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# **expandas Documentation**

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## Data Handling

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### 1.1 Data Preparation

This section describes how to prepare basic data format named `ModelFrame`. `ModelFrame` defines a metadata to specify target (response variable) and data (explanatory variable / features). Using these metadata, `ModelFrame` can call other statistics/ML functions in more simple way.

You can create `ModelFrame` as the same manner as `pandas.DataFrame`. The below example shows how to create basic `ModelFrame`, which DOESN'T have target values.

```
>>> import expandas as expd

>>> df = expd.ModelFrame({'A': [1, 2, 3], 'B': [2, 3, 4],
...                       'C': [3, 4, 5]}, index=['A', 'B', 'C'])
>>> df
   A  B  C
A  1  2  3
B  2  3  4
C  3  4  5

>>> type(df)
<class 'expandas.core.frame.ModelFrame'>
```

You can check whether the created `ModelFrame` has target values using `ModelFrame.has_target()` function.

```
>>> df.has_target()
False
```

Target values can be specifyied via `target` keyword. You can simply pass a column name to be handled as target. Target column name can be confirmed via `target_name` property.

```
>>> df2 = expd.ModelFrame({'A': [1, 2, 3], 'B': [2, 3, 4],
...                       'C': [3, 4, 5]}, target='A')
>>> df2
   A  B  C
0  1  2  3
1  2  3  4
2  3  4  5

>>> df2.has_target()
True

>>> df2.target_name
'A'
```

Also, you can pass any list-likes to be handled as a target. In this case, target column will be named as ".target".

```
>>> df3 = expd.ModelFrame({'A': [1, 2, 3], 'B': [2, 3, 4],
...                       'C': [3, 4, 5]}, target=[4, 5, 6])
>>> df3
   .target  A  B  C
0         4  1  2  3
1         5  2  3  4
2         6  3  4  5

>>> df3.has_target()
True

>>> df3.target_name
'.target'
```

Also, you can pass `pandas.DataFrame` and `pandas.Series` as data and target.

```
>>> import pandas as pd
df4 = expd.ModelFrame({'A': [1, 2, 3], 'B': [2, 3, 4],
...                   'C': [3, 4, 5]}, target=pd.Series([4, 5, 6]))
>>> df4
   .target  A  B  C
0         4  1  2  3
1         5  2  3  4
2         6  3  4  5

>>> df4.has_target()
True

>>> df4.target_name
'.target'
```

---

**Note:** Target values are mandatory to perform operations which require response variable, such as regression and supervised learning.

---

## 1.2 Data Manipulation

You can access to each property as the same as `pandas.DataFrame`. Sliced results will be `ModelSeries` (simple wrapper for `pandas.Series` to support some data manipulation) or `ModelFrame`

```
>>> df
   A  B  C
A  1  2  3
B  2  3  4
C  3  4  5

>>> sliced = df['A']
>>> sliced
A    1
B    2
C    3
Name: A, dtype: int64

>>> type(sliced)
<class 'expandas.core.series.ModelSeries'>
```



```
>>> subset = df[['A', 'B']]
>>> subset
   A  B
A  1  2
B  2  3
C  3  4

>>> type(subset)
<class 'expandas.core.frame.DataFrame'>
```

ModelFrame has a special properties `data` to access data (features) and `target` to access target.

```
>>> df2
   A  B  C
0  1  2  3
1  2  3  4
2  3  4  5

>>> df2.target_name
'A'

>>> df2.data
   B  C
0  2  3
1  3  4
2  4  5

>>> df2.target
0    1
1    2
2    3
Name: A, dtype: int64
```

You can update data and target via properties, in addition to standard `pandas.DataFrame` ways.

```
>>> df2.target = [9, 9, 9]
>>> df2
   A  B  C
0  9  2  3
1  9  3  4
2  9  4  5

>>> df2.data = pd.DataFrame({'X': [1, 2, 3], 'Y': [4, 5, 6]})
>>> df2
   A  X  Y
0  9  1  4
1  9  2  5
2  9  3  6

>>> df2['X'] = [0, 0, 0]
>>> df2
   A  X  Y
0  9  0  4
1  9  0  5
2  9  0  6
```

You can change target column specifying `target_name` property. Specifying a column which doesn't exist in ModelFrame results in target column to be data column.

```
>>> df2.target_name
'A'

>>> df2.target_name = 'X'
>>> df2.target_name
'X'

>>> df2.target_name = 'XXXX'
>>> df2.has_target()
False

>>> df2.data
   A  X  Y
0  9  0  4
1  9  0  5
2  9  0  6
```

---

## Use scikit-learn

---

This section describes how to use `scikit-learn` functionalities via `expandas`.

### 2.1 Basics

You can create `ModelFrame` instance from `scikit-learn` datasets directly.

```
>>> import expandas as expd
>>> import sklearn.datasets as datasets

>>> df = expd.ModelFrame(datasets.load_iris())
>>> df.head()
```

	.target	sepal length (cm)	sepal width (cm)	petal length (cm)	\
0	0	5.1	3.5	1.4	
1	0	4.9	3.0	1.4	
2	0	4.7	3.2	1.3	
3	0	4.6	3.1	1.5	
4	0	5.0	3.6	1.4	

  

	petal width (cm)
0	0.2
1	0.2
2	0.2
3	0.2
4	0.2

```
# make columns be readable
>>> df.columns = ['.target', 'sepal length', 'sepal width', 'petal length', 'petal width']

ModelFrame has accessor methods which makes easier access to scikit-learn namespace.

>>> df.cluster.KMeans
<class 'sklearn.cluster.k_means_.KMeans'>
```

Following table shows `scikit-learn` module and corresponding `ModelFrame` module. Some accessors has its abbreviated versions.

---

**Note:** Currently, `ModelFrame` can handle target which consists from a single column. Modules which uses multiple target columns cannot be handled automatically, and marked with (*WIP*).

---

scikit-learn	ModelFrame accessor
sklearn.cluster	ModelFrame.cluster
sklearn.covariance	ModelFrame.covariance
sklearn.cross_validation	ModelFrame.cross_validation, crv
sklearn.decomposition	ModelFrame.decomposition
sklearn.datasets	(not accesible from accessor)
sklearn.dummy	ModelFrame.dummy
sklearn.ensemble	ModelFrame.ensemble
sklearn.feature_extraction	ModelFrame.feature_extraction
sklearn.gaussian_process	ModelFrame.gaussian_process (WIP)
sklearn.grid_search	ModelFrame.grid_search
sklearn.isotonic	ModelFrame.isotonic
sklearn.kernel_approximation	ModelFrame.kernel_approximation
sklearn.lda	ModelFrame.lda
sklearn.linear_model	ModelFrame.linear_model
sklearn.manifold	ModelFrame.manifold
sklearn.metrics	ModelFrame.metrics
sklearn.mixture	ModelFrame.mixture
sklearn.multiclass	ModelFrame.multiclass
sklearn.naive_bayes	ModelFrame.naive_bayes
sklearn.neighbors	ModelFrame.neighbors
sklearn.cross_decomposition	ModelFrame.cross_decomposition (WIP)
sklearn.pipeline	ModelFrame.pipeline
sklearn.preprocessing	ModelFrame.preprocessing, pp
sklearn.qda	ModelFrame.qda
sklearn.semi_supervised	ModelFrame.semi_supervised
sklearn.svm	ModelFrame.svm
sklearn.tree	ModelFrame.tree
sklearn.utils	(not accesible from accessor)

Thus, you can instantiate each estimator via ModelFrame accessors. Once create an estimator, you can pass it to ModelFrame.fit then predict. ModelFrame automatically uses its data and target properties for each operations.

```
>>> estimator = df.cluster.KMeans(n_clusters=3)
>>> df.fit(estimator)
```

```
>>> predicted = df.predict(estimator)
>>> predicted
0      1
1      1
2      1
...
147     2
148     2
149     0
Length: 150, dtype: int32
```

ModelFrame preserves the most recently used estimator in estimator attribute, and predicted results in predicted attribute.

```
>>> df.estimator
KMeans(copy_x=True, init='k-means++', max_iter=300, n_clusters=3, n_init=10,
       n_jobs=1, precompute_distances=True, random_state=None, tol=0.0001,
       verbose=0)

>>> df.predicted
```

```
0      1
1      1
2      1
...
147     2
148     2
149     0
Length: 150, dtype: int32
```

ModelFrame has following methods corresponding to various `scikit-learn` estimators. The last results are saved as corresponding ModelFrame properties.

ModelFrame method	ModelFrame property
ModelFrame.fit	(None)
ModelFrame.transform	(None)
ModelFrame.fit_transform	(None)
ModelFrame.inverse_transform	(None)
ModelFrame.predict	ModelFrame.predicted
ModelFrame.fit_predict	ModelFrame.predicted
ModelFrame.score	(None)
ModelFrame.predict_proba	ModelFrame.proba
ModelFrame.predict_log_proba	ModelFrame.log_proba
ModelFrame.decision_function	ModelFrame.decision

**Note:** If you access to a property before calling ModelFrame methods, ModelFrame automatically calls corresponding method of the latest estimator and return the result.

