

>>

Examples

sklearn.metrics.silhouette_score

sklearn.metrics.silhouette_score(X, labels, metric='euclidean',
sample_size=None, random_state=None, **kwds) [source]

Compute the mean Silhouette Coefficient of all samples.

The Silhouette Coefficient is calculated using the mean intra-cluster distance (a) and the mean nearest-cluster distance (b) for each sample. The Silhouette Coefficient for a sample is $(b - a) / \max(a, b)$. To clarify, b is the distance between a sample and the nearest cluster that the sample is not a part of. Note that Silhouette Coefficient is only defined if number of labels is $2 \le n$ _labels n_samples - 1.

This function returns the mean Silhouette Coefficient over all samples. To obtain the values for each sample, use **silhouette samples**.

The best value is 1 and the worst value is -1. Values near 0 indicate overlapping clusters. Negative values generally indicate that a sample has been assigned to the wrong cluster, as a different cluster is more similar.

Read more in the User Guide.

Parameters:

X : array [n_samples_a, n_samples_a] if metric == "precomputed", or, [n_samples_a, n_features] otherwise

Array of pairwise distances between samples, or a feature array.

labels : array, shape = [n_samples]

Predicted labels for each sample.

metric : string, or callable

The metric to use when calculating distance between instances in a feature array. If metric is a string, it must be one of the options allowed by

metrics.pairwise.pairwise_distances. If X is the distance array itself, use metric="precomputed".

sample size: int or None

The size of the sample to use when computing the Silhouette Coefficient on a random subset of the data. If sample size is None, no sampling is used.

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

The generator used to randomly select a subset of samples. 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. Used when sample size is not None.

**kwds : optional keyword parameters

Any further parameters are passed directly to the distance function. If using a scipy.spatial.distance metric, the parameters are still metric dependent. See the scipy docs for usage examples.

Returns: silk

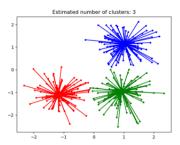
silhouette: float

Mean Silhouette Coefficient for all samples.

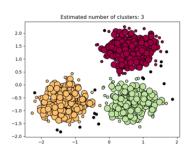
References

- [1] Peter J. Rousseeuw (1987). "Silhouettes: a Graphical Aid to the Interpretation and Validation of Cluster Analysis". Computational and Applied Mathematics 20: 53-65.
- [2] Wikipedia entry on the Silhouette Coefficient

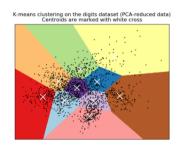
Examples using sklearn.metrics.silhouette score



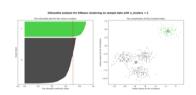
Demo of affinity propagation clustering algorithm



Demo of DBSCAN clustering algorithm



A demo of K-Means clustering on the handwritten digits data



Selecting the number of clusters with silhouette analysis on KMeans clustering



Clustering text documents using k-means

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