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	City	Temperature
0	Nagpur	-2
1	Kanpur	10
2	Allahabad	18
3	Kannuaj	34
4	Allahabad	-8
5	Kanpur	-4
6	Kanpur	36
7	Kanpur	45

Absolute numeric values in a frovedis dataframe:

For example,

```
# abs() demo
print(fdf1['Temperature'].abs())
```

Output

index	Temperature
0	2
1	10
2	18
3	34

4	8
5	4
6	36
7	45

Return Value

It returns a frovedis DataFrame instance.

56.1.3.2 2. DataFrame.add(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be added with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform addition operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform addition operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs addition between two operands. It is equivalent to **'dataframe + other'**.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)
```

```
# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Add a scalar value using operator version:

For example,

```
print(fdf1 + 10)
```

Output

index	points	total
0	15	20
1	16	21
2	14	22

Add a scalar value using method version:

For example,

```
fdf1.add(10).show()
```

Output

index	points	total
0	15	20
1	16	21
2	14	22

In both versions, all column elements (axis = 1 by default) are added with a scalar value.

Creating two frovedis dataframes to perform addition:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
```

```
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Add two dataframes using operator version:

For example,

```
print(fdf1 + fdf2)
```

Output

index	points	total
0	7	17
1	9	20
2	12	23

Add two dataframes using method version:

For example,

```
fdf1.add(other = fdf2).show()
```

Output

index	points	total
0	7	17
1	9	20
2	12	23

In both versions, only common columns in both dataframes are added. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes in order to use fill_value parameter during addition:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Add two dataframes and using fill_value parameter:

For example,

```
# add() demo on two dataframes using method version and fill_value = 10
fdf1.add(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	7	17
1	9	21
2	12	20

Here, only common columns in both dataframes are added, excluding the missing values. Other column elements are added with the `fill_value = 10` (excluding missing values) in resultant dataframe.

Return Value

It returns a `frovedis DataFrame` which contains the result of arithmetic operation.

56.1.3.3 3. `DataFrame.div(other, axis = 'columns', level = None, fill_value = None)`

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- `frovedis DataFrame`. It must not be an empty dataframe.

Any of these is considered as the value to be divided over the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform division operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns'**: perform division operation on the columns. Currently, `axis = 1` is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs floating division operation between two operands. It is equivalent to '`dataframe / other`'.

It is an alias of `truediv()`.

Creating `frovedis DataFrame` from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}
```



```
# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Divide a scalar value using operator version:

For example,

```
print(fdf1 / 10)
```

Output

index	points	total
0	0.5	1
1	0.6	1.1
2	0.4	1.19999

Divide a scalar value using method version:

For example,

```
fdf1.div(10).show()
```

Output

index	points	total
0	0.5	1
1	0.6	1.1
2	0.4	1.19999

In both versions, all column elements (axis = 1 by default) are divided by a scalar value.

Creating two frovedis dataframes to perform division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
```

```
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Divide two dataframes using operator version:

For example,

```
print(fdf1 / fdf2)
```

Output

index	points	total
0	2.5	1.42857
1	2	1.22222
2	0.5	1.0909

Divide two dataframes using method version:

For example,

```
fdf1.div(other = fdf2).show()
```

Output

index	points	total
0	2.5	1.42857
1	2	1.22222
2	0.5	1.0909

In both versions, only common columns in both dataframes are divided. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes in order to use fill_value parameter during division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Divide two dataframes and using fill_value parameter:

For example,

```
# div() demo on two dataframes using method version and fill_value = 10
fdf1.div(other = fdf2, fill_value = 10).show()
```

Output

```
index  points  total
0      2.5    1.42857
1       2     1.1
2      0.5     1
```

Here, only common columns in both dataframes are divided, excluding the missing values. Other column elements are divided with the `fill_value = 10` (excluding the missing values) in resultant dataframe.

Return Value

It returns a `frovedis DataFrame` which contains the result of arithmetic operation.

56.1.3.4 4. `DataFrame.floordiv(other, axis = 'columns', level = None, fill_value = None)`

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be divided over the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform division operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform division operation on the columns. Currently, `axis = 1` is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs integer division operation between operands. It is equivalent to `'dataframe // other'`.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
```

```

        "points": [5, 6, 4],
        "total": [10, 11, 12]
    }

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Floor Division on a scalar value using operator version:

For example,

```
print(fdf1 // 10)
```

Output

index	points	total
0	0	1
1	0	1
2	0	1

Floor Division on a scalar value using method version:

For example,

```
fdf1.floordiv(10).show()
```

Output

index	points	total
0	0	1
1	0	1
2	0	1

In both versions, all column elements (axis = 1 by default) are divided by a scalar value. Also, resultant dataframe column elements will contain floor integer value.

Creating two frovedis dataframes to perform floor division:

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

```

```
# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Floor Division on two dataframes using operator version:

For example,

```
print(fdf1 // fdf2)
```

Output

index	points	total
0	2	1
1	2	1
2	0	1

Floor Division on two dataframes using method version:

For example,

```
fdf1.floordiv(other = fdf2).show()
```

Output

index	points	total
0	2	1
1	2	1
2	0	1

In both versions, only common columns in both dataframes are divided. Other are replaced with NaN values in resultant dataframe (`fill_value = None` by default).

Creating two frovedis dataframes in order to use `fill_value` parameter during floor division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL

2 8 NULL

Floor Division on two dataframes and using fill_value parameter:

For example,

```
# floordiv() demo on two dataframes using method version and fill_value = 10
fdf1.floordiv(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	2	1
1	2	1
2	0	1

Here, only common columns in both dataframes are divided, excluding the missing values. Other column elements are divided with the fill_value = 10 (excluding the missing values) in resultant dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.5 5. DataFrame.mod(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to perform modulo operation with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform modulo operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform modulo operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs modulo operation between two operands. It is equivalent to 'dataframe % other'.