Following example shows to perform PCA, then revert principal components back to original space.

```
>>> estimator = df.decomposition.PCA()
>>> df.fit(estimator)

>>> transformed = df.transform(estimator)
>>> transformed.head()
   .target      0      1      2      3
0      0 -2.684207 -0.326607  0.021512  0.001006
1      0 -2.715391  0.169557  0.203521  0.099602
2      0 -2.889820  0.137346 -0.024709  0.019305
3      0 -2.746437  0.311124 -0.037672 -0.075955
4      0 -2.728593 -0.333925 -0.096230 -0.063129

>>> type(transformed)
<class 'expandas.core.frame.ModelFrame'>

>>> transformed.inverse_transform(estimator)
   .target      0      1      2      3
0      0  5.1  3.5  1.4  0.2
1      0  4.9  3.0  1.4  0.2
2      0  4.7  3.2  1.3  0.2
3      0  4.6  3.1  1.5  0.2
4      0  5.0  3.6  1.4  0.2
..      ...      ...      ...      ...
145     2  6.7  3.0  5.2  2.3
146     2  6.3  2.5  5.0  1.9
147     2  6.5  3.0  5.2  2.0
148     2  6.2  3.4  5.4  2.3
149     2  5.9  3.0  5.1  1.8
```

[150 rows x 5 columns]

---

**Note:** columns information will be lost once transformed to principal components.

---

If `ModelFrame` both has target and predicted values, the model evaluation can be performed using functions available in `ModelFrame.metrics`.

```
>>> estimator = df.svm.SVC()
>>> df.fit(estimator)

>>> df.predict(estimator)
0      0
1      0
2      0
...
147     2
148     2
149     2
Length: 150, dtype: int64

>>> df.predicted
0      0
1      0
2      0
...
147     2
148     2
149     2
Length: 150, dtype: int64

>>> df.metrics.confusion_matrix()
Predicted   0    1    2
Target
0           50    0    0
1            0   48    2
2            0    0   50
```

## 2.2 Use module level functions

Some `scikit-learn` modules define functions which handle data without instanciating estimators. You can call these functions from accessor methods directly, and `ModelFrame` will pass corresponding data on background. Following example shows to use `sklearn.cluster.k_means` function to perform K-means.

---

**Important:** When you use module level function, `ModelFrame.predicted` WILL NOT be updated. Thus, using estimator is recommended.

---

```
# no need to pass data explicitly
# sklearn.cluster.kmeans returns centroids, cluster labels and inertia
>>> c, l, i = df.cluster.k_means(n_clusters=3)
>>> l
0      1
1      1
2      1
...
```

```
147     2
148     2
149     0
Length: 150, dtype: int32
```

## 2.3 Pipeline

ModelFrame can handle pipeline as the same as normal estimators.

```
>>> estimators = [('reduce_dim', df.decomposition.PCA()),
...               ('svm', df.svm.SVC())]
>>> pipe = df.pipeline.Pipeline(estimators)
>>> df.fit(pipe)

>>> df.predict(pipe)
0     0
1     0
2     0
...
147    2
148    2
149    2
Length: 150, dtype: int64
```

Above expression is the same as below:

```
>>> df2 = df.copy()
>>> df2 = df2.fit_transform(df2.decomposition.PCA())
>>> svm = df2.svm.SVC()
>>> df2.fit(svm)
SVC(C=1.0, cache_size=200, class_weight=None, coef0=0.0, degree=3, gamma=0.0,
    kernel='rbf', max_iter=-1, probability=False, random_state=None,
    shrinking=True, tol=0.001, verbose=False)
>>> df2.predict(svm)
0     0
1     0
2     0
...
147    2
148    2
149    2
Length: 150, dtype: int64
```

## 2.4 Cross Validation

scikit-learn has some classes for cross validation. `cross_validation.train_test_split` splits data to training and test set. You can access to the function via `cross_validation` accessor.

```
>>> train_df, test_df = df.cross_validation.train_test_split()
>>> train_df
   .target  sepal length  sepal width  petal length  petal width
0         0          4.8          3.4          1.9          0.2
1         1          6.3          3.3          4.7          1.6
2         0          4.8          3.4          1.6          0.2
```

```

3          2          7.7          2.6          6.9          2.3
4          0          5.4          3.4          1.7          0.2
..      ...      ...      ...      ...      ...
107         0          5.1          3.7          1.5          0.4
108         1          6.7          3.1          4.7          1.5
109         0          4.7          3.2          1.3          0.2
110         0          5.8          4.0          1.2          0.2
111         0          5.1          3.5          1.4          0.2

```

[112 rows x 5 columns]

```

>>> test_df
   .target  sepal length  sepal width  petal length  petal width
0         2           6.3           2.7           4.9           1.8
1         0           4.5           2.3           1.3           0.3
2         2           5.8           2.8           5.1           2.4
3         0           4.3           3.0           1.1           0.1
4         0           5.0           3.0           1.6           0.2
..      ...      ...      ...      ...      ...
33        1           6.7           3.1           4.4           1.4
34        0           4.6           3.6           1.0           0.2
35        1           5.7           3.0           4.2           1.2
36        1           5.9           3.0           4.2           1.5
37        2           6.4           2.8           5.6           2.1

```

[38 rows x 5 columns]

Also, there are some iterative classes which returns indexes for training sets and test sets. You can slice `ModelFrame` using these indexes.

```

>>> kf = df.cross_validation.KFold(n=150, n_folds=3)
>>> for train_index, test_index in kf:
...     print('training set shape: ', df.iloc[train_index, :].shape,
...           'test set shape: ', df.iloc[test_index, :].shape)
('training set shape: ', (100, 5), 'test set shape: ', (50, 5))
('training set shape: ', (100, 5), 'test set shape: ', (50, 5))
('training set shape: ', (100, 5), 'test set shape: ', (50, 5))

```

For further simplification, `ModelFrame.cross_validation.iterate` can accept such iterators and returns `ModelFrame` corresponding to training and test data.

```

>>> kf = df.cross_validation.KFold(n=150, n_folds=3)
>>> for train_df, test_df in df.cross_validation.iterate(kf):
...     print('training set shape: ', train_df.shape,
...           'test set shape: ', test_df.shape)
('training set shape: ', (100, 5), 'test set shape: ', (50, 5))
('training set shape: ', (100, 5), 'test set shape: ', (50, 5))
('training set shape: ', (100, 5), 'test set shape: ', (50, 5))

```

## 2.5 Grid Search

You can perform grid search using `ModelFrame.fit`.

```

>>> tuned_parameters = [{'kernel': ['rbf'], 'gamma': [1e-3, 1e-4],
...                       'C': [1, 10, 100]},
...                      {'kernel': ['linear'], 'C': [1, 10, 100]}]

```



```
>>> df = expd.ModelFrame(datasets.load_digits())
>>> cv = df.grid_search.GridSearchCV(df.svm.SVC(C=1), tuned_parameters,
...                                 cv=5, scoring='precision')

>>> df.fit(cv)

>>> cv.best_estimator_
SVC(C=10, cache_size=200, class_weight=None, coef0=0.0, degree=3, gamma=0.001,
    kernel='rbf', max_iter=-1, probability=False, random_state=None,
    shrinking=True, tol=0.001, verbose=False)
```

In addition, `ModelFrame.grid_search` has a `describe` function to organize each grid search result as `pd.DataFrame` accepting estimator.

```
>>> df.grid_search.describe(cv)
   mean      std    C  gamma kernel
0  0.974108  0.013139    1  0.0010   rbf
1  0.951416  0.020010    1  0.0001   rbf
2  0.975372  0.011280   10  0.0010   rbf
3  0.962534  0.020218   10  0.0001   rbf
4  0.975372  0.011280  100  0.0010   rbf
5  0.964695  0.016686  100  0.0001   rbf
6  0.951811  0.018410    1     NaN  linear
7  0.951811  0.018410   10     NaN  linear
8  0.951811  0.018410  100     NaN  linear
```

API:



---

## expandas.core package

---

### 3.1 Submodules

### 3.2 expandas.core.accessor module

**class** `expandas.core.accessor.AccessorMethods` (*df, module\_name=None, attrs=None*)

Bases: `object`

Accessor to related functionalities.

### 3.3 expandas.core.frame module

**class** `expandas.core.frame.ModelFrame` (*data, target=None, \*args, \*\*kwargs*)

Bases: `pandas.core.frame.DataFrame`

Data structure subclassing `pandas.DataFrame` to define a metadata to specify target (response variable) and data (explanatory variable / features).

