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Examples

sklearn.svm.SVC

class sklearn.svm.SVC(C=1.0, kernel='rbf', degree=3, gamma='auto_deprecated', 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) [source]

C-Support Vector Classification.

The implementation is based on libsvm. The fit time scales at least quadratically with the number of samples and may be impractical beyond tens of thousands of samples. For large datasets consider using sklearn.linear_model.LinearSVC or sklearn.linear_model.SGDClassifier instead, possibly after a sklearn.kernel_approximation.Nystroem transformer.

The multiclass support is handled according to a one-vs-one scheme.

For details on the precise mathematical formulation of the provided kernel functions and how gamma, coef0 and degree affect each other, see the corresponding section in the narrative documentation: Kernel functions.

Read more in the User Guide.

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).

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'.

Current default is 'auto' which uses 1 / n_features, if gamma='scale' is passed then it uses 1 / (n_features * X.var()) as value of gamma. The current default of gamma, 'auto', will change to 'scale' in version 0.22. 'auto_deprecated', a deprecated version of 'auto' is used as a default indicating that no explicit value of gamma was passed.

coef0 : float, optional (default=0.0)

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

shrinking : boolean, optional (default=True)

Whether to use the shrinking heuristic.

probability : boolean, optional (default=False)

Whether to enable probability estimates. This must be enabled prior to calling fit, and will slow down that method.

tol: float, optional (default=1e-3)

Tolerance for stopping criterion.

cache_size: float, optional

Specify the size of the kernel cache (in MB).

class_weight : {dict, 'balanced'}, optional

Set the parameter C of class i to class_weight[i]*C for SVC. If not given, all classes are supposed to have weight one. The "balanced" mode uses the values of y to automatically adjust weights inversely proportional to class frequencies in the input data as n_samples / (n_classes * np.bincount(y))

verbose: bool, default: False

Enable verbose output. Note that this setting takes advantage of a per-process runtime setting in libsvm that, if enabled, may not work properly in a multithreaded context.

max iter: int, optional (default=-1)

Hard limit on iterations within solver, or -1 for no limit.

decision function shape: 'ovo', 'ovr', default='ovr'

Whether to return a one-vs-rest ('ovr') decision function of shape (n_samples, n_classes) as all other classifiers, or the original one-vs-one ('ovo') decision function of libsvm which has shape (n_samples, n_classes * (n_classes - 1) / 2). However, one-vs-one ('ovo') is always used as multi-class strategy.

Changed in version 0.19: decision function shape is 'ovr' by default.

New in version 0.17: decision function shape='ovr' is recommended.

Changed in version 0.17: Deprecated decision function shape='ovo' and None.

random_state : int, RandomState instance or None, optional (default=None)

The seed of the pseudo random number generator used when shuffling the data for probability estimates. If int, random_state is the seed used by the random number generator; If RandomState instance, random_state is the random number generator; If None, the random number generator is the RandomState instance used by np.random.

Attributes: support : array-like, shape = [n SV]

Indices of support vectors.

>>

support_vectors_: array-like, shape = [n_SV, n_features]
Support vectors.

n_support_: array-like, dtype=int32, shape = [n_class] Number of support vectors for each class.

dual_coef_ : array, shape = [n_class-1, n_SV]

Coefficients of the support vector in the decision function. For multiclass, coefficient for all 1-vs-1 classifiers. The layout of the coefficients in the multiclass case is somewhat non-trivial. See the section about multi-class classification in the SVM section of the User Guide for details.

coef_ : array, shape = [n_class * (n_class-1) / 2, n_features]

Weights assigned to the features (coefficients in the primal problem). This is only available in the case of a linear kernel.

coef is a readonly property derived from dual coef and support vectors .

intercept_ : *array, shape* = [*n_class* * (*n_class-1*) / 2]

Constants in decision function.

fit_status_: int

0 if correctly fitted, 1 otherwise (will raise warning)

probA_ : array, shape = [n_class * (n_class-1) / 2]

probB_: array, shape = [n_class * (n_class-1) / 2]

If probability=True, the parameters learned in Platt scaling to produce probability estimates from decision values. If probability=False, an empty array. Platt scaling uses the logistic function 1 / (1 + exp(decision_value * probA_ + probB_)) where probA_ and probB_ are learned from the dataset [R20c70293ef72-2]. For more information on the multiclass case and training procedure see section 8 of [R20c70293ef72-1].