Creating frovedis DataFrame from pandas DataFrame:

For example,


```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame()

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Modulo on a scalar value using operator version:

For example,

```
print(fdf1 % 10)
```

Output

index	points	total
0	5	0
1	6	1
2	4	2

Modulo on a scalar value using method version:

For example,

```
fdf1.mod(10).show()
```

Output

index	points	total
0	5	0
1	6	1
2	4	2

In both versions, modulo operation is performed on all column elements (axis = 1 by default) by a scalar value.

Creating two frovedis dataframes to perform modulo:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
```

```

data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```

# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()

```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Modulo on two dataframes using operator version:

For example,

```
print(fdf1 % fdf2)
```

Output

index	points	total
0	1	3
1	0	2
2	4	1

Creating two frovedis dataframes to perform modulo, use `fill_value` parameter too:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Modulo on two dataframes using method version:

For example,

```
fdf1.mod(other = fdf2).show()
```

Output

index	points	total
-------	--------	-------

0	1	3
1	0	2
2	4	1

In both versions, modulo is performed on only common columns in both dataframes. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Modulo on two dataframes and using fill_value parameter:

For example,

```
# mod() demo on two dataframes using method version and fill_value = 10
fdf1.mod(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	1	3
1	0	1
2	4	0

Here, modulo is performed on only common columns in both dataframes, excluding the missing values. Modulo is performed on other column elements using the value 10 (excluding the missing values) in resultant dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.6 6. DataFrame.mul(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be multiplied with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform multiplication operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform multiplication operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs multiplication operation between two operands. It is equivalent to **‘dataframe * other’**.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Multiply a scalar value using operator version:

For example,

```
print(fdf1 * 10)
```

Output

index	points	total
0	50	100
1	60	110
2	40	120

Multiply a scalar value using method version:

For example,

```
fdf1.mul(10).show()
```

Output

index	points	total
0	50	100
1	60	110
2	40	120

In both versions, all column elements (axis = 1 by default) are multiplied with a scalar value.

Creating two frovedis dataframes to perform multiplication:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Multiply two dataframes using operator version:

For example,

```
print(fdf1 * fdf2)
```

Output

index	points	total
0	10	70

1	18	99
2	32	132

Multiply two dataframes using method version:

For example,

```
# mul() demo on two dataframes using method version
fdf1.mul(other = fdf2).show()
```

Output

index	points	total
0	10	70
1	18	99
2	32	132

In both versions, only common columns in both dataframes are multiplied. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes in order to use fill_value parameter during multiplication:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Multiply two dataframes and using fill_value parameter:

For example,

```
# mul() demo on two dataframes using method version and fill_value = 10
fdf1.mul(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	10	70
1	18	110
2	32	100

Here, only common columns in both dataframes are multiplied, excluding the missing values. Other column elements are multiplied with the value 10 (excluding the missing values) in resultant dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.7 7. DataFrame.pow(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to perform exponential power operation with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform exponential power operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform exponential power operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element

needed for successful dataframe alignment, with this value before computation. (Default: None)
Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs exponential power operation between two operands. It is equivalent to ‘`dataframe ** other`’.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame()

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Exponential power operation on a scalar value using operator version:

For example,

```
print(fdf1 ** 2)
```

Output

index	points	total
0	10	20
1	12	22
2	8	24

Exponential power operation on a scalar value using method version:

For example,

```
fdf1.pow(2).show()
```

Output

index	points	total
0	10	20
1	12	22
2	8	24

In both versions, exponential power operation is performed on all column elements (axis = 1 by default) by a scalar value.

Creating two frovedis dataframes to perform exponential power operation:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Exponential power operation on two dataframes using operator version:

For example,

```
print(fdf1 ** fdf2)
```

Output

index	points	total
0	10	70
1	18	99
2	32	132

Exponential power operation on two dataframes using method version:

For example,

```
fdf1.pow(other = fdf2).show()
```

Output

index	points	total
0	10	70
1	18	99
2	32	132

In both versions, exponential power operation on only common columns in both dataframes. Exponential power operation on other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during exponential power operation:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
```

```

    }

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()

```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Exponential power operation on two dataframes and using fill_value parameter:

For example,

```

# pow() on two dataframes using method version and fill_value = 10
fdf1.pow(other = fdf2, fill_value = 10).show()

```

Output

index	points	total
0	25	1.00000e+07
1	216	2.59374e+10
2	65536	1.00000e+10

Here, exponential power operation is performed on only common columns in both dataframes, excluding the missing values. Exponential power operation on other column elements is performed with the value 10 (excluding the missing values) in resultant dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.8 8. DataFrame.sub(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be subtracted with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform subtraction operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform subtraction operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs subtraction operation between two operands. It is equivalent to '**dataframe - other**'.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Subtract a scalar value using operator version:

For example,

```
print(fdf1 - 10)
```

Output

index	points	total
0	-5	0
1	-4	1
2	-6	2

Subtract a scalar value using method version:

For example,

```
fdf1.sub(10).show()
```

Output

```
index  points  total
0      -5      0
1      -4      1
2      -6      2
```

In both versions, all column elements (axis = 1 by default) are subtracted by a scalar value.

Creating two frovedis dataframes to perform subtraction:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

```
index  points  total
0       5     10
1       6     11
2       4     12
```

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

```
index  points  total
```

0	2	7
1	3	9
2	8	11

Subtract two dataframes using operator version:

For example,

```
print(fdf1 - fdf2)
```

Output

index	points	total
0	3	3
1	3	2
2	-4	1

Subtract two dataframes using method version:

For example,

```
fdf1.sub(other = fdf2).show()
```

Output

index	points	total
0	3	3
1	3	2
2	-4	1

In both versions, only common columns in both dataframes are subtracted. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during subtraction:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Subtract two dataframes and using fill_value parameter:

For example,

```
# sub() demo on two dataframes using method version and fill_value = 10
fdf1.sub(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	3	3
1	3	1
2	-4	0

Here, only common columns in both dataframes are subtracted, excluding the missing values. Other column elements are subtracted with the value 10 (excluding the missing values) in resultant dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.9 9. DataFrame.truediv(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.

- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be divided with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform division operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform division operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs floating division operation between two operands. It is equivalent to '**dataframe / other**'.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [8, 5, 9],
    "total": [3, 2, 1]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	8	3
1	5	2
2	9	1

Floating Division on a scalar value using operator version:

For example,

```
print(fdf1 / 10)
```

Output

index	points	total
0	0.8	0.3
1	0.5	0.2
2	0.9	0.1

Floating Division on a scalar value using method version:

For example,

```
fdf1.truediv(10).show()
```

Output

index	points	total
0	0.8	0.3
1	0.5	0.2
2	0.9	0.1

In both versions, all column elements (axis = 1 by default) are divided by a scalar value.

Creating two frovedis dataframes to perform floating division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [8, 5, 9],
    "total": [3, 2, 1]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	8	3
1	5	2
2	9	1

For example,

```
# a dictionary
data2 = {
    "points": [4, 7, 2],
    "total": [9, 1, 9]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	4	9
1	7	1
2	2	9

Floating Division on two dataframes using operator version:

For example,

```
print(fdf1 / fdf2)
```

Output

index	points	total
0	2	0.333333
1	0.714285	2
2	4.5	0.111111

Floating Division on two dataframes using method version:

For example,

```
fdf1.truediv(other = fdf2).show()
```

Output

index	points	total
0	2	0.333333
1	0.714285	2
2	4.5	0.111111

In both versions, only common columns in both dataframes are divided. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during floating division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10

```
1      6      11
2      4      NULL
```

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

```
index  points  total
0      2      7
1      3      NULL
2      8      NULL
```

Floating Division on two dataframes and using fill_value parameter:

For example,

```
# truediv() demo on two dataframes using method version and fill_value = 10
fdf1.truediv(other = fdf2, fill_value = 10).show()
```

Output

```
index  points  total
0      2.5    1.42857
1      2      1.1
2      0.5    1
```

Here, only common columns in both dataframes are divided, excluding the missing values. Other column elements are divided with the value 10 (excluding the missing values) in resultant dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.10 10. DataFrame.radd(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.

- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be added with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse addition operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse addition operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse addition operation between two operands. It is equivalent to '**other + dataframe**'.

Currently, it does not perform reverse addition of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

```
index  points  total
0      5      10
1      6      11
2      4      12
```

Reverse addition on a scalar value using method version:

For example,

```
# radd() demo with scalar value using method version
fdf1.radd(10).show()
```

Output

index	points	total
0	15	20
1	16	21
2	14	22

Here, it adds the scalar to all columns in dataframe (axis = 1 by default).

Creating two frovedis dataframes to perform reverse addition:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For examples,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7

1	3	9
2	8	11

Reverse addition on two dataframes using method version:

For example,

```
# radd() demo on two dataframes using method version
fdf1.radd(other = fdf2).show()
```

Output

index	points	total
0	7	17
1	9	20
2	12	23

Here, only common columns in both dataframes are added. Column values in other dataframe are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse addition:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse addition on two dataframes and using fill_value parameter:

For example,

```
# radd() demo on two dataframes using method version and fill_value = 10
fdf1.radd(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	7	17
1	9	21
2	12	20

Here, only common columns in both dataframes are added excluding the missing values. The fill_value = 10 is added to both column values in the dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.11 11. DataFrame.rdiv(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be divided with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse division operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse division operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element

needed for successful dataframe alignment, with this value before computation. (Default: None)
Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse floating division operation between two operands. It is equivalent to ‘**other / dataframe**’.

It is an alias of `rtruediv()`.

Currently, it does not perform reverse division of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Reverse division on a scalar value using method version:

For example,

```
# rdiv() demo with scalar value using method version
fdf1.rdiv(10).show()
```

Output

index	points	total
0	2	1
1	1.66666	0.90909
2	2.5	0.833333

Here, it uses the scalar to perform division on all column elements in dataframe (axis = 1 by default).

Creating two frovedis dataframes to perform reverse division:

For example,

```
import pandas as pd
```

```
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Reverse division on two dataframes using method version:

For example,

```
# rdiv() demo on two dataframes using method version
fdf1.rdiv(other = fdf2).show()
```

Output

index	points	total
0	0.4	0.7

1	0.5	0.818181
2	2	0.916666

Here, only common columns in both dataframes are divided. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse division on two dataframes and using fill_value parameter:

For example,

```
# rdiv() demo on two dataframes using method version and fill_value = 10
fdf1.rdiv(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	0.4	0.7
1	0.5	0.90909
2	2	1

Here, only common columns in both dataframes are divided, excluding the missing values. The fill_value = 10 is used to divide over column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.12 12. DataFrame.rfloordiv(other, axis = 'columns', level = None, fill_value = None)**Parameters**

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be divided with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse division operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse division operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse floating division operation between two operands. It is equivalent to '**other // dataframe**'.

Currently, it does not perform reverse division of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Reverse floor division on a scalar value using method version:

For example,

```
# rfloordiv() demo with scalar value using method version
fdf1.rfloordiv(10).show()
```

Output

index	points	total
0	2	1
1	1	0
2	2	0

Here, it uses the scalar to perform division on all column elements (axis = 1 by default). Also, resultant dataframe column elements will contain floor integer value.

Creating two frovedis dataframes to perform reverse floor division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)
```

```
# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Reverse floor division on two dataframes using method version:

For example,

```
# rfloordiv() demo on two dataframes using method version
fdf1.rfloordiv(other = fdf2).show()
```

Output

index	points	total
0	0	0
1	0	0
2	2	0

Here, only common columns in both dataframes are divided. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse floor division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
```

```
# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}
```

```
# create pandas dataframe
pdf1 = pd.DataFrame(data1)
```

```
# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)
```

```
# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse floor division on two dataframes and using fill_value parameter:

For example,

```
# rfloordiv() demo on two dataframes using method version and fill_value = 10
fdf1.rfloordiv(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	0	0
1	0	0
2	2	1

Here, only common columns in both dataframes are divided, excluding the missing values. The `fill_value = 10` is used to divide over column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.13 13. DataFrame.rmod(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to perform modulo operation with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse modulo operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse modulo operation on the columns. Currently, `axis = 1` is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse modulo operation between two operands. It is equivalent to `'other % dataframe'`.

Currently, it does not perform reverse modulo of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [50, 40, 20]
}

# create pandas dataframe
```



```
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Reverse modulo on a scalar value using method version:

For example,

