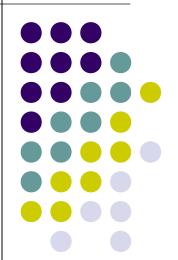
Pattern Recognition Introduction to ScikitLearn

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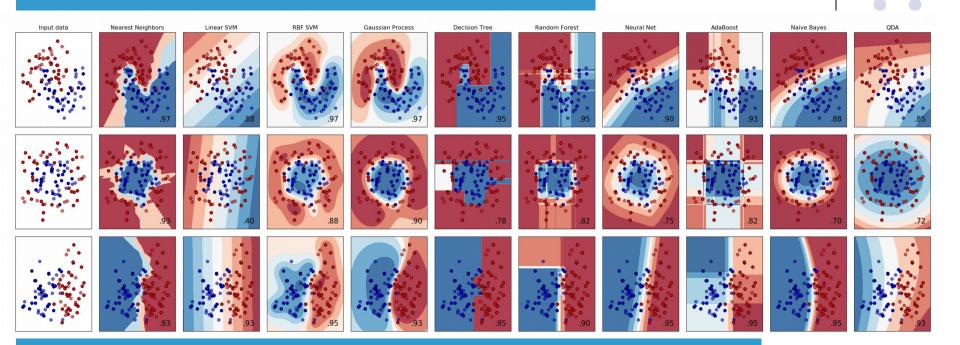




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scikit-learn

Machine Learning in Python



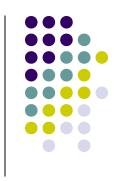
- Simple and efficient tools for data mining and data analysis
- Accessible to everybody, and reusable in various contexts
- Built on NumPy, SciPy, and matplotlib
- Open source, commercially usable BSD license



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- C-support vector classification
- Implementation based on Symlib

sklearn.svm.SVC

class sklearn.svm. **SVC** (C=1.0, kernel='rbf', degree=3, gamma='auto', coef0=0.0, shrinking=True, probability=False, tol=0.001, cache_size=200, class_weight=None, verbose=False, max_iter=-1, decision_function_shape='ovr', random_state=None)



Parameters: C: float, optional (default=1.0)

Penalty parameter C of the error term.

kernel: string, optional (default='rbf')

Specifies the kernel type to be used in the algorithm. It must be one of 'linear', 'poly', 'rbf', 'sigmoid', 'precomputed' or a callable. If none is given, 'rbf' will be used. If a callable is given it is used to pre-compute the kernel matrix from data matrices; that matrix should be an array of shape (n_samples, n_samples).

degree : int, optional (default=3)

Degree of the polynomial kernel function ('poly'). Ignored by all other kernels.

gamma: float, optional (default='auto')

Kernel coefficient for 'rbf', 'poly' and 'sigmoid'. If gamma is 'auto' then 1/n_features will be used instead.

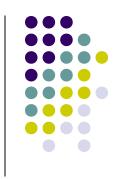
coef0 : float, optional (default=0.0)

Independent term in kernel function. It is only significant in 'poly' and 'sigmoid'.

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- linear: $\langle x, x' \rangle$.
- polynomial: $(\gamma\langle x,x'\rangle+r)^d$. d is specified by keyword degree , r by coef0 .
- rbf: $\exp(-\gamma||x-x'||^2)$. γ is specified by keyword gamma, must be greater than 0.
- sigmoid $(\tanh(\gamma\langle x,x'\rangle+r))$, where r is specified by coef0.



fit (X, y, sample weight=None)

[source]

Fit the SVM model according to the given training data.

Parameters: X: {array-like, sparse matrix}, shape (n_samples, n_features)

Training vectors, where n samples is the number of samples and n features is the number of features. For kernel="precomputed", the expected shape of X is (n samples, n samples).

y: array-like, shape (n samples,)

Target values (class labels in classification, real numbers in regression)

sample_weight : array-like, shape (n_samples,)

Per-sample weights. Rescale C per sample. Higher weights force the classifier to put more emphasis on these points.

Returns:

self: object

Returns self.

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predict (X)

Perform classification on samples in X.

For an one-class model, +1 or -1 is returned.

Parameters: X : {array-like, sparse matrix}, shape (n_samples, n_features)

For kernel="precomputed", the expected shape of X is [n_samples_test, n_sam-

ples_train]

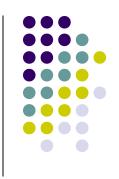
Returns: y_pred : array, shape (n_samples,)

Class labels for samples in X.



sklearn.svm.NuSVC

class sklearn.svm. **NuSVC** (nu=0.5, kernel='rbf', degree=3, gamma='auto', coef0=0.0, shrinking=True, probability=False, tol=0.001, cache_size=200, class_weight=None, verbose=False, max_iter=-1, decision_function_shape='ovr', random_state=None)



Nu-Support Vector Classification.

Similar to SVC but uses a parameter to control the number of support vectors.

The implementation is based on libsvm.

Read more in the User Guide.

Parameters: nu : float, optional (default=0.5)

An upper bound on the fraction of training errors and a lower bound of the fraction of support vectors. Should be in the interval (0, 1].

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