**Parameters** **data** : same as `pandas.DataFrame`

**target** : str or array-like

Column name or values to be used as target

**args** : arguments passed to `pandas.DataFrame`

**kwargs** : keyword arguments passed to `pandas.DataFrame`

#### Attributes

<code>T</code>	Transpose index and columns
<code>at</code>	
<code>axes</code>	
<code>blocks</code>	Internal property, property synonym for <code>as_blocks()</code>
<code>crv</code>	Property to access <code>sklearn.cross_validation</code>
<code>data</code>	Return data (explanatory variable / features)
<code>decision</code>	Return current estimator's decision function
<code>dtypes</code>	Return the dtypes in this object

Continued on next page

Table 3.1 – continued from previous page

<code>empty</code>	True if NDFrame is entirely empty [no items]
<code>estimator</code>	Return most recently used estimator
<code>ftypes</code>	Return the ftypes (indication of sparse/dense and dtype) in this object.
<code>iat</code>	
<code>iloc</code>	
<code>ix</code>	
<code>loc</code>	
<code>log_proba</code>	Return current estimator's log probabilities
<code>ndim</code>	Number of axes / array dimensions
<code>pp</code>	Property to access <code>sklearn.preprocessing</code>
<code>predicted</code>	Return current estimator's predicted results
<code>proba</code>	Return current estimator's probabilities
<code>shape</code>	
<code>size</code>	number of elements in the NDFrame
<code>target</code>	Return target (response variable)
<code>target_name</code>	Return target column name
<code>values</code>	Numpy representation of NDFrame

cluster
covariance
cross_decomposition
cross_validation
decomposition
dummy
ensemble
feature_extraction
feature_selection
gaussian_process
grid_search
is_copy
isotonic
kernel_approximation
lda
linear_model
manifold
metrics
mixture
multiclass
naive_bayes
neighbors
pipeline
preprocessing
qda
semi_supervised
svm
tree

## Methods

<code>abs()</code>	Return an object with absolute value taken.
--------------------	---

Table 3.2 – cont

<code>add(other[, axis, level, fill_value])</code>	Binary operator add with support to substitute a fill_value for missing data in
<code>add_prefix(prefix)</code>	Concatenate prefix string with panel items names.
<code>add_suffix(suffix)</code>	Concatenate suffix string with panel items names
<code>align(other[, join, axis, level, copy, ...])</code>	Align two object on their axes with the
<code>all([axis, bool_only, skipna, level])</code>	Return whether all elements are True over requested axis
<code>any([axis, bool_only, skipna, level])</code>	Return whether any element is True over requested axis
<code>append(other[, ignore_index, verify_integrity])</code>	Append columns of other to end of this frame's columns and index, returning
<code>apply(func[, axis, broadcast, raw, reduce, args])</code>	Applies function along input axis of DataFrame.
<code>applymap(func)</code>	Apply a function to a DataFrame that is intended to operate elementwise, i.e.
<code>as_blocks()</code>	Convert the frame to a dict of dtype -> Constructor Types that each has a hom
<code>as_matrix([columns])</code>	Convert the frame to its Numpy-array representation.
<code>asfreq(freq[, method, how, normalize])</code>	Convert all TimeSeries inside to specified frequency using DateOffset objects
<code>astype(dtype[, copy, raise_on_error])</code>	Cast object to input numpy.dtype
<code>at_time(time[, asof])</code>	Select values at particular time of day (e.g.
<code>between_time(start_time, end_time[, ...])</code>	Select values between particular times of the day (e.g., 9:00-9:30 AM)
<code>bfill([axis, inplace, limit, downcast])</code>	Synonym for NDFrame.fillna(method='bfill')
<code>bool()</code>	Return the bool of a single element PandasObject
<code>boxplot([column, by, ax, fontsize, rot, ...])</code>	Make a box plot from DataFrame column optionally grouped by some colum
<code>clip([lower, upper, out])</code>	Trim values at input threshold(s)
<code>clip_lower(threshold)</code>	Return copy of the input with values below given value truncated
<code>clip_upper(threshold)</code>	Return copy of input with values above given value truncated
<code>combine(other, func[, fill_value, overwrite])</code>	Add two DataFrame objects and do not propagate NaN values, so if for a
<code>combineAdd(other)</code>	Add two DataFrame objects and do not propagate
<code>combineMult(other)</code>	Multiply two DataFrame objects and do not propagate NaN values, so if
<code>combine_first(other)</code>	Combine two DataFrame objects and default to non-null values in frame calli
<code>compound([axis, skipna, level])</code>	Return the compound percentage of the values for the requested axis
<code>consolidate([inplace])</code>	Compute NDFrame with "consolidated" internals (data of each dtype groupe
<code>convert_objects([convert_dates, ...])</code>	Attempt to infer better dtype for object columns
<code>copy([deep])</code>	Make a copy of this object
<code>corr([method, min_periods])</code>	Compute pairwise correlation of columns, excluding NA/null values
<code>corrwith(other[, axis, drop])</code>	Compute pairwise correlation between rows or columns of two DataFrame ob
<code>count([axis, level, numeric_only])</code>	Return Series with number of non-NA/null observations over requested axis.
<code>cov([min_periods])</code>	Compute pairwise covariance of columns, excluding NA/null values
<code>cummax([axis, dtype, out, skipna])</code>	Return cumulative max over requested axis.
<code>cummin([axis, dtype, out, skipna])</code>	Return cumulative min over requested axis.
<code>cumprod([axis, dtype, out, skipna])</code>	Return cumulative prod over requested axis.
<code>cumsum([axis, dtype, out, skipna])</code>	Return cumulative sum over requested axis.
<code>decision_function(estimator, *args, **kwargs)</code>	Call estimator's decision_function method.
<code>describe([percentile_width, percentiles, ...])</code>	Generate various summary statistics, excluding NaN values.
<code>diff([periods])</code>	1st discrete difference of object
<code>div(other[, axis, level, fill_value])</code>	Binary operator truediv with support to substitute a fill_value for missing data
<code>divide(other[, axis, level, fill_value])</code>	Binary operator truediv with support to substitute a fill_value for missing data
<code>dot(other)</code>	Matrix multiplication with DataFrame or Series objects
<code>drop(labels[, axis, level, inplace])</code>	Return new object with labels in requested axis removed
<code>drop_duplicates(*args, **kwargs)</code>	Return DataFrame with duplicate rows removed, optionally only
<code>dropna([axis, how, thresh, subset, inplace])</code>	Return object with labels on given axis omitted where alternately any
<code>duplicated(*args, **kwargs)</code>	Return boolean Series denoting duplicate rows, optionally only
<code>eq(other[, axis, level])</code>	Wrapper for flexible comparison methods eq
<code>equals(other)</code>	Determines if two NDFrame objects contain the same elements.
<code>eval(expr, **kwargs)</code>	Evaluate an expression in the context of the calling DataFrame instance.
<code>ffill([axis, inplace, limit, downcast])</code>	Synonym for NDFrame.fillna(method='ffill')
<code>fillna([value, method, axis, inplace, ...])</code>	Fill NA/NaN values using the specified method

Table 3.2 – cont

<code>filter([items, like, regex, axis])</code>	Restrict the info axis to set of items or wildcard
<code>first(offset)</code>	Convenience method for subsetting initial periods of time series data
<code>first_valid_index()</code>	Return label for first non-NA/null value
<code>fit(estimator, *args, **kwargs)</code>	Call estimator’s fit method.
<code>fit_predict(estimator, *args, **kwargs)</code>	Call estimator’s fit_predict method.
<code>fit_transform(estimator, *args, **kwargs)</code>	Call estimator’s fit_transform method.
<code>floordiv(other[, axis, level, fill_value])</code>	Binary operator floordiv with support to substitute a fill_value for missing data
<code>from_csv(path[, header, sep, index_col, ...])</code>	Read delimited file into DataFrame
<code>from_dict(data[, orient, dtype])</code>	Construct DataFrame from dict of array-like or dicts
<code>from_items(items[, columns, orient])</code>	Convert (key, value) pairs to DataFrame.
<code>from_records(data[, index, exclude, ...])</code>	Convert structured or record ndarray to DataFrame
<code>ge(other[, axis, level])</code>	Wrapper for flexible comparison methods ge
<code>get(key[, default])</code>	Get item from object for given key (DataFrame column, Panel slice, etc.).
<code>get_dtype_counts()</code>	Return the counts of dtypes in this object
<code>get_ftype_counts()</code>	Return the counts of ftypes in this object
<code>get_value(index, col[, takeable])</code>	Quickly retrieve single value at passed column and index
<code>get_values()</code>	same as values (but handles sparseness conversions)
<code>groupby([by, axis, level, as_index, sort, ...])</code>	Group series using mapper (dict or key function, apply given function
<code>gt(other[, axis, level])</code>	Wrapper for flexible comparison methods gt
<code>has_data()</code>	Return whether ModelFrame has data
<code>has_target()</code>	Return whether ModelFrame has target
<code>head([n])</code>	Returns first n rows
<code>hist(data[, column, by, grid, xlabelsize, ...])</code>	Draw histogram of the DataFrame’s series using matplotlib / pylab.
<code>icol(i)</code>	
<code>idxmax([axis, skipna])</code>	Return index of first occurrence of maximum over requested axis.
<code>idxmin([axis, skipna])</code>	Return index of first occurrence of minimum over requested axis.
<code>iget_value(i, j)</code>	
<code>info([verbose, buf, max_cols, memory_usage, ...])</code>	Concise summary of a DataFrame.
<code>insert(loc, column, value[, allow_duplicates])</code>	Insert column into DataFrame at specified location.
<code>interpolate([method, axis, limit, inplace, ...])</code>	Interpolate values according to different methods.
<code>inverse_transform(estimator, *args, **kwargs)</code>	Call estimator’s inverse_transform method.
<code>irow(i[, copy])</code>	
<code>isin(values)</code>	Return boolean DataFrame showing whether each element in the DataFrame
<code>isnull()</code>	Return a boolean same-sized object indicating if the values are null
<code>iteritems()</code>	Iterator over (column, series) pairs
<code>iterkv(*args, **kwargs)</code>	iteritems alias used to get around 2to3. Deprecated
<code>iterrows()</code>	Iterate over rows of DataFrame as (index, Series) pairs.
<code>itertuples([index])</code>	Iterate over rows of DataFrame as tuples, with index value
<code>join(other[, on, how, lsuffix, rsuffix, sort])</code>	Join columns with other DataFrame either on index or on a key column.
<code>keys()</code>	Get the ‘info axis’ (see Indexing for more)
<code>kurt([axis, skipna, level, numeric_only])</code>	Return unbiased kurtosis over requested axis
<code>kurtosis([axis, skipna, level, numeric_only])</code>	Return unbiased kurtosis over requested axis
<code>last(offset)</code>	Convenience method for subsetting final periods of time series data
<code>last_valid_index()</code>	Return label for last non-NA/null value
<code>le(other[, axis, level])</code>	Wrapper for flexible comparison methods le
<code>load(path)</code>	Deprecated.
<code>lookup(row_labels, col_labels)</code>	Label-based “fancy indexing” function for DataFrame.
<code>lt(other[, axis, level])</code>	Wrapper for flexible comparison methods lt
<code>mad([axis, skipna, level])</code>	Return the mean absolute deviation of the values for the requested axis
<code>mask(cond)</code>	Returns copy whose values are replaced with nan if the
<code>max([axis, skipna, level, numeric_only])</code>	This method returns the maximum of the values in the object.
<code>mean([axis, skipna, level, numeric_only])</code>	Return the mean of the values for the requested axis

