# sklearn.feature selection.SelectKBest

class sklearn.feature\_selection.SelectKBest(score\_func=<function f\_classif>, k=10)

[source]

Select features according to the k highest scores.

Read more in the User Guide.

#### **Parameters:**

### score\_func : callable

Function taking two arrays X and y, and returning a pair of arrays (scores, pvalues) or a single array with scores. Default is f\_classif (see below "See also"). The default function only works with classification tasks.

### k: int or "all", optional, default=10

Number of top features to select. The "all" option bypasses selection, for use in a parameter search.

### **Attributes:**

### scores\_: array-like of shape (n\_features,)

Scores of features.

### pvalues\_: array-like of shape (n\_features,)

p-values of feature scores, None if score\_func returned only scores.

#### See also:

#### f classif

ANOVA F-value between label/feature for classification tasks.

#### mutual info classif

Mutual information for a discrete target.

#### chi2

Chi-squared stats of non-negative features for classification tasks.

#### <u>f\_regressior</u>

F-value between label/feature for regression tasks.

### mutual info regression

Mutual information for a continuous target.

#### <u>SelectPercentile</u>

Select features based on percentile of the highest scores.

### <u>SelectFpr</u>

Select features based on a false positive rate test.

### SelectFdr

Select features based on an estimated false discovery rate.

### <u>SelectFwe</u>

Select features based on family-wise error rate.

#### <u>GenericUnivariateSelect</u>

Univariate feature selector with configurable mode.

### Notes

Ties between features with equal scores will be broken in an unspecified way.

### **Examples**

```
>>> from sklearn.datasets import load_digits
>>> from sklearn.feature_selection import SelectKBest, chi2
>>> X, y = load_digits(return_X_y=True)
>>> X.shape
(1797, 64)
>>> X_new = SelectKBest(chi2, k=20).fit_transform(X, y)
>>> X_new.shape
(1797, 20)
```

#### Methods

	fit(self X y)		Run score function on (X, y) and get the appropriate features.
Тс	oggle Menu	m(self, X[, y])	Fit to data, then transform it.

<pre>get params(self[, deep])</pre>	Get parameters for this estimator.
<pre>get support(self[, indices])</pre>	Get a mask, or integer index, of the features selected
<pre>inverse transform(self, X)</pre>	Reverse the transformation operation
<pre>set_params(self, \*\*params)</pre>	Set the parameters of this estimator.
<pre>transform(self, X)</pre>	Reduce X to the selected features.

<u>\_\_init\_\_(self, score\_func=<function f\_classif at 0x7f4d964b6320>, k=10)</u>

[source]

Initialize self. See help(type(self)) for accurate signature.

fit(self, X, y) [source]

Run score function on (X, y) and get the appropriate features.

#### **Parameters:**

### X: array-like of shape (n\_samples, n\_features)

The training input samples.

### y: array-like of shape (n\_samples,)

The target values (class labels in classification, real numbers in regression).

### **Returns:**

self : object

 ${\tt fit\_transform}(self, X, y = None, **fit\_params)$ 

[source]

Fit to data, then transform it.

Fits transformer to X and y with optional parameters fit\_params and returns a transformed version of X.

#### **Parameters:**

### X: numpy array of shape [n\_samples, n\_features]

Training set.

### y: numpy array of shape [n\_samples]

Target values.

### \*\*fit\_params: dict

Additional fit parameters.

### **Returns:**

### X\_new: numpy array of shape [n\_samples, n\_features\_new]

Transformed array.

get\_params(self, deep=True)

[source]

Get parameters for this estimator.

### **Parameters:**

#### deep: bool, default=True

If True, will return the parameters for this estimator and contained subobjects that are estimators.

### **Returns:**

### params: mapping of string to any

Parameter names mapped to their values.

get\_support(self, indices=False)

[source]

Get a mask, or integer index, of the features selected

### **Parameters:**

ా: boolean (default False)

Igle Menu , the return value will be an array of integers, rather than a boolean mask.

### Returns:

#### support: array

An index that selects the retained features from a feature vector. If indices is False, this is a boolean array of shape [# input features], in which an element is True iff its corresponding feature is selected for retention. If indices is True, this is an integer array of shape [# output features] whose values are indices into the input feature vector.

inverse\_transform(self, X)
[source]

Reverse the transformation operation

#### **Parameters:**

#### X : array of shape [n\_samples, n\_selected\_features]

The input samples.

#### **Returns:**

### X\_r: array of shape [n\_samples, n\_original\_features]

x with columns of zeros inserted where features would have been removed by transform.

set\_params(self, \*\*params) [source]

Set the parameters of this estimator.

The method works on simple estimators as well as on nested objects (such as pipelines). The latter have parameters of the form <component>\_\_<parameter> so that it's possible to update each component of a nested object.

#### Parameters:

#### \*\*params: dict

Estimator parameters.

#### **Returns:**

### self : object

Estimator instance.

 ${\sf transform}(\textit{self}, \textit{X}) \hspace*{2cm} [\mathsf{source}]$ 

Reduce X to the selected features.

### **Parameters:**

## X : array of shape [n\_samples, n\_features]

The input samples.

#### Returns:

# X\_r: array of shape [n\_samples, n\_selected\_features]

The input samples with only the selected features.

# Examples using sklearn.feature\_selection.SelectKBest



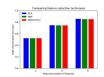
Pipeline Anova SVM



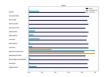
Univariate Feature
Selection



Concatenating multiple feature extraction methods



Selecting dimensionality reduction with Pipeline and GridSearchCV



Classification of text documents using sparse features