

# binspec

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binary_peaks	<i>Find binary peaks</i>
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## Description

Find peaks in window of size  $2 \times \text{neighbors} + 1$  and label m/z integers within the error as peaks. Returns vector of peak m/z integers.

## Usage

```
binary_peaks(df, neighbors, error = 0)
```

## Arguments

df	Data frame of m/z and intensities
neighbors	Number of neighboring m/z values to compare on right and left
error	m/z Decimal error value

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classifier_accuracies	<i>Classifier Accuracies</i>
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## Description

Find the best classifier using leave-one-out cross validation (svm) and out-of-bag error (random forests). Returns a list of classifier results

## Usage

```
classifier_accuracies(peaks, labels, min_peak_percentage)
```

## Arguments

peaks	Boolean matrix of mass spectra rows with m/z columns, indicating if an m/z value corresponds to a peak.
labels	The correct classifications of the peaks.
minpeaks	How many "true" values must show up for a given m/z value for it to be considered a feature.

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combine_peaks	<i>Combine peak vectors</i>
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### Description

Create a binary matrix, each column represents an m/z value, and each row represents a mass spectra. The value indicates whether or not the m/z of this spectra is a peak.

### Usage

```
combine_peaks(list_mz_peaks)
```

### Arguments

list_mz_peaks	List of m/z peak vectors
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naive_feature_importance	<i>Rank importance of features</i>
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### Description

Given a matrix of binary peaks and each row's corresponding labels, this function takes returns the absolute difference between the proportion of times an m/z value was labeled as a peak within each of the two classes.

### Usage

```
naive_feature_importance(peaks, labels)
```

### Arguments

peaks	A matrix of peaks
labels	A factor vector of labels whose length is equal to the number of rows of peaks

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round_df	<i>Round data frame</i>
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### Description

Round all m/z and intensity values to integers.

### Usage

```
round_df(df)
```

### Arguments

df	Data frame
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svm\_rf*SVM and RF Accuracies*

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**Description**

Given a vector of neighbor values and a vector of the minimum number of peaks to be considered, this function finds the peak mz values for a data set by running `binary_peaks` using each of the neighbor vector values, runs SVM and RF on the peaks for each of the `min_peak_count` values, and returns the accuracies of each test in a table. The table's rows are the number of neighbors, and the columns are the `min_peak_count` values.

**Usage**

```
svm_rf(list_of_dfs, labels, neighbors, min_peaks_percentage)
```

**Arguments**

<code>list_of_dfs</code>	The first data frame of mz values and frequencies
<code>labels</code>	The labels of the two states the first data frame's values could be classified as
<code>neighbors</code>	A vector of the number of neighbors to be considered in the <code>binary_peaks</code> function
<code>min_peaks_count</code>	A vector of the minimum numbers of peaks to be considered in the <code>classifier_accuracies</code> function

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