Table 3.2 – cont

<code>median([axis, skipna, level, numeric_only])</code>	Return the median of the values for the requested axis
<code>memory_usage([index])</code>	Memory usage of DataFrame columns.
<code>merge(right[, how, on, left_on, right_on, ...])</code>	Merge DataFrame objects by performing a database-style join operation by co
<code>min([axis, skipna, level, numeric_only])</code>	This method returns the minimum of the values in the object.
<code>mod(other[, axis, level, fill_value])</code>	Binary operator mod with support to substitute a fill_value for missing data in
<code>mode([axis, numeric_only])</code>	Gets the mode of each element along the axis selected.
<code>mul(other[, axis, level, fill_value])</code>	Binary operator mul with support to substitute a fill_value for missing data in
<code>multiply(other[, axis, level, fill_value])</code>	Binary operator mul with support to substitute a fill_value for missing data in
<code>ne(other[, axis, level])</code>	Wrapper for flexible comparison methods ne
<code>notnull()</code>	Return a boolean same-sized object indicating if the values are
<code>pct_change([periods, fill_method, limit, freq])</code>	Percent change over given number of periods.
<code>pivot([index, columns, values])</code>	Reshape data (produce a “pivot” table) based on column values.
<code>pivot_table(*args, **kwargs)</code>	Create a spreadsheet-style pivot table as a DataFrame.
<code>plot(data[, x, y, kind, ax, subplots, ...])</code>	Make plots of DataFrame using matplotlib / pylab.
<code>pop(item)</code>	Return item and drop from frame.
<code>pow(other[, axis, level, fill_value])</code>	Binary operator pow with support to substitute a fill_value for missing data in
<code>predict(estimator, *args, **kwargs)</code>	Call estimator’s predict method.
<code>predict_log_proba(estimator, *args, **kwargs)</code>	Call estimator’s predict_log_proba method.
<code>predict_proba(estimator, *args, **kwargs)</code>	Call estimator’s predict_proba method.
<code>prod([axis, skipna, level, numeric_only])</code>	Return the product of the values for the requested axis
<code>product([axis, skipna, level, numeric_only])</code>	Return the product of the values for the requested axis
<code>quantile([q, axis, numeric_only])</code>	Return values at the given quantile over requested axis, a la numpy.percentile
<code>query(expr, **kwargs)</code>	Query the columns of a frame with a boolean expression.
<code>radd(other[, axis, level, fill_value])</code>	Binary operator radd with support to substitute a fill_value for missing data in
<code>rank([axis, numeric_only, method, ...])</code>	Compute numerical data ranks (1 through n) along axis.
<code>rdiv(other[, axis, level, fill_value])</code>	Binary operator rtruediv with support to substitute a fill_value for missing da
<code>reindex([index, columns])</code>	Conform DataFrame to new index with optional filling logic, placing NA/NaN
<code>reindex_axis(labels[, axis, method, level, ...])</code>	Conform input object to new index with optional filling logic, placing NA/NaN
<code>reindex_like(other[, method, copy, limit])</code>	return an object with matching indicies to myself
<code>rename([index, columns])</code>	Alter axes input function or functions.
<code>rename_axis(mapper[, axis, copy, inplace])</code>	Alter index and / or columns using input function or functions.
<code>reorder_levels(order[, axis])</code>	Rearrange index levels using input order.
<code>replace([to_replace, value, inplace, limit, ...])</code>	Replace values given in ‘to_replace’ with ‘value’.
<code>resample(rule[, how, axis, fill_method, ...])</code>	Convenience method for frequency conversion and resampling of regular time
<code>reset_index([level, drop, inplace, ...])</code>	For DataFrame with multi-level index, return new DataFrame with labeling in
<code>rfloordiv(other[, axis, level, fill_value])</code>	Binary operator rfloordiv with support to substitute a fill_value for missing da
<code>rmod(other[, axis, level, fill_value])</code>	Binary operator rmod with support to substitute a fill_value for missing data in
<code>rmul(other[, axis, level, fill_value])</code>	Binary operator rmul with support to substitute a fill_value for missing data in
<code>rpow(other[, axis, level, fill_value])</code>	Binary operator rpow with support to substitute a fill_value for missing data in
<code>rsub(other[, axis, level, fill_value])</code>	Binary operator rsub with support to substitute a fill_value for missing data in
<code>rtruediv(other[, axis, level, fill_value])</code>	Binary operator rtruediv with support to substitute a fill_value for missing da
<code>save(path)</code>	Deprecated.
<code>score(estimator, *args, **kwargs)</code>	Call estimator’s score method.
<code>select(crit[, axis])</code>	Return data corresponding to axis labels matching criteria
<code>select_dtypes([include, exclude])</code>	Return a subset of a DataFrame including/excluding columns based on their c
<code>sem([axis, skipna, level, ddof])</code>	Return unbiased standard error of the mean over requested axis.
<code>set_axis(axis, labels)</code>	public version of axis assignment
<code>set_index(keys[, drop, append, inplace, ...])</code>	Set the DataFrame index (row labels) using one or more existing columns.
<code>set_value(index, col, value[, takeable])</code>	Put single value at passed column and index
<code>shift([periods, freq, axis])</code>	Shift index by desired number of periods with an optional time freq
<code>skew([axis, skipna, level, numeric_only])</code>	Return unbiased skew over requested axis
<code>slice_shift([periods, axis])</code>	Equivalent to <i>shift</i> without copying data.

Table 3.2 – cont

<code>sort([columns, axis, ascending, inplace, ...])</code>	Sort DataFrame either by labels (along either axis) or by the values in
<code>sort_index([axis, by, ascending, inplace, ...])</code>	Sort DataFrame either by labels (along either axis) or by the values in
<code>sortlevel([level, axis, ascending, inplace, ...])</code>	Sort multilevel index by chosen axis and primary level.
<code>squeeze()</code>	squeeze length 1 dimensions
<code>stack([level, dropna])</code>	Pivot a level of the (possibly hierarchical) column labels, returning a DataFrame
<code>std([axis, skipna, level, ddof])</code>	Return unbiased standard deviation over requested axis.
<code>sub(other[, axis, level, fill_value])</code>	Binary operator sub with support to substitute a fill_value for missing data in
<code>subtract(other[, axis, level, fill_value])</code>	Binary operator sub with support to substitute a fill_value for missing data in
<code>sum([axis, skipna, level, numeric_only])</code>	Return the sum of the values for the requested axis
<code>swapaxes(axis1, axis2[, copy])</code>	Interchange axes and swap values axes appropriately
<code>swaplevel(i, j[, axis])</code>	Swap levels i and j in a MultiIndex on a particular axis
<code>tail([n])</code>	Returns last n rows
<code>take(indices[, axis, convert, is_copy])</code>	Analogous to ndarray.take
<code>to_clipboard([excel, sep])</code>	Attempt to write text representation of object to the system clipboard This can
<code>to_csv(*args, **kwargs)</code>	Write DataFrame to a comma-separated values (csv) file
<code>to_dense()</code>	Return dense representation of NDFrame (as opposed to sparse)
<code>to_dict(*args, **kwargs)</code>	Convert DataFrame to dictionary.
<code>to_excel(*args, **kwargs)</code>	Write DataFrame to a excel sheet
<code>to_gbq(destination_table[, project_id, ...])</code>	Write a DataFrame to a Google BigQuery table.
<code>to_hdf(path_or_buf, key, **kwargs)</code>	activate the HDFStore
<code>to_html([buf, columns, col_space, colSpace, ...])</code>	Render a DataFrame as an HTML table.
<code>to_json([path_or_buf, orient, date_format, ...])</code>	Convert the object to a JSON string.
<code>to_latex([buf, columns, col_space, ...])</code>	Render a DataFrame to a tabular environment table.
<code>to_msgpack([path_or_buf])</code>	msgpack (serialize) object to input file path
<code>to_panel()</code>	Transform long (stacked) format (DataFrame) into wide (3D, Panel) format.
<code>to_period([freq, axis, copy])</code>	Convert DataFrame from DatetimeIndex to PeriodIndex with desired
<code>to_pickle(path)</code>	Pickle (serialize) object to input file path
<code>to_records([index, convert_datetime64])</code>	Convert DataFrame to record array.
<code>to_sparse([fill_value, kind])</code>	Convert to SparseDataFrame
<code>to_sql(name, con[, flavor, schema, ...])</code>	Write records stored in a DataFrame to a SQL database.
<code>to_stata(fname[, convert_dates, ...])</code>	A class for writing Stata binary dta files from array-like objects
<code>to_string([buf, columns, col_space, ...])</code>	Render a DataFrame to a console-friendly tabular output.
<code>to_timestamp([freq, how, axis, copy])</code>	Cast to DatetimeIndex of timestamps, at <i>beginning</i> of period
<code>to_wide(*args, **kwargs)</code>	
<code>transform(estimator, *args, **kwargs)</code>	Call estimator's transform method.
<code>transpose()</code>	Transpose index and columns
<code>truediv(other[, axis, level, fill_value])</code>	Binary operator truediv with support to substitute a fill_value for missing data
<code>truncate([before, after, axis, copy])</code>	Truncates a sorted NDFrame before and/or after some particular dates.
<code>tshift([periods, freq, axis])</code>	Shift the time index, using the index's frequency if available
<code>tz_convert(tz[, axis, level, copy])</code>	Convert the axis to target time zone.
<code>tz_localize(*args, **kwargs)</code>	Localize tz-naive TimeSeries to target time zone
<code>unstack([level])</code>	Pivot a level of the (necessarily hierarchical) index labels, returning a DataFrame
<code>update(other[, join, overwrite, ...])</code>	Modify DataFrame in place using non-NA values from passed DataFrame.
<code>var([axis, skipna, level, ddof])</code>	Return unbiased variance over requested axis.
<code>where(cond[, other, inplace, axis, level, ...])</code>	Return an object of same shape as self and whose corresponding entries are fr
<code>xs(key[, axis, level, copy, drop_level])</code>	Returns a cross-section (row(s) or column(s)) from the Series/DataFrame.