See also:

SVR

Support Vector Machine for Regression implemented using libsvm.

LinearSVC

Scalable Linear Support Vector Machine for classification implemented using liblinear. Check the See also section of LinearSVC for more comparison element.

References

[R20c70293ef72-1] LIBSVM: A Library for Support Vector Machines

[R20c70293ef72- Platt, John (1999). "Probabilistic outputs for support vector machines and comparison to regularizedlikelihood methods."

Examples

```
>>> import numpy as np
>>> X = np.array([[-1, -1], [-2, -1], [1, 1], [2, 1]])
>>> y = np.array([1, 1, 2, 2])
>>> from sklearn.svm import SVC
>>> clf = SVC(gamma='auto')
>>> clf.fit(X, y)
SVC(C=1.0, cache_size=200, class_weight=None, coef0=0.0,
    decision_function_shape='ovr', degree=3, gamma='auto', kernel='rbf',
    max_iter=-1, probability=False, random_state=None, shrinking=True,
    tol=0.001, verbose=False)
>>> print(clf.predict([[-0.8, -1]]))
[1]
```

Methods

${\tt decision_function}({\tt Self},{\sf X})$	Evaluates the decision function for the samples in X.
<pre>fit(self, X, y[, sample_weight])</pre>	Fit the SVM model according to the given training data.
<pre>get_params(self[, deep])</pre>	Get parameters for this estimator.
<pre>predict(self, X)</pre>	Perform classification on samples in X.
<pre>score(self, X, y[, sample_weight])</pre>	Returns the mean accuracy on the given test data and labels.
<pre>set_params(self, **params)</pre>	Set the parameters of this estimator.

__init__(self, C=1.0, kernel='rbf', degree=3, gamma='auto_deprecated', 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) [source]

```
\label{eq:cource} \begin{split} \mathsf{decision\_function}(\mathit{self}, X) \end{split} \qquad \qquad [\mathsf{source}] \end{split}
```

Evaluates the decision function for the samples in X.

Parameters: X : array-like, shape (n_samples, n_features)

Returns: X: array-like, shape (n_samples, n_classes * (n_classes-1) / 2)

Returns the decision function of the sample for each class in the model. If decision_function_shape='ovr', the shape is (n_samples, n_classes).