```
# rmod() demo with scalar value using method version
fdf1.rmod(10).show()
```

Output

index	points	total
0	0	0
1	4	10
2	2	10

Here, it uses the scalar to perform modulo operation on all column elements (axis = 1 by default).

Creating two frovedis dataframes to perform reverse modulo:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Reverse modulo on two dataframes using method version:

For example,

```
# rmod() demo on two dataframes using method version
fdf1.rmod(other = fdf2).show()
```

Output

index	points	total
0	2	7
1	3	9
2	0	11

Here, modulo is performed on only common columns in both dataframes. Other are replaced with NaN values in resultant dataframe (`fill_value = None` by default).

Creating two frovedis dataframes and use `fill_value` parameter during reverse modulo:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
```

```
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse modulo on two dataframes and using fill_value parameter:

For example,

```
# rmod() demo on two dataframes using method version and fill_value = 10
fdf1.rmod(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	2	7
1	3	10
2	0	0

Here, modulo is performed on only common columns in both dataframes, excluding the missing values. The fill_value = 10 is used to perform modulo over column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.14 14. DataFrame.rmul(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be multiplied with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse multiplication operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse multiplication operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse multiplication operation between two operands. It is equivalent to **'other * dataframe'**.

Currently, it does not perform reverse multiplication of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11

```
2      4      12
```

Reverse multiplication on a scalar value using method version:

For example,

```
# rmul() demo with scalar value using method version
fdf1.rmul(10).show()
```

Output

```
index  points  total
0      50     100
1      60     110
2      40     120
```

Here, it uses the scalar to perform multiplication on all column elements in dataframe (axis = 1 by default).

Creating two frovedis dataframes to perform reverse multiplication:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
```

```
# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}
```

```
# create pandas dataframe
pdf1 = pd.DataFrame(data1)
```

```
# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)
```

```
# display the frovedis dataframe
fdf1.show()
```

Output

```
index  points  total
0      5      10
1      6      11
2      4      12
```

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Reverse multiplication on two dataframes using method version:

For example,

```
# rmul() demo on two dataframes using method version
fdf1.rmul(other = fdf2).show()
```

Output

index	points	total
0	10	70
1	18	99
2	32	132

Here, only common columns in both dataframes are multiplied. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse multiplication:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
```

```

        "points": [2, 3, 8],
        "total": [7, np.nan, np.nan]
    }

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()

```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse multiplication on two dataframes and using fill_value parameter:

For example,

```

# rmul() demo on two dataframes using method version and fill_value = 10
fdf1.rmul(other = fdf2, fill_value = 10).show()

```

Output

index	points	total
0	10	70
1	18	110
2	32	100

Here, only common columns in both dataframes are multiplied, excluding the missing values. The fill_value = 10 is multiplied with column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.15 15. DataFrame.rpow(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to perform exponential power operation with the current dataframe. **axis**: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse exponential power operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns'**: perform reverse exponential power operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse exponential power operation between two operands. It is equivalent to '**other ** dataframe**'.

Currently, it does not perform reverse exponential power operation of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Reverse exponential power operation on a scalar value using method version:

For example,

```
# rpow() demo with scalar value using method version
fdf1.rpow(2).show()
```

Output

index	points	total
0	32	1024
1	64	2048

2 16 4096

Here, it uses the scalar to perform exponential power operation on all column elements in dataframe (axis = 1 by default).

Creating two frovedis dataframes to perform reverse exponential power operation:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Reverse exponential power operation on two dataframes using method version:

For example,

```
# rpow() demo on two dataframes using method version
fdf1.rpow(other = fdf2).show()
```

Output

index	points	total
0	10	70
1	18	99
2	32	132

Here, exponential power operation is performed on only common columns in both dataframes. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse exponential power operation:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse exponential power operation on two dataframes and using fill_value parameter:

For example,

```
# rpow() demo on two dataframes using method version and fill_value = 10
fdf1.rpow(other = fdf2, fill_value = 10).show()
```

Output

index	points	total
0	32	2.82475e+08
1	729	9.99999e+10
2	4096	1.00000e+10

Here, exponential power operation is performed on only common columns in both dataframes, excluding the missing values. The fill_value = 10 is used to perform exponential power operation on column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a frovedis DataFrame which contains the result of arithmetic operation.

56.1.3.16 16. DataFrame.rsub(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be subtracted with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse subtraction operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse subtraction operation on the columns. Currently, axis = 1 is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse subtraction operation between two operands. It is equivalent to **‘other - dataframe’**.

Currently, it does not perform reverse subtraction of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

Reverse subtraction on a scalar value using method version:

For example,

```
# rsub() with scalar value using method version
fdf1.rsub(10).show()
```

Output

index	points	total
0	5	0
1	4	-1
2	6	-2

Here, it subtracts the scalar to all columns in dataframe (axis = 1 by default).

Creating two frovedis dataframes to perform reverse subtraction:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
```

```
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, 12]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	12

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, 9, 11]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	9
2	8	11

Reverse subtraction on two dataframes using method version:

For example,

```
# rsub() on two dataframes using method version
fdf1.rsub(other = fdf2).show()
```

Output

index	points	total
0	-3	-3
1	-3	-2
2	4	-1

Here, only common columns in both dataframes are subtracted. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse subtraction:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```
# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse subtraction on two dataframes and using fill_value parameter:

For example,

```
# rsub() on two dataframes using method version and fill_value = 10
fdf1.rsub(other = fdf2, fill_value = 10).show()
```

Output

```
index  points  total
0      -3     -3
1      -3     -1
2       4       0
```

Here, only common columns in both dataframes are subtracted, excluding the missing values. The `fill_value = 10` is subtracted on column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a `frovedis DataFrame` which contains the result of arithmetic operation.

56.1.3.17 17. DataFrame.rtruediv(other, axis = 'columns', level = None, fill_value = None)

Parameters

other: It can accept single or multiple element data structure like the following:

- Number
- List having 1 dimension. Currently, this method supports operation on only list of numeric values.
- A numpy ndarray having 1 dimension. Currently, this method supports operation on only an array of numeric values.
- pandas DataFrame. It must not be an empty dataframe.
- pandas Series
- frovedis DataFrame. It must not be an empty dataframe.