**cluster = None**

**covariance = None**

**cross\_decomposition = None**

**cross\_validation = None**



**crv**  
Property to access `sklearn.cross_validation`

**data**  
Return data (explanatory variable / features)  
**Returns data** : `ModelFrame`

**decision**  
Return current estimator's decision function  
**Returns probabilities** : `ModelFrame`

**decision\_function** (*estimator*, \*args, \*\*kwargs)  
Call estimator's `decision_function` method.  
**Parameters args** : arguments passed to `decision_function` method  
**kwargs** : keyword arguments passed to `decision_function` method  
**Returns returned** : decisions

**decomposition** = `None`

**dummy** = `None`

**ensemble** = `None`

**estimator**  
Return most recently used estimator  
**Returns estimator** : estimator

**feature\_extraction** = `None`

**feature\_selection** = `None`

**fit** (*estimator*, \*args, \*\*kwargs)  
Call estimator's `fit` method.  
**Parameters args** : arguments passed to `fit` method  
**kwargs** : keyword arguments passed to `fit` method  
**Returns returned** : `None` or fitted estimator

**fit\_predict** (*estimator*, \*args, \*\*kwargs)  
Call estimator's `fit_predict` method.  
**Parameters args** : arguments passed to `fit_predict` method  
**kwargs** : keyword arguments passed to `fit_predict` method  
**Returns returned** : predicted result

**fit\_transform** (*estimator*, \*args, \*\*kwargs)  
Call estimator's `fit_transform` method.  
**Parameters args** : arguments passed to `fit_transform` method  
**kwargs** : keyword arguments passed to `fit_transform` method  
**Returns returned** : transformed result

**gaussian\_process** = `None`

**grid\_search** = `None`

**has\_data()**  
Return whether `ModelFrame` has data

**Returns** `has_data` : bool

**has\_target()**  
Return whether `ModelFrame` has target

**Returns** `has_target` : bool

**inverse\_transform(*estimator*, \**args*, \*\**kwargs*)**  
Call estimator's `inverse_transform` method.

**Parameters** `args` : arguments passed to `inverse_transform` method

**kwargs** : keyword arguments passed to `inverse_transform` method

**Returns** `returned` : transformed result

**isotonic** = None

**kernel\_approximation** = None

**lda** = None

**linear\_model** = None

**log\_proba**  
Return current estimator's log probabilities

**Returns** `probabilities` : `ModelFrame`

**manifold** = None

**metrics** = None

**mixture** = None

**multiclass** = None

**naive\_bayes** = None

**neighbors** = None

**pipeline** = None

**pp**  
Property to access `sklearn.preprocessing`

**predict(*estimator*, \**args*, \*\**kwargs*)**  
Call estimator's `predict` method.

**Parameters** `args` : arguments passed to `predict` method

**kwargs** : keyword arguments passed to `predict` method

**Returns** `returned` : predicted result

**predict\_log\_proba(*estimator*, \**args*, \*\**kwargs*)**  
Call estimator's `predict_log_proba` method.

**Parameters** `args` : arguments passed to `predict_log_proba` method

**kwargs** : keyword arguments passed to `predict_log_proba` method

**Returns** `returned` : probabilities

**predict\_proba(*estimator*, \**args*, \*\**kwargs*)**  
Call estimator's `predict_proba` method.

**Parameters** **args** : arguments passed to predict\_proba method

**kwargs** : keyword arguments passed to predict\_proba method

**Returns** **returned** : probabilities

**predicted**  
Return current estimator's predicted results

**Returns** **predicted** : `ModelSeries`

**preprocessing** = `None`

**proba**  
Return current estimator's probabilities

**Returns** **probabilities** : `ModelFrame`

**qda** = `None`

**score** (*estimator*, \**args*, \*\**kwargs*)  
Call estimator's score method.

**Parameters** **args** : arguments passed to score method

**kwargs** : keyword arguments passed to score method

**Returns** **returned** : score

**semi\_supervised** = `None`

**svm** = `None`

**target**  
Return target (response variable)

**Returns** **target** : `ModelSeries`

**target\_name**  
Return target column name

**Returns** **target** : object

**transform** (*estimator*, \**args*, \*\**kwargs*)  
Call estimator's transform method.

**Parameters** **args** : arguments passed to transform method

**kwargs** : keyword arguments passed to transform method

**Returns** **returned** : transformed result

**tree** = `None`

## 3.4 expandas.core.series module

**class** `expandas.core.series.ModelSeries` (*data=None*, *index=None*, *dtype=None*, *name=None*,  
*copy=False*, *fastpath=False*)  
 Bases: `pandas.core.series.Series`  
 Wrapper for `pandas.Series` to support `sklearn.preprocessing`

## Attributes

T	return the transpose, which is by definition self
at	
axes	
base	return the base object if the memory of the underlying data is shared
blocks	Internal property, property synonym for as_blocks()
data	return the data pointer of the underlying data
dtype	return the dtype object of the underlying data
dtypes	return the dtype object of the underlying data
empty	True if NDFrame is entirely empty [no items]
flags	return the ndarray.flags for the underlying data
ftype	return if the data is sparseldense
ftypes	return if the data is sparseldense
iat	
iloc	
imag	
is_time_series	
itemsizes	return the size of the dtype of the item of the underlying data
ix	
loc	
nbytes	return the number of bytes in the underlying data
ndim	return the number of dimensions of the underlying data, by definition 1
pp	Property to access sklearn.preprocessing
real	
shape	return a tuple of the shape of the underlying data
size	return the number of elements in the underlying data
strides	return the strides of the underlying data
values	Return Series as ndarray

cat	
dt	
is_copy	
preprocessing	
str	

## Methods

abs()	Return an object with absolute value taken.
add(other[, level, fill_value, axis])	Binary operator add with support to substitute a fill_value for missing data
add_prefix(prefix)	Concatenate prefix string with panel items names.
add_suffix(suffix)	Concatenate suffix string with panel items names
align(other[, join, axis, level, copy, ...])	Align two object on their axes with the
all([axis, bool_only, skipna, level])	Return whether all elements are True over requested axis
any([axis, bool_only, skipna, level])	Return whether any element is True over requested axis
append(to_append[, verify_integrity])	Concatenate two or more Series.
apply(func[, convert_dtype, args])	Invoke function on values of Series.
argmax([axis, out, skipna])	Index of first occurrence of maximum of values.
argmin([axis, out, skipna])	Index of first occurrence of minimum of values.
argsort([axis, kind, order])	Overrides ndarray.argsort.
as_blocks()	Convert the frame to a dict of dtype -> Constructor Types that each has a homog
as_matrix([columns])	Convert the frame to its Numpy-array representation.

