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scipy.cluster.hierarchy.dendrogram¶

scipy.cluster.hierarchy.dendrogram(*Z*, *p*=30, truncate_mode=None, color_threshold=None, get_leaves=True, orientation='top', labels=None, count_sort=False, distance_sort=False, show_leaf_counts=True, no_plot=False, no_labels=False, color_list=None, leaf_font_size=None, leaf_rotation=None, leaf_label_func=None, no_leaves=False, show_contracted=False, link_color_func=None, ax=None) [source]

(http://github.com/scipy/scipy/blob/v0.14.0/scipy/cluster/hierarchy.py#L1894)

Plots the hierarchical clustering as a dendrogram.

The dendrogram illustrates how each cluster is composed by drawing a U-shaped link between a non-singleton cluster and its children. The height of the top of the U-link is the distance between its children clusters. It is also the cophenetic distance between original observations in the two children clusters. It is expected that the distances in Z[:,2] be monotonic, otherwise crossings appear in the dendrogram.

Parameters: Z:ndarray

The linkage matrix encoding the hierarchical clustering to render as a dendrogram. See the linkage function for more information on the format of z.

p: int, optional

The p parameter for truncate mode.

truncate_mode : str, optional

The dendrogram can be hard to read when the original observation matrix from which the linkage is derived is large. Truncation is used to condense the dendrogram. There are several modes:

None/'none'

No truncation is performed (Default).

'lastp'

The last p non-singleton formed in the linkage are the only non-leaf nodes in the linkage; they correspond to rows Z[n-p-2:end] in Z. All other non-singleton clusters are contracted into leaf nodes.

'mlab'

This corresponds to MATLAB(TM) behavior. (not implemented yet)

'level'/'mtica'

No more than p levels of the dendrogram tree are displayed. This corresponds to Mathematica(TM) behavior.

color_threshold : double, optional

For brevity, let t be the color_threshold. Colors all the descendent links below a cluster node k the same color if k is the first node below the cut threshold t. All links connecting nodes with distances greater than or equal to the threshold are colored blue. If t is less than or equal to zero, all nodes are colored blue. If color_threshold is None or 'default', corresponding with MATLAB(TM) behavior, the threshold is set to

0.7*max(Z[:,2]).

get_leaves : bool, optional

Includes a list R['leaves']=H in the result dictionary. For each i, H[i] == j, cluster node j appears in position i in the left-to-right traversal of the leaves, where j < 2n-1 and i < n.

orientation: str, optional

The direction to plot the dendrogram, which can be any of the following strings:

'top'

Plots the root at the top, and plot descendent links going downwards. (default).

'bottom'

Plots the root at the bottom, and plot descendent links going upwards.

'left'

Plots the root at the left, and plot descendent links going right.

'right'

Plots the root at the right, and plot descendent links going left.

labels: ndarray, optional

By default labels is None so the index of the original observation is used to label the leaf nodes. Otherwise, this is an n-sized list (or tuple). The labels[i] value is the text to put under the i th leaf node only if it corresponds to an original observation and not a non-singleton cluster.

count_sort : str or bool, optional

For each node n, the order (visually, from left-to-right) n's two descendent links are plotted is determined by this parameter, which can be any of the following values:

False

Nothing is done.

'ascending' Or True

The child with the minimum number of original objects in its cluster is plotted first.

'descendent'

The child with the maximum number of original objects in its cluster is plotted first.

Note distance_sort and count_sort cannot both be True.

distance_sort : str or bool, optional

For each node n, the order (visually, from left-to-right) n's two descendent links are plotted is determined by this parameter, which can be any of the following values:

False

Nothing is done.

'ascending' Or True

The child with the minimum distance between its direct descendents is plotted first.

'descending'

The child with the maximum distance between its direct descendents is plotted first.

Note distance_sort and count_sort cannot both be True.

show_leaf_counts : bool, optional

When True, leaf nodes representing k>1 original observation are labeled with the number of observations they contain in parentheses.

no_plot : bool, optional

When True, the final rendering is not performed. This is useful if only the data structures computed for the rendering are needed or if matplotlib is not available.

no_labels : bool, optional

When True, no labels appear next to the leaf nodes in the rendering of the dendrogram.

leaf_label_rotation : double, optional

Specifies the angle (in degrees) to rotate the leaf labels. When unspecified, the rotation based on the number of nodes in the dendrogram (default is 0).

leaf font size: int, optional

Specifies the font size (in points) of the leaf labels. When unspecified, the size based on the number of nodes in the dendrogram.

leaf_label_func : lambda or function, optional

When leaf_label_func is a callable function, for each leaf with cluster index k < 2n-1. The function is expected to return a string with the label for the leaf.

Indices k < n correspond to original observations while indices k > n correspond to non-singleton clusters.

For example, to label singletons with their node id and nonsingletons with their id, count, and inconsistency coefficient, simply do:

```
>>> # First define the leaf label function.
>>> def llf(id):
...     if id < n:
...         return str(id)
...     else:
>>>         return '[%d %d %1.2f]' % (id, count, R
[n-id,3])
>>>
>>> # The text for the leaf nodes is going to be bi
g so force
>>> # a rotation of 90 degrees.
>>> dendrogram(Z, leaf_label_func=llf, leaf_rotatio
n=90)
```

show_contracted: bool, optional

When True the heights of non-singleton nodes contracted into a leaf node are plotted as crosses along the link connecting that leaf node. This really is only useful when truncation is used (see truncate mode parameter).

link_color_func : callable, optional

If given, *link_color_function* is called with each non-singleton id corresponding to each U-shaped link it will paint. The function is expected to return the color to paint the link, encoded as a matplotlib color string code. For example:

```
>>> dendrogram(Z, link_color_func=lambda k: colors[k])
```

colors the direct links below each untruncated non-singleton node k using colors[k].

ax: matplotlib Axes instance, optional

If None and *no_plot* is not True, the dendrogram will be plotted on the current axes. Otherwise if *no_plot* is not True the dendrogram will be plotted on the given Axes instance. This can be useful if the dendrogram is part of a more complex figure.

Returns:

R: dict

A dictionary of data structures computed to render the dendrogram. Its has the following keys:

'icoords'

A list of lists [I1, I2, ..., Ip] where Ik is a list of 4 independent variable coordinates corresponding to the line that represents the k'th link painted.

'dcoords'

A list of lists [12, 12, ..., Ip] where Ik is a list of 4 independent variable coordinates corresponding to the line that represents the k'th link painted.

'ivl'

A list of labels corresponding to the leaf nodes.

'leaves'

For each i, $\mathtt{H[i]} == \mathtt{j}$, cluster node \mathtt{j} appears in position \mathtt{i} in the left-to-right traversal of the leaves, where j < 2n-1 and i < n. If \mathtt{j} is less than \mathtt{n} , the \mathtt{i} -th leaf node corresponds to an original observation. Otherwise, it corresponds to a non-singleton cluster.

Previous topic

scipy.cluster.hierarchy.to_mlab_linkage (scipy.cluster.hierarchy.to_mlab_linkage.html)

Next topic

scipy.cluster.hierarchy.ClusterNode (scipy.cluster.hierarchy.ClusterNode.html)