Any of these is considered as the value to be divided with the current dataframe.

axis: It accepts an integer or string object as parameter. It is used to decide whether to perform reverse division operation along the indices or by column labels. (Default: 'columns')

- **1 or 'columns':** perform reverse division operation on the columns. Currently, `axis = 1` is supported in this method.

level: This is an unused parameter. (Default: None)

fill_value: It accepts scalar values or None. It fills existing missing (NaN) values, and any new element needed for successful dataframe alignment, with this value before computation. (Default: None)

Irrespective of the specified value, if data in both corresponding dataframe locations is missing, then the result will be missing (contains NaNs).

Purpose

It performs reverse floating division operation between two operands. It is equivalent to `'other / dataframe'`.

Currently, it does not perform reverse division of scalar using operator version. Only method version is supported.

Creating frovedis DataFrame from pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
```

```
# a dictionary
data1 = {
    "points": [8, 5, 9],
    "total": [3, 2, 1]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()
```

Output

index	points	total
0	8	3
1	5	2
2	9	1

Reverse floating division on a scalar value using method version:

For example,

```
# rtruediv() demo with scalar value using method version
fdf1.rtruediv(10).show()
```

Output

index	points	total
0	1.25	3.33333
1	2	5
2	1.11111	10

Here, it uses the scalar to perform division on all column elements in dataframe (axis = 1 by default).

Creating two frovedis dataframes to perform reverse floating division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [8, 5, 9],
    "total": [3, 2, 1]
}

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
```



```
fdf1.show()
```

Output

index	points	total
0	8	3
1	5	2
2	9	1

For example,

```
# a dictionary
data2 = {
    "points": [4, 7, 2],
    "total": [9, 1, 9]
}
```

```
# create pandas dataframe
pdf2 = pd.DataFrame(data2)
```

```
# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)
```

```
# display the frovedis dataframe
fdf2.show()
```

Output

index	points	total
0	4	9
1	7	1
2	2	9

Reverse floating division on two dataframes using method version:

For example,

```
# rtruediv() demo on two dataframes using method version
fdf1.rtruediv(other = fdf2).show()
```

Output

index	points	total
0	0.5	3
1	1.39999	0.5
2	0.222222	9

Here, only common columns in both dataframes are divided. Other are replaced with NaN values in resultant dataframe (fill_value = None by default).

Creating two frovedis dataframes and use fill_value parameter during reverse floating division:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
data1 = {
    "points": [5, 6, 4],
    "total": [10, 11, np.nan]
```

```

    }

# create pandas dataframe
pdf1 = pd.DataFrame(data1)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	points	total
0	5	10
1	6	11
2	4	NULL

For example,

```

# a dictionary
data2 = {
    "points": [2, 3, 8],
    "total": [7, np.nan, np.nan]
}

# create pandas dataframe
pdf2 = pd.DataFrame(data2)

# create frovedis dataframe
fdf2 = fdf.DataFrame(pdf2)

# display the frovedis dataframe
fdf2.show()

```

Output

index	points	total
0	2	7
1	3	NULL
2	8	NULL

Reverse floating division on two dataframes and using fill_value parameter:

For example,

```

# rtruediv() demo on two dataframes using method version and fill_value = 10
fdf1.rtruediv(other = fdf2, fill_value = 10).show()

```

Output

index	points	total
0	0.4	0.7
1	0.5	0.90909
2	2	1

Here, only common columns in both dataframes are divided, excluding the missing values. The fill_value = 10 is divided with column values in other dataframe (excluding the missing values) and stored in new dataframe.

Return Value

It returns a `frovedis DataFrame` which contains the result of arithmetic operation.

56.2 SEE ALSO

- **DataFrame - Introduction**
- **DataFrame - Generic Functions**
- **DataFrame - Conversion Functions**
- **DataFrame - Sorting Functions**
- **DataFrame - Aggregate Functions**

Chapter 57

FrovedisGroupedDataframe

57.1 NAME

FrovedisGroupedDataframe - A python class for handling grouped dataframes. These are returned by groupby calls.

57.2 SYNOPSIS

```
frovedis.dataframe.grouped_df.FrovedisGroupedDataframe(df = None)
```

57.3 DESCRIPTION

FrovedisGroupedDataframe instance contains information about the grouped dataframe. After columns are grouped, various aggregations can be performed like groupwise average, groupwise variance, etc.

In FrovedisGroupedDataframe, currently aggregation operations is performed along the rows only.

This module provides a client-server implementation, where the client application is a normal python program. The FrovedisGroupedDataframe interface is almost same as pandas DataFrameGroupBy interface, but it doesn't have any dependency on pandas. It can be used simply even if the system doesn't have pandas installed. The FrovedisGroupedDataframe instance is created when groupby interface is called for frovedis Dataframe instance. Thus, in this implementation, a python client can interact with a frovedis server sending the required python dataframe for performing query at frovedis side. Python dataframe is converted into frovedis compatible dataframe internally and the python client side call is linked with the respective frovedis side call to get the job done at frovedis server.

57.3.1 Detailed description

57.3.1.1 1. FrovedisGroupedDataframe(df = None)

Parameters

df: It accepts only frovedis dataframe as parameter. (Default: None)

When it is None (not specified explicitly), an empty FrovedisGroupedDataframe instance is created.

Purpose

It is used for a specific purpose. It's instance is created in order to hold result of groupby method calls. This instance can then further be used with aggregate functions such as mean(), sem(), etc.

Creating FrovedisGroupedDataframe instance using groupby operation:

For example,

```
# create frovedis dataframe
import pandas as pd
import frovedis.dataframe as fdf
peopleDF = {
    'Ename' : ['Michael', 'Andy', 'Tanaka', 'Raul', 'Yuta'],
    'Age' : [29, 30, 27, 19, 31],
    'Country' : ['USA', 'England', 'Japan', 'France', 'Japan'],
    'isMale': [False, False, False, False, True]
}
pdf1 = pd.DataFrame(peopleDF)
fdf1 = fdf.DataFrame(pdf1)

# create FrovedisGroupedDataframe object
g_df = fdf1.groupby('Country')
```

The groupby call returns a FrovedisGroupedDataframe instance.