Table 3.4 – continued from previous page

<code>asfreq(freq[, method, how, normalize])</code>	Convert all TimeSeries inside to specified frequency using DateOffset objects.
<code>asof(when)</code>	Return last good (non-NaN) value in TimeSeries if value is NaN for requested date
<code>astype(dtype[, copy, raise_on_error])</code>	Cast object to input numpy.dtype
<code>at_time(time[, asof])</code>	Select values at particular time of day (e.g.
<code>autocorr()</code>	Lag-1 autocorrelation
<code>between(left, right[, inclusive])</code>	Return boolean Series equivalent to <code>left &lt;= series &lt;= right</code> .
<code>between_time(start_time, end_time[, ...])</code>	Select values between particular times of the day (e.g., 9:00-9:30 AM)
<code>bfill([axis, inplace, limit, downcast])</code>	Synonym for <code>NDFrame.fillna(method='bfill')</code>
<code>bool()</code>	Return the bool of a single element PandasObject
<code>clip([lower, upper, out])</code>	Trim values at input threshold(s)
<code>clip_lower(threshold)</code>	Return copy of the input with values below given value truncated
<code>clip_upper(threshold)</code>	Return copy of input with values above given value truncated
<code>combine(other, func[, fill_value])</code>	Perform elementwise binary operation on two Series using given function
<code>combine_first(other)</code>	Combine Series values, choosing the calling Series's values first.
<code>compound([axis, skipna, level])</code>	Return the compound percentage of the values for the requested axis
<code>compress(condition[, axis, out])</code>	Return selected slices of an array along given axis as a Series
<code>consolidate([inplace])</code>	Compute NDFrame with "consolidated" internals (data of each dtype grouped together)
<code>convert_objects([convert_dates, ...])</code>	Attempt to infer better dtype for object columns
<code>copy([deep])</code>	Make a copy of this object
<code>corr(other[, method, min_periods])</code>	Compute correlation with <i>other</i> Series, excluding missing values
<code>count([level])</code>	Return number of non-NA/null observations in the Series
<code>cov(other[, min_periods])</code>	Compute covariance with Series, excluding missing values
<code>cummax([axis, dtype, out, skipna])</code>	Return cumulative max over requested axis.
<code>cummin([axis, dtype, out, skipna])</code>	Return cumulative min over requested axis.
<code>cumprod([axis, dtype, out, skipna])</code>	Return cumulative prod over requested axis.
<code>cumsum([axis, dtype, out, skipna])</code>	Return cumulative sum over requested axis.
<code>describe([percentile_width, percentiles, ...])</code>	Generate various summary statistics, excluding NaN values.
<code>diff([periods])</code>	1st discrete difference of object
<code>div(other[, level, fill_value, axis])</code>	Binary operator <code>truediv</code> with support to substitute a <code>fill_value</code> for missing data
<code>divide(other[, level, fill_value, axis])</code>	Binary operator <code>truediv</code> with support to substitute a <code>fill_value</code> for missing data
<code>dot(other)</code>	Matrix multiplication with DataFrame or inner-product with Series
<code>drop(labels[, axis, level, inplace])</code>	Return new object with labels in requested axis removed
<code>drop_duplicates([take_last, inplace])</code>	Return Series with duplicate values removed
<code>dropna([axis, inplace])</code>	Return Series without null values
<code>duplicated([take_last])</code>	Return boolean Series denoting duplicate values
<code>eq(other)</code>	
<code>equals(other)</code>	Determines if two NDFrame objects contain the same elements.
<code>factorize([sort, na_sentinel])</code>	Encode the object as an enumerated type or categorical variable
<code>ffill([axis, inplace, limit, downcast])</code>	Synonym for <code>NDFrame.fillna(method='ffill')</code>
<code>fillna([value, method, axis, inplace, ...])</code>	Fill NA/NaN values using the specified method
<code>filter([items, like, regex, axis])</code>	Restrict the info axis to set of items or wildcard
<code>first(offset)</code>	Convenience method for subsetting initial periods of time series data
<code>first_valid_index()</code>	Return label for first non-NA/null value
<code>floordiv(other[, level, fill_value, axis])</code>	Binary operator <code>floordiv</code> with support to substitute a <code>fill_value</code> for missing data
<code>from_array(arr[, index, name, dtype, copy, ...])</code>	
<code>from_csv(path[, sep, parse_dates, header, ...])</code>	Read delimited file into Series
<code>ge(other)</code>	
<code>get(key[, default])</code>	Get item from object for given key (DataFrame column, Panel slice, etc.).
<code>get_dtype_counts()</code>	Return the counts of dtypes in this object
<code>get_ftype_counts()</code>	Return the counts of ftypes in this object
<code>get_value(label[, takeable])</code>	Quickly retrieve single value at passed index label
<code>get_values()</code>	same as <code>values</code> (but handles sparseness conversions); is a view

Table 3.4 – continued from previous page

<code>groupby([by, axis, level, as_index, sort, ...])</code>	Group series using mapper (dict or key function, apply given function
<code>gt(other)</code>	return if I have any nans; enables various perf speedups
<code>hasnans()</code>	Returns first <i>n</i> rows
<code>head([n])</code>	Draw histogram of the input series using matplotlib
<code>hist([by, ax, grid, xlabelsize, xrot, ...])</code>	Index of first occurrence of maximum of values.
<code>idxmax([axis, out, skipna])</code>	Index of first occurrence of minimum of values.
<code>idxmin([axis, out, skipna])</code>	Return the <i>i</i> -th value or values in the Series by location
<code>iget(i[, axis])</code>	Return the <i>i</i> -th value or values in the Series by location
<code>iget_value(i[, axis])</code>	Interpolate values according to different methods.
<code>interpolate([method, axis, limit, inplace, ...])</code>	Return the <i>i</i> -th value or values in the Series by location
<code>irow(i[, axis])</code>	Return a boolean Series showing whether each element in the Series is <i>exa</i>
<code>isin(values)</code>	Return a boolean same-sized object indicating if the values are null
<code>isnull()</code>	return the first element of the underlying data as a python scalar
<code>item()</code>	Lazily iterate over (index, value) tuples
<code>iteritems()</code>	iteritems alias used to get around 2to3. Deprecated
<code>iterkv(*args, **kwargs)</code>	Alias for index
<code>keys()</code>	Return unbiased kurtosis over requested axis
<code>kurt([axis, skipna, level, numeric_only])</code>	Return unbiased kurtosis over requested axis
<code>kurtosis([axis, skipna, level, numeric_only])</code>	Convenience method for subsetting final periods of time series data
<code>last(offset)</code>	Return label for last non-NA/null value
<code>last_valid_index()</code>	
<code>le(other)</code>	Deprecated.
<code>load(path)</code>	
<code>lt(other)</code>	
<code>mad([axis, skipna, level])</code>	Return the mean absolute deviation of the values for the requested axis
<code>map(arg[, na_action])</code>	Map values of Series using input correspondence (which can be
<code>mask(cond)</code>	Returns copy whose values are replaced with nan if the
<code>max([axis, skipna, level, numeric_only])</code>	This method returns the maximum of the values in the object.
<code>mean([axis, skipna, level, numeric_only])</code>	Return the mean of the values for the requested axis
<code>median([axis, skipna, level, numeric_only])</code>	Return the median of the values for the requested axis
<code>min([axis, skipna, level, numeric_only])</code>	This method returns the minimum of the values in the object.
<code>mod(other[, level, fill_value, axis])</code>	Binary operator mod with support to substitute a fill_value for missing data
<code>mode()</code>	Returns the mode(s) of the dataset.
<code>mul(other[, level, fill_value, axis])</code>	Binary operator mul with support to substitute a fill_value for missing data
<code>multiply(other[, level, fill_value, axis])</code>	Binary operator mul with support to substitute a fill_value for missing data
<code>ne(other)</code>	
<code>nlargest([n, take_last])</code>	Return the largest <i>n</i> elements.
<code>nonzero()</code>	Return the indices of the elements that are non-zero
<code>notnull()</code>	Return a boolean same-sized object indicating if the values are
<code>nsmallest([n, take_last])</code>	Return the smallest <i>n</i> elements.
<code>nunique([dropna])</code>	Return number of unique elements in the object.
<code>order([na_last, ascending, kind, ...])</code>	Sorts Series object, by value, maintaining index-value link.
<code>pct_change([periods, fill_method, limit, freq])</code>	Percent change over given number of periods.
<code>plot(data[, kind, ax, figsize, use_index, ...])</code>	Make plots of Series using matplotlib / pylab.
<code>pop(item)</code>	Return item and drop from frame.
<code>pow(other[, level, fill_value, axis])</code>	Binary operator pow with support to substitute a fill_value for missing data
<code>prod([axis, skipna, level, numeric_only])</code>	Return the product of the values for the requested axis
<code>product([axis, skipna, level, numeric_only])</code>	Return the product of the values for the requested axis
<code>ptp([axis, out])</code>	
<code>put(*args, **kwargs)</code>	return a ndarray with the values put
<code>quantile([q])</code>	Return value at the given quantile, a la numpy.percentile.
<code>radd(other[, level, fill_value, axis])</code>	Binary operator radd with support to substitute a fill_value for missing data