Notes

If decision_function_shape='ovo', the function values are proportional to the distance of the samples X to the separating hyperplane. If the exact distances are required, divide the function values by the norm of the weight vector (coef_). See also this question for further details. If decision_function_shape='ovr', the decision function is a monotonic transformation of ovo decision function.

```
fit(self, 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

Notes

If X and y are not C-ordered and contiguous arrays of np.float64 and X is not a scipy.sparse.csr_matrix, X and/or y may be copied.

If X is a dense array, then the other methods will not support sparse matrices as input.

get_params(self, deep=True)

[source]

Get parameters for this estimator.

Parameters: deep: boolean, optional

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.

predict(self, X)

[source]

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_samples_train]

Returns: y pred : array, shape (n samples,)

Class labels for samples in X.

predict_log_proba

Compute log probabilities of possible outcomes for samples in X.

The model need to have probability information computed at training time: fit with attribute probability set to True.

Parameters: X: array-like, shape (n samples, n features)

For kernel="precomputed", the expected shape of X is [n_samples_test, n_samples_train]

Returns:

T: array-like, shape (n samples, n classes)

Returns the log-probabilities of the sample for each class in the model. The columns correspond to the classes in sorted order, as they appear in the attribute classes.

Notes

The probability model is created using cross validation, so the results can be slightly different than those obtained by predict. Also, it will produce meaningless results on very small datasets.

predict_proba

Compute probabilities of possible outcomes for samples in X.

The model need to have probability information computed at training time: fit with attribute probability set to True.

Parameters: X : array-like, shape (n samples, n features)

For kernel="precomputed", the expected shape of X is [n_samples_test, n_samples_train]

Returns: T: array-like, shape (n_samples, n_classes)

Returns the probability of the sample for each class in the model. The columns correspond to the classes in sorted order, as they appear in the attribute classes .

Notes

The probability model is created using cross validation, so the results can be slightly different than those obtained by predict. Also, it will produce meaningless results on very small datasets.

score(self, X, y, sample weight=None)

[source]

Returns the mean accuracy on the given test data and labels.

In multi-label classification, this is the subset accuracy which is a harsh metric since you require for each sample that each label set be correctly predicted.

Parameters: X : array-like, shape = (n_samples, n_features)

Test samples.

y : array-like, shape = (n_samples) or (n_samples, n_outputs)

True labels for X.

sample_weight : array-like, shape = [n_samples], optional
Sample weights.

Returns: score : float

Mean accuracy of self.predict(X) wrt. y.

set_params(self, **params)

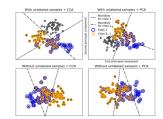
[source]

Set the parameters of this estimator.

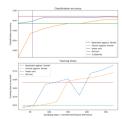
The method works on simple estimators as well as on nested objects (such as pipelines). The latter have parameters of the form <component>__<parameter> so that it's possible to update each component of a nested object.

Returns: self

Examples using sklearn.svm.SVC



Multilabel classification

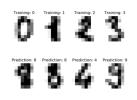


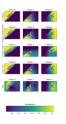
Explicit feature map approximation for RBF kernels



Faces recognition example using eigenfaces and SVMs







Libsvm GUI

Recognizing hand-written digits

Plot classification probability





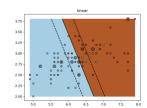


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Classifier comparison

Concatenating multiple feature extraction methods

Plot the decision boundaries of a VotingClassifier

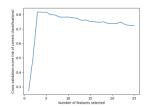


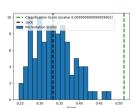


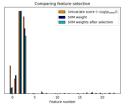
Cross-validation on Digits
Dataset Exercise

SVM Exercise

Recursive feature elimination



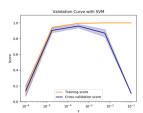




Recursive feature elimination with crossvalidation

Test with permutations the significance of a classification score

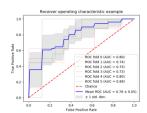
Univariate Feature Selection



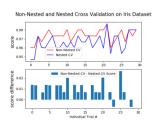
Plotting Validation Curves



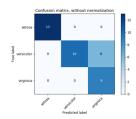
Parameter estimation using grid search with cross-validation



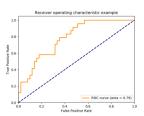
Receiver Operating
Characteristic (ROC) with
cross validation



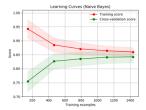
Nested versus non-nested cross-validation



Confusion matrix



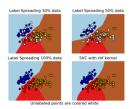
Receiver Operating Characteristic (ROC)



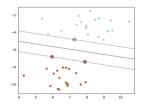
Plotting Learning Curves



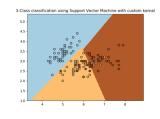
Feature discretization



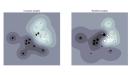
Decision boundary of label propagation versus SVM on the Iris dataset



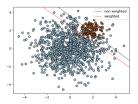
SVM: Maximum margin separating hyperplane



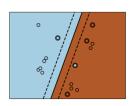
SVM with custom kernel



SVM: Weighted samples



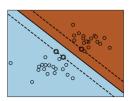
SVM: Separatinghyperplane for unbalanced classes



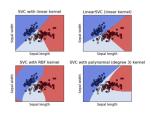
SVM-Kernels



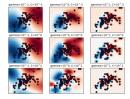
SVM-Anova: SVM with univariate feature selection



SVM Margins Example



Plot different SVM classifiers in the iris dataset



RBF SVM parameters