Multiple columns can be used in groupby operation to create this instance:

For example,

```
# using multiple columns to create FrovedisGroupedDataframe object
g_df = fdf1.groupby(['Age', 'Country'])
```

In order to select one of the grouped column:

For example,

```
print(g_df['Country'])
```

Output

```
Country
England
France
Japan
USA
```

In order to select multiple columns from the grouped dataframe:

For example,

```
print(g_df[['Age', 'Country']])
```

Output

index	Age	Country
0	19	France
1	27	Japan
2	29	USA
3	30	England
4	31	Japan

Return Value

It simply returns “self” reference.

57.3.1.2 2. `release()`

Purpose

This method acts like a destructor. It is used to release dataframe pointer from server heap and it resets all its attributes to None.

For example,

```
g_df.release()
```

Return Value

It returns nothing.

57.3.2 Public Member Functions

FrovedisGroupedDataFrame provides a lot of utilities to perform various operations.

57.3.2.1 List of Aggregate Functions

1. **agg()** - it aggregates using one or more operations over the specified axis. It is an alias for `aggregate()`.
2. **aggregate()** - it aggregates using one or more operations over the specified axis. The alias `agg()` can be used instead.
3. **count()** - it computes count of group, excluding missing values.
4. **max()** - it computes maximum of group values.
5. **mean()** - it computes mean of groups, excluding missing values.
6. **min()** - it computes minimum of group values.
7. **sem()** - it computes standard error of the mean of groups, excluding missing values.
8. **size()** - it computes group sizes.
9. **sum()** - it computes sum of group values.
10. **var()** - it computes variance of groups, excluding missing values.

57.4 SEE ALSO

- Using aggregate functions on **FrovedisGroupedDataFrame**
- Introduction to **frovedis DataFrame**

Chapter 58

FrovedisGroupedDataFrame Aggregate Functions

58.1 NAME

FrovedisGroupedDataFrame Aggregate Functions - aggregate operations performed on grouped dataframe are being illustrated here.

58.1.1 DESCRIPTION

Once FrovedisGroupedDataframe instance is created, several aggregation operations can be performed on it such as `max()`, `min()`, `sem()`, `var()`, etc.

The aggregation operation will basically compute summary for each group. Some examples:

- Compute group sums, means, var or sem.
- Compute group sizes / counts.
- Compute max or min in the group.

Also, aggregation functions can be chained along with `groupby()` calls in `frovedis`.

58.1.2 Public Member Functions

```
1. agg(func, *args, **kwargs)
2. aggregate(func, *args, **kwargs)
3. count(numeric_only = True)
4. groupby()
5. max(numeric_only = True, min_count = -1)
6. mean(numeric_only = True)
7. min(numeric_only = True, min_count = -1)
8. sem(ddof = 1.0)
9. size(numeric_only = True)
10. sum(numeric_only = True, min_count = 0)
11. var(ddof = 1.0)
```

58.1.3 Detailed Description

58.1.3.1 1. FrovedisGroupedDataFrame.agg(func, *args, **kwargs)

Parameters

func: Names of functions to use for aggregating the data. The input to be used with the function must be a frovedis DataFrame instance having atleast one numeric column.

Accepted combinations for this parameter are: - A string function name such as 'max', 'min', etc.

- list of functions and/or function names, For example, ['max', 'mean'].

- dictionary with keys as column labels and values as function name or list of such functions.

For Example, {'Age': ['max', 'min', 'mean'], 'Ename': ['count']}

***args**: This is an unused parameter.

****kwargs**: This is an unused parameter.

Purpose

It computes an aggregate operation based on the condition specified in 'func'. It is an alias for aggregate().

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Aggregate the function on the grouped dataframe where func is a function string name:

For example,

```
# agg() demo with func as a function string name
fdf1.groupby('Qualification').agg('min').show()
```

Output

Qualification	min_Age	min_Score
B.Tech	22	23
Phd	24	34

It displays a frovedis dataframe containing numeric column(s) with newly computed minimum of each groups.

Aggregate the function on the grouped dataframe where func is a dictionary:

For example,

```
# agg() demo with func as a dictionary
fdf1.groupby('Qualification').agg({'Age': ['max','min','mean'], 'Score': ['sum']}).show()
```

Output

Qualification	max_Age	min_Age	mean_Age	sum_Score
B.Tech	36	22	29.25	108
Phd	33	24	29	13

Aggregate the function on the grouped dataframe where func is a list of functions:

For example,

```
# agg() demo where func is a list of functions
fdf1.groupby('Qualification').agg(['min', 'max']).show()
```

Output

Qualification	min_Age	max_Age	min_Score	max_Score
B.Tech	22	36	23	50
Phd	24	33	34	52

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.2 2. FrovedisGroupedDataFrame.aggregate(func, *args, **kwargs)

Parameters

func: Names of functions to use for aggregating the data. The input to be used with the function must be a frovedis DataFrame instance having atleast one numeric column.

Accepted combinations for this parameter are: - A string function name such as 'max', 'min', etc.

- list of functions and/or function names, For example, ['max', 'mean'].

- dictionary with keys as column labels and values as function name or list of such functions.