Table 3.4 – continued from previous page

<code>rank([method, na_option, ascending, pct])</code>	Compute data ranks (1 through n).
<code>ravel([order])</code>	Return the flattened underlying data as an ndarray
<code>rdiv(other[, level, fill_value, axis])</code>	Binary operator <code>rtruediv</code> with support to substitute a <code>fill_value</code> for missing data
<code>reindex([index])</code>	Conform Series to new index with optional filling logic, placing NA/NaN in location for compatibility with higher dims
<code>reindex_axis(labels[, axis])</code>	return an object with matching indicies to myself
<code>reindex_like(other[, method, copy, limit])</code>	Alter axes input function or functions.
<code>rename([index])</code>	Alter index and / or columns using input function or functions.
<code>rename_axis(mapper[, axis, copy, inplace])</code>	Rearrange index levels using input order.
<code>reorder_levels(order)</code>	return a new Series with the values repeated <code>reps</code> times
<code>repeat(reps)</code>	Replace values given in 'to_replace' with 'value'.
<code>replace([to_replace, value, inplace, limit, ...])</code>	Convenience method for frequency conversion and resampling of regular time-series
<code>resample(rule[, how, axis, fill_method, ...])</code>	Analogous to the <code>pandas.DataFrame.reset_index()</code> function, see documentation
<code>reset_index([level, drop, name, inplace])</code>	return an ndarray with the values shape
<code>reshape(*args, **kwargs)</code>	Binary operator <code>rfloordiv</code> with support to substitute a <code>fill_value</code> for missing data
<code>rfloordiv(other[, level, fill_value, axis])</code>	Binary operator <code>rmod</code> with support to substitute a <code>fill_value</code> for missing data
<code>rmod(other[, level, fill_value, axis])</code>	Binary operator <code>rmul</code> with support to substitute a <code>fill_value</code> for missing data
<code>rmul(other[, level, fill_value, axis])</code>	Return <i>a</i> with each element rounded to the given number of decimals.
<code>round([decimals, out])</code>	Binary operator <code>rpow</code> with support to substitute a <code>fill_value</code> for missing data
<code>rpow(other[, level, fill_value, axis])</code>	Binary operator <code>rsub</code> with support to substitute a <code>fill_value</code> for missing data
<code>rsub(other[, level, fill_value, axis])</code>	Binary operator <code>rtruediv</code> with support to substitute a <code>fill_value</code> for missing data
<code>rtruediv(other[, level, fill_value, axis])</code>	Deprecated.
<code>save(path)</code>	Find indices where elements should be inserted to maintain order.
<code>searchsorted(v[, side, sorter])</code>	Return data corresponding to axis labels matching criteria
<code>select(crit[, axis])</code>	Return unbiased standard error of the mean over requested axis.
<code>sem([axis, skipna, level, ddof])</code>	public version of axis assignment
<code>set_axis(axis, labels)</code>	Quickly set single value at passed label.
<code>set_value(label, value[, takeable])</code>	Shift index by desired number of periods with an optional time frequency
<code>shift([periods, freq, axis])</code>	Return unbiased skew over requested axis
<code>skew([axis, skipna, level, numeric_only])</code>	Equivalent to <i>shift</i> without copying data.
<code>slice_shift([periods, axis])</code>	Sort values and index labels by value.
<code>sort([axis, ascending, kind, na_position, ...])</code>	Sort object by labels (along an axis)
<code>sort_index([ascending])</code>	Sort Series with MultiIndex by chosen level.
<code>sortlevel([level, ascending, sort_remaining])</code>	squeeze length 1 dimensions
<code>squeeze()</code>	Return unbiased standard deviation over requested axis.
<code>std([axis, skipna, level, ddof])</code>	Binary operator <code>sub</code> with support to substitute a <code>fill_value</code> for missing data
<code>sub(other[, level, fill_value, axis])</code>	Binary operator <code>sub</code> with support to substitute a <code>fill_value</code> for missing data
<code>subtract(other[, level, fill_value, axis])</code>	Return the sum of the values for the requested axis
<code>sum([axis, skipna, level, numeric_only])</code>	Interchange axes and swap values axes appropriately
<code>swapaxes(axis1, axis2[, copy])</code>	Swap levels <i>i</i> and <i>j</i> in a MultiIndex
<code>swaplevel(i, j[, copy])</code>	Returns last <i>n</i> rows
<code>tail([n])</code>	return Series corresponding to requested indices
<code>take(indices[, axis, convert, is_copy])</code>	Attempt to write text representation of object to the system clipboard This can be used to copy data to the clipboard
<code>to_clipboard([excel, sep])</code>	Write Series to a comma-separated values (csv) file
<code>to_csv(path[, index, sep, na_rep, ...])</code>	Return dense representation of NDFrame (as opposed to sparse)
<code>to_dense()</code>	Convert Series to {label -> value} dict
<code>to_dict()</code>	Convert Series to DataFrame
<code>to_frame([name])</code>	activate the HDFStore
<code>to_hdf(path_or_buf, key, **kwargs)</code>	Convert the object to a JSON string.
<code>to_json([path_or_buf, orient, date_format, ...])</code>	msgpack (serialize) object to input file path
<code>to_msgpack([path_or_buf])</code>	Convert TimeSeries from DatetimeIndex to PeriodIndex with desired frequency
<code>to_period([freq, copy])</code>	Pickle (serialize) object to input file path
<code>to_pickle(path)</code>	



Table 3.4 – continued from previous page

<code>to_sparse([kind, fill_value])</code>	Convert Series to SparseSeries
<code>to_sql(name, con[, flavor, schema, ...])</code>	Write records stored in a DataFrame to a SQL database.
<code>to_string([buf, na_rep, float_format, ...])</code>	Render a string representation of the Series
<code>to_timestamp([freq, how, copy])</code>	Cast to DatetimeIndex of timestamps, at <i>beginning</i> of period
<code>tolist()</code>	Convert Series to a nested list
<code>transpose()</code>	return the transpose, which is by definition self
<code>truediv(other[, level, fill_value, axis])</code>	Binary operator truediv with support to substitute a fill_value for missing data
<code>truncate([before, after, axis, copy])</code>	Truncates a sorted NDFrame before and/or after some particular dates.
<code>tshift([periods, freq, axis])</code>	Shift the time index, using the index's frequency if available
<code>tz_convert(tz[, axis, level, copy])</code>	Convert the axis to target time zone.
<code>tz_localize(*args, **kwargs)</code>	Localize tz-naive TimeSeries to target time zone
<code>unique()</code>	Return array of unique values in the object.
<code>unstack([level])</code>	Unstack, a.k.a.
<code>update(other)</code>	Modify Series in place using non-NA values from passed Series.
<code>valid([inplace])</code>	
<code>value_counts([normalize, sort, ascending, ...])</code>	Returns object containing counts of unique values.
<code>var([axis, skipna, level, ddof])</code>	Return unbiased variance over requested axis.
<code>view([dtype])</code>	
<code>where(cond[, other, inplace, axis, level, ...])</code>	Return an object of same shape as self and whose corresponding entries are from
<code>xs(key[, axis, level, copy, drop_level])</code>	Returns a cross-section (row(s) or column(s)) from the Series/DataFrame.

**pp**

Property to access `sklearn.preprocessing`

**preprocessing = None**

**to\_frame** (*name=None*)

Convert Series to DataFrame

**Parameters** **name** : object, default None

The passed name should substitute for the series name (if it has one).

**Returns** **data\_frame** : DataFrame

## 3.5 Module contents





---

## expandas.skaccessors package

---

### 4.1 Subpackages

#### 4.1.1 expandas.skaccessors.test package

##### Submodules

expandas.skaccessors.test.test\_cluster module

expandas.skaccessors.test.test\_covariance module

expandas.skaccessors.test.test\_cross\_decomposition module

expandas.skaccessors.test.test\_cross\_validation module

expandas.skaccessors.test.test\_decomposition module

expandas.skaccessors.test.test\_dummy module

expandas.skaccessors.test.test\_ensemble module

expandas.skaccessors.test.test\_feature\_extraction module

expandas.skaccessors.test.test\_feature\_selection module

expandas.skaccessors.test.test\_gaussian\_process module

expandas.skaccessors.test.test\_grid\_search module

expandas.skaccessors.test.test\_isotonic module

expandas.skaccessors.test.test\_kernel\_approximation module

expandas.skaccessors.test.test\_lda module

expandas.skaccessors.test.test\_linear\_model module

expandas.skaccessors.test.test\_manifold module

expandas.skaccessors.test.test\_metrics module

expandas.skaccessors.test.test\_mixture module

expandas.skaccessors.test.test\_multiclass module

Accessor to `sklearn.cluster`.

#### Attributes

bicluster	
-----------	--

#### Methods

---

```

affinity_propagation(*args, **kwargs)
dbscan(*args, **kwargs)
k_means(n_clusters, *args, **kwargs)
mean_shift(*args, **kwargs)
spectral_clustering(*args, **kwargs)

```

---

**affinity\_propagation** (\*args, \*\*kwargs)

**bicluster** = None

**dbscan** (\*args, \*\*kwargs)

**k\_means** (n\_clusters, \*args, \*\*kwargs)

**mean\_shift** (\*args, \*\*kwargs)

**spectral\_clustering** (\*args, \*\*kwargs)

## 4.4 expandas.skaccessors.covariance module

**class** `expandas.skaccessors.covariance.CovarianceMethods` (*df*, *module\_name=None*, *attrs=None*)

Bases: `expandas.core.accessor.AccessorMethods`

Accessor to `sklearn.covariance`.

#### Methods

---

```

empirical_covariance(*args, **kwargs)
ledoit_wolf(*args, **kwargs)
oas(*args, **kwargs)

```

---

**empirical\_covariance** (\*args, \*\*kwargs)

**ledoit\_wolf** (\*args, \*\*kwargs)

**oas** (\*args, \*\*kwargs)

## 4.5 expandas.skaccessors.cross\_validation module

**class** expandas.skaccessors.cross\_validation.**CrossValidationMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.cross\_validation.