For Example, {'Age': ['max','min','mean'], 'Ename': ['count']}

***args:** This is an unused parameter.

****kwargs:** This is an unused parameter.

Purpose

It computes an aggregate operation based on the condition specified in 'func'.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```

import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()

```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Aggregate the function on the grouped dataframe where func is a function string name:

For example,

```

# aggregate() demo using FrovedisGroupedDataframe instance and 'func' as a function string name
fdf1.groupby('Qualification').agg('max').show()

```

Output

Qualification	max_Age	max_Score
B.Tech	36	50
Phd	33	52

It displays a frovedis dataframe containing numeric column(s) with newly computed minimum of each groups.

Aggregate the function on the grouped dataframe where func is a dictionary:

For example,

```

# aggregate() demo using FrovedisGroupedDataframe instance and 'func' as a dictionary
fdf1.groupby('Qualification').agg({'Age': ['count'], 'Score': ['max', 'min']}).show()

```

Output

Qualification	count_Age	max_Score	min_Score
---------------	-----------	-----------	-----------

B.Tech	4	50	23
Phd	4	52	34

Aggregate the function on the grouped dataframe where func is a list of functions:

For example,

```
# aggregate() demo using FrovedisGroupedDataframe instance and 'func' is a list of functions
fdf1.groupby('Qualification').agg(['mean', 'sum']).show()
```

Output

Qualification	mean_Age	sum_Age	mean_Score	sum_Score
B.Tech	29.25	117	36	108
Phd	29	116	43.6666	131

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.3 3. FrovedisGroupedDataFrame.count(numeric_only = True)

Parameters

numeric_only: It accepts a boolean parameter that specifies whether or not to use only numeric column data as input. (Default: True)

Purpose

It computes count of groups, excluding the missing values.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34

2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Count groups in each column of the grouped dataframe:

For example,

```
# count() demo using FrovedisGroupedDataframe instance
fdf1.groupby('Qualification').count().show()
```

Output

Qualification	count_Age	count_Score	count_Name	count_City
B.Tech	4	3	4	4
Phd	4	3	4	4

It displays a frovedis dataframe containing the newly computed count of each groups.

Also, it excludes the missing value(s) in ‘Score’ column while computing count of groups ‘B.Tech’ and ‘Phd’.

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.4 4. DataFrame.groupby(by = None, axis = 0, level = None, as_index = True, sort = True, group_keys = True, squeeze = False, observed = False, dropna = True)

Parameters

by: It accepts a string object or an iterable to determine the groups on which groupby operation will be applied. Currently, groupby operation will be applied along the index levels. It must be provided, otherwise it will raise an exception. (Default: None)

axis: It accepts an integer as parameter. It is used to decide whether to perform groupby operation along the indices or by column labels. (Default: 0)

Currently, axis = 0 is supported by this method.

level: This is an unused parameter. (Default: None)

as_index: This is an unused parameter. (Default: True)

sort: This is an unused parameter. (Default: True)

group_keys: This is an unused parameter. (Default: True)

squeeze: This is an unused parameter. (Default: False)

observed: This is an unused parameter. (Default: False)

dropna: It accepts a boolean parameter. It is used to remove missing values (NaNs) from the frovedis DataFrame during groupby operation. Currently, it removes missing values along the index levels. (Default: True)

Purpose

This method can be used to group large amounts of data and compute operations on these groups.

The parameters: “level”, “as_index”, “sort”, “group_keys”, “squeeze” and “observed” is simply kept in to make the interface uniform to the pandas DataFrame.groupby(). This is not used anywhere within the frovedis implementation.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```

import pandas as pd
import numpy as np
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', np.nan, 'Phd', 'B.Tech', 'Phd', np.nan],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# create pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# create frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# display the frovedis dataframe
fdf1.show()

```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	NULL	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	NULL	NULL

Performing groupby operation on the frovedis dataframe:

For example,

```
fdf1.groupby('Qualification')
```

This will perform groupby operation on the dataframe over ‘**Qualification**’ column data.

Using groupby() to perform aggregation on resultant grouped dataframe. Also dropna = True:

For example,

```
fdf1.groupby('Qualification', dropna = True).agg({'Score': 'count'})
```

Output

Qualification	count_Score
B.Tech	3
Phd	2

Here, it excludes **NULL** group since missing values were dropped during groupby().

Using groupby() to perform aggregation on resultant grouped dataframe. Also dropna = False:

For example,

```
fdf1.groupby('Qualification', dropna = False).agg({'Score': 'count'})
```

Output

```
Qualification    count_Score
B.Tech          3
Phd              2
NULL            1
```

Here, it includes **NULL** as new group since missing values were not dropped during `groupby()`.

Return Value

It returns a `FrovedisGroupedDataFrame` instance. This instance is then further used to perform aggregate operations.

58.1.3.5 5. `DataFrame.max(numeric_only = True, min_count = -1)`

Parameters

numeric_only: It accepts a boolean parameter that specifies whether or not to use only numeric columns for aggregation. (Default: True)

min_count: This is an unused parameter. (Default: -1)

Purpose

It computes the maximum of group values.

The parameter: “`min_count`” is simply kept in to to make the interface uniform to the pandas `GroupBy.max()`. This is not used anywhere within the `frovedis` implementation.

Currently, this method only displays result for dataframe having atleast one numeric column.

Constructing `frovedis DataFrame` from a `pandas DataFrame`:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

To calculate maximum value in each group of grouped dataframe:

For example,

```
# max() demo using FrovedisGroupedDataframe instance
fdf1.groupby('Qualification').max().show()
```

Output

Qualification	max_Age	max_Score
B.Tech	36	50
Phd	33	52

It displays a frovedis dataframe containing numeric column(s) with newly computed maximum of each groups.

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.6 6. FrovedisGroupedDataFrame.mean(numeric_only = True)

Parameters

numeric_only: It accepts a boolean parameter that specifies whether or not to use only numeric column data as input. (Default: True)

Purpose

It computes mean of groups, excluding the missing values.

Currently, this method only displays result for dataframe having atleast one numeric column.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannauj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)
```

```
# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannauj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Compute mean of each group:

For example,

```
# mean() demo using FrovedisGroupedDataframe instance
fdf1.groupby('Qualification').mean().show()
```

Output

Qualification	mean_Age	mean_Score
B.Tech	29.25	36
Phd	29	43.6666

It displays a frovedis dataframe containing numeric column(s) with newly computed mean of each groups.

Also, it excludes the missing value(s) in ‘Score’ column while computing mean of groups ‘B.Tech’ and ‘Phd’.

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.7 7. FrovedisGroupedDataFrame.min(numeric_only = True, min_count = -1)

Parameters

numeric_only: It accepts a boolean parameter that specifies whether or not to use only numeric columns for aggregation. (Default: True)

min_count: This is an unused parameter. (Default: -1)

Purpose

It computes the minimum of group values.

The parameter: “min_count” is simply kept in to to make the interface uniform to the pandas GroupBy.min(). This is not used anywhere within the frovedis implementation.