### Methods

---

```
StratifiedShuffleSplit(*args, **kwargs)
check_cv(cv, *args, **kwargs)
cross_val_score(estimator, *args, **kwargs)
iterate(cv)
permutation_test_score(estimator, *args, ...)
train_test_split(*args, **kwargs)
```

---

```
StratifiedShuffleSplit (*args, **kwargs)
check_cv (cv, *args, **kwargs)
cross_val_score (estimator, *args, **kwargs)
iterate (cv)
permutation_test_score (estimator, *args, **kwargs)
train_test_split (*args, **kwargs)
```

## 4.6 expandas.skaccessors.decomposition module

**class** expandas.skaccessors.decomposition.**DecompositionMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.decomposition.

### Methods

---

```
dict_learning(n_components, alpha, *args, ...)
dict_learning_online(*args, **kwargs)
fastica(*args, **kwargs)
sparse_encode(dictionary, *args, **kwargs)
```

---

```
dict_learning (n_components, alpha, *args, **kwargs)
dict_learning_online (*args, **kwargs)
fastica (*args, **kwargs)
sparse_encode (dictionary, *args, **kwargs)
```

## 4.7 expandas.skaccessors.ensemble module

**class** expandas.skaccessors.ensemble.**EnsembleMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.ensemble.

### Attributes

---

[`partial\_dependence`](#)

---

### `partial_dependence`

**class** expandas.skaccessors.ensemble.**PartialDependenceMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

### Methods

---

[`partial\_dependence\(gbrt, target\_variables, ...\)`](#)  
[`plot\_partial\_dependence\(gbrt, features, \*\*kwargs\)`](#)

---

**partial\_dependence** (*gbrt*, *target\_variables*, *\*\*kwargs*)

**plot\_partial\_dependence** (*gbrt*, *features*, *\*\*kwargs*)

## 4.8 expandas.skaccessors.feature\_extraction module

**class** expandas.skaccessors.feature\_extraction.**FeatureExtractionMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.feature\_extraction.

### Attributes

image	
text	

**image** = None

**text** = None

## 4.9 expandas.skaccessors.feature\_selection module

```
class expandas.skaccessors.feature_selection.FeatureSelectionMethods (df, module_name=None,
                                                                    attr=None,
                                                                    trs=None)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.feature_selection.
```

## 4.10 expandas.skaccessors.gaussian\_process module

```
class expandas.skaccessors.gaussian_process.GaussianProcessMethods (df, module_name=None,
                                                                    attr=None,
                                                                    attrs=None)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.gaussian_process.
```

### Attributes

---

```
correlation_models
regression_models
```

---

**correlation\_models**

**regression\_models**

```
class expandas.skaccessors.gaussian_process.RegressionModelsMethods (df, module_name=None,
                                                                    attr=None,
                                                                    trs=None)

Bases: expandas.core.accessor.AccessorMethods
```

## 4.11 expandas.skaccessors.grid\_search module

```
class expandas.skaccessors.grid_search.GridSearchMethods (df, module_name=None, attr=None,
                                                                    trs=None)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.grid_search.
```

### Methods

---

```
describe(estimator) Return cross validation results as pd.DataFrame.
```

---

**describe** (*estimator*)  
Return cross validation results as pd.DataFrame.



## 4.12 expandas.skaccessors.isotonic module

**class** expandas.skaccessors.isotonic.**IsotonicMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.isotonic.

### Attributes

---

IsotonicRegression

---

### Methods

---

check\_increasing(\*args, \*\*kwargs)  
isotonic\_regression(\*args, \*\*kwargs)

---

### IsotonicRegression

**check\_increasing** (\*args, \*\*kwargs)

**isotonic\_regression** (\*args, \*\*kwargs)

## 4.13 expandas.skaccessors.linear\_model module

**class** expandas.skaccessors.linear\_model.**LinearModelMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.linear\_model.

### Methods

---

lars\_path(\*args, \*\*kwargs)  
lasso\_path(\*args, \*\*kwargs)  
lasso\_stability\_path(\*args, \*\*kwargs)  
orthogonal\_mp\_gram(\*args, \*\*kwargs)

---

**lars\_path** (\*args, \*\*kwargs)

**lasso\_path** (\*args, \*\*kwargs)

**lasso\_stability\_path** (\*args, \*\*kwargs)

**orthogonal\_mp\_gram** (\*args, \*\*kwargs)

## 4.14 expandas.skaccessors.manifold module

**class** expandas.skaccessors.manifold.**ManifoldMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.manifold.

### Methods

---

[`locally\_linear\_embedding\(n\_neighbors, ...\)`](#)  
[`spectral\_embedding\(\*args, \*\*kwargs\)`](#)

---

**locally\_linear\_embedding** (*n\_neighbors*, *n\_components*, *\*args*, *\*\*kwargs*)

**spectral\_embedding** (*\*args*, *\*\*kwargs*)

## 4.15 expandas.skaccessors.metrics module

**class** expandas.skaccessors.metrics.**MetricsMethods** (*df*, *module\_name=None*, *attrs=None*)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.metrics.

### Attributes

---

[`pairwise`](#)

---

### Methods

---

`auc([kind, reorder])`  
`average_precision_score(*args, **kwargs)`  
`confusion_matrix(*args, **kwargs)`  
`consensus_score(*args, **kwargs)`  
`f1_score(*args, **kwargs)`  
`fbeta_score(beta, *args, **kwargs)`  
`hinge_loss(*args, **kwargs)`  
`log_loss(*args, **kwargs)`  
`precision_recall_curve(*args, **kwargs)`  
`precision_recall_fscore_support(*args, **kwargs)`  
`precision_score(*args, **kwargs)`  
`recall_score(*args, **kwargs)`  
`roc_auc_score(*args, **kwargs)`  
`roc_curve(*args, **kwargs)`  
`silhouette_samples(*args, **kwargs)`  
[`silhouette\_score\(\*args, \*\*kwargs\)`](#)

---

```

auc (kind='roc', reorder=False, **kwargs)
average_precision_score (*args, **kwargs)
confusion_matrix (*args, **kwargs)
consensus_score (*args, **kwargs)
f1_score (*args, **kwargs)
fbeta_score (beta, *args, **kwargs)
hinge_loss (*args, **kwargs)
log_loss (*args, **kwargs)
pairwise
precision_recall_curve (*args, **kwargs)
precision_recall_fscore_support (*args, **kwargs)
precision_score (*args, **kwargs)
recall_score (*args, **kwargs)
roc_auc_score (*args, **kwargs)
roc_curve (*args, **kwargs)
silhouette_samples (*args, **kwargs)
silhouette_score (*args, **kwargs)

```

## 4.16 expandas.skaccessors.multiclass module

**class** expandas.skaccessors.multiclass.**MultiClassMethods** (*df*, *module\_name*=None, *attrs*=None)

Bases: expandas.core.accessor.AccessorMethods

Accessor to sklearn.multiclass.

### Attributes

---

```

OneVsOneClassifier
OneVsRestClassifier
OutputCodeClassifier

```

---

### Methods

---

```

fit_ecoc(*args, **kwargs)
fit_ovo(*args, **kwargs)
fit_ovr(*args, **kwargs)
predict_ecoc(*args, **kwargs)
predict_ovo(*args, **kwargs)
predict_ovr(*args, **kwargs)

```

---

```
OneVsOneClassifier
OneVsRestClassifier
OutputCodeClassifier
fit_ecoc (*args, **kwargs)
fit_ovo (*args, **kwargs)
fit_ovr (*args, **kwargs)
predict_ecoc (*args, **kwargs)
predict_ovo (*args, **kwargs)
predict_ovr (*args, **kwargs)
```

## 4.17 expandas.skaccessors.neighbors module

```
class expandas.skaccessors.neighbors.NeighborsMethods (df, module_name=None,
                                                         attrs=None)
    Bases: expandas.core.accessor.AccessorMethods
    Accessor to sklearn.neighbors.
```

## 4.18 expandas.skaccessors.pipeline module

```
class expandas.skaccessors.pipeline.PipelineMethods (df, module_name=None,
                                                         attrs=None)
    Bases: expandas.core.accessor.AccessorMethods
    Accessor to sklearn.pipeline.
```

### Attributes

---

```
make_pipeline
make_union
```

---

```
make_pipeline
make_union
```

## 4.19 expandas.skaccessors.preprocessing module

```
class expandas.skaccessors.preprocessing.PreprocessingMethods (df,
                                                                module_name=None,
                                                                attrs=None)
    Bases: expandas.core.accessor.AccessorMethods
    Accessor to sklearn.preprocessing.
```

## Methods

---

`add_dummy_feature([value])`

---

`add_dummy_feature(value=1.0)`

## 4.20 expandas.skaccessors.svm module

**class** `expandas.skaccessors.svm.SVMMethods(df, module_name=None, attrs=None)`

Bases: `expandas.core.accessor.AccessorMethods`

Accessor to `sklearn.svm`.

## Attributes

---

`liblinear`  
`libsvm`  
`libsvm_sparse`

---

## Methods

---

`l1_min_c(*args, **kwargs)`

---

`l1_min_c(*args, **kwargs)`

`liblinear`

`libsvm`

`libsvm_sparse`

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