Currently, this method only displays result for dataframe having atleast one numeric column.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf
```

```
# a dictionary
peopleDF = {
    'Name':['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age':[27, 24, 22, 32, 33, 36, 27, 32],
    'City':['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification':['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

To calculate minimum value in each group of grouped dataframe:

For example,

```
# min() demo using FrovedisGroupedDataframe instance
fdf1.groupby('Qualification').min().show()
```

Output

Qualification	min_Age	min_Score
B.Tech	22	23
Phd	24	34

It displays a frovedis dataframe containing numeric column(s) with newly computed minimum of each groups.

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.8 8. FrovedisGroupedDataFrame.sem(ddof = 1.0)

Parameters

ddof: It accepts an integer parameter that specifies the delta degrees of freedom. (Default: 1.0)

Purpose

It computes standard error of the mean of groups, excluding missing values.

Currently, this method only displays result for dataframe having atleast one numeric column.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
             'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

Compute standard error of the mean of each group:

For example,

```
# sem() demo using FrovedisGroupedDataframe instance and default ddof value
fdf1.groupby('Qualification').sem().show()
```

Output

Qualification	sem_Age	sem_Score
B.Tech	3.03795	7.81024
Phd	2.12132	5.23874

It displays a frovedis dataframe with numeric column(s) containing the newly computed standard error of mean for each groups.

Also, it excludes the missing value(s) in ‘Score’ column while computing standard error of mean for the groups ‘B.Tech’ and ‘Phd’.

Using ddof parameter to compute standard error of the mean of each group:

For example,

```
# sem() demo using ddof = 2
fdf1.groupby('Qualification').sem(ddof = 2).show()
```

Output

Qualification	sem_Age	sem_Score
B.Tech	3.72071	11.0453
Phd	2.59807	7.4087

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.9 9. FrovedisGroupedDataFrame.size(numeric_only = True)

Parameters

numeric_only: It accepts a boolean parameter that specifies whether or not to use only numeric column data as input. (Default: True)

Purpose

It computes group sizes, including the missing values.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

To compute size of groups for the grouped dataframe:

For example,

```
# size() demo using FrovedisGroupedDataframe instance
fdf1.groupby('Qualification').size().show()
```

Output

```
Qualification  size_Qualification
B.Tech         4
Phd            4
```

It displays a frovedis dataframe containing a column with the newly computed size of each group.

Also, it does not exclude the missings values while computing group size.

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.10 10. FrovedisGroupedDataFrame.sum(numeric_only = True, min_count = 0)

Parameters

numeric_only: It accepts a boolean parameter that specifies whether or not to use only numeric column data as input. (Default: True)

min_count: This is an unused parameter. (Default: 0)

Purpose

It computes the sum of group values.

The parameter: “min_count” is simply kept in to to make the interface uniform to the pandas GroupBy.sum(). This is not used anywhere within the frovedis implementation.

Currently, this method only displays result for dataframe having atleast one numeric column.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)

# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

To compute sum of group values for the grouped dataframe:

For example,

```
# sum() demo using FrovedisGroupedDataframe instance
fdf1.groupby('Qualification').sum().show()
```

Output

Qualification	sum_Age	sum_Score
B.Tech	117	108
Phd	116	131

It displays a frovedis dataframe containing numeric column(s) with newly computed sum of each groups.

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.1.3.11 11. FrovedisGroupedDataFrame.var(ddof = 1.0)

Parameters

ddof: It accepts an integer parameter that specifies the delta degrees of freedom. (Default: 1.0)

Purpose

It computes the variance of groups, excluding missing values.

Currently, this method only displays result for dataframe having atleast one numeric column.

Constructing frovedis DataFrame from a pandas DataFrame:

For example,

```
import pandas as pd
import frovedis.dataframe as fdf

# a dictionary
peopleDF = {
    'Name': ['Jai', 'Anuj', 'Jai', 'Princi', 'Gaurav', 'Anuj', 'Princi', 'Abhi'],
    'Age': [27, 24, 22, 32, 33, 36, 27, 32],
    'City': ['Nagpur', 'Kanpur', 'Allahabad', 'Kannuaj', 'Allahabad',
            'Kanpur', 'Kanpur', 'Kanpur'],
    'Qualification': ['B.Tech', 'Phd', 'B.Tech', 'Phd', 'Phd', 'B.Tech', 'Phd', 'B.Tech'],
    'Score': [23, 34, 35, 45, np.nan, 50, 52, np.nan]
}

# convert to pandas dataframe
pdf1 = pd.DataFrame(peopleDF)
```

```
# convert to frovedis dataframe
fdf1 = fdf.DataFrame(pdf1)

# displaying created frovedis dataframe
fdf1.show()
```

Output

index	Name	Age	City	Qualification	Score
0	Jai	27	Nagpur	B.Tech	23
1	Anuj	24	Kanpur	Phd	34
2	Jai	22	Allahabad	B.Tech	35
3	Princi	32	Kannuaj	Phd	45
4	Gaurav	33	Allahabad	Phd	NULL
5	Anuj	36	Kanpur	B.Tech	50
6	Princi	27	Kanpur	Phd	52
7	Abhi	32	Kanpur	B.Tech	NULL

To compute variance of groups in the grouped dataframe:

For example,

```
# var() demo using FrovedisGroupedDataframe instance and default ddof value
fdf1.groupby('Qualification').var().show()
```

Output

Qualification	var_Age	var_Score
B.Tech	36.9166	183
Phd	18	82.3333

It displays a frovedis dataframe with numeric column(s) containing the newly computed variance for each groups.

Also, it excludes the missing value in ‘Score’ column while computing variance of groups ‘B.Tech’ and ‘Phd’.

Using ddof parameter to compute variance of each group:

For example,

```
# var() demo using ddof = 2
fdf1.groupby('Qualification').var(ddof = 2).show()
```

Output

Qualification	var_Age	var_Score
B.Tech	55.375	366
Phd	27	164.666

Return Value

It returns a new frovedis DataFrame instance with the result of the specified aggregate functions.

58.2 SEE ALSO

- Introduction to FrovedisGroupedDataFrame
- Introduction to frovedis DataFrame