Visualizing hierarchies

UNSUPERVISED LEARNING IN PYTHON



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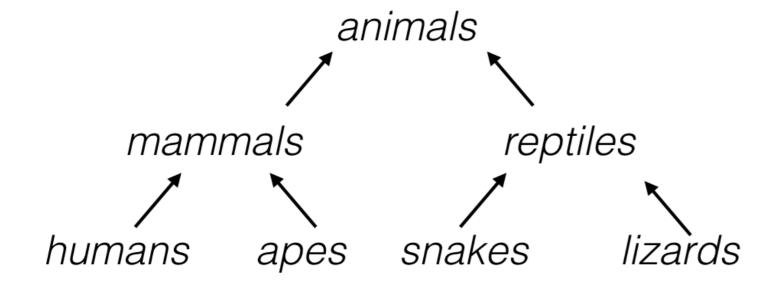


Visualizations communicate insight

- "t-SNE": Creates a 2D map of a dataset (later)
- "Hierarchical clustering" (this video)

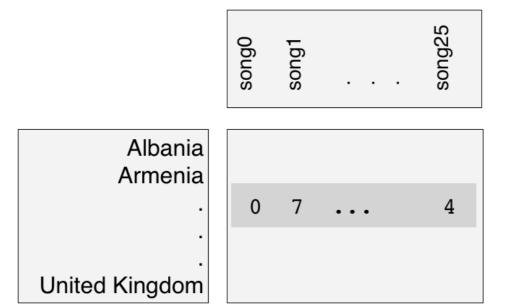
A hierarchy of groups

- Groups of living things can form a hierarchy
- Clusters are contained in one another



Eurovision scoring dataset

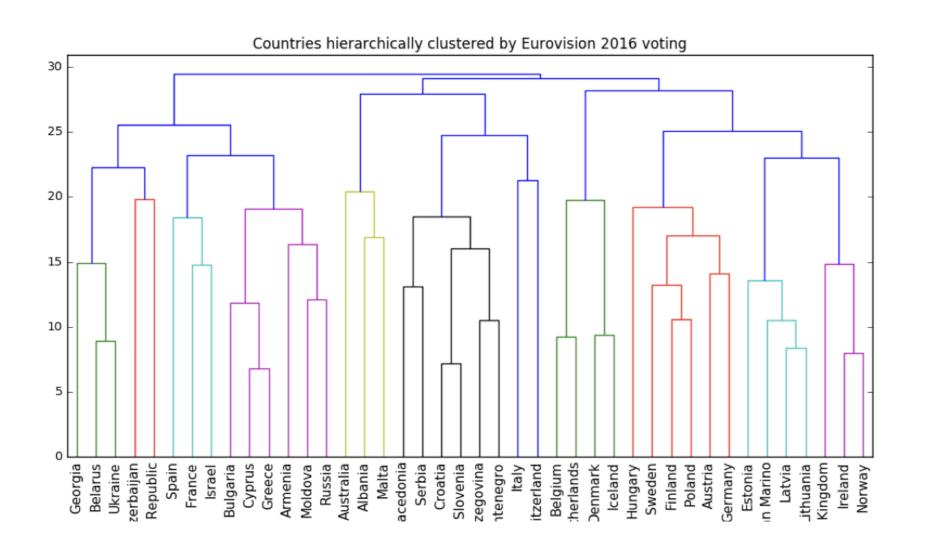
- Countries gave scores to songs performed at the Eurovision 2016
- 2D array of scores
- Rows are countries, columns are songs



¹ https://www.eurovision.tv/page/results



Hierarchical clustering of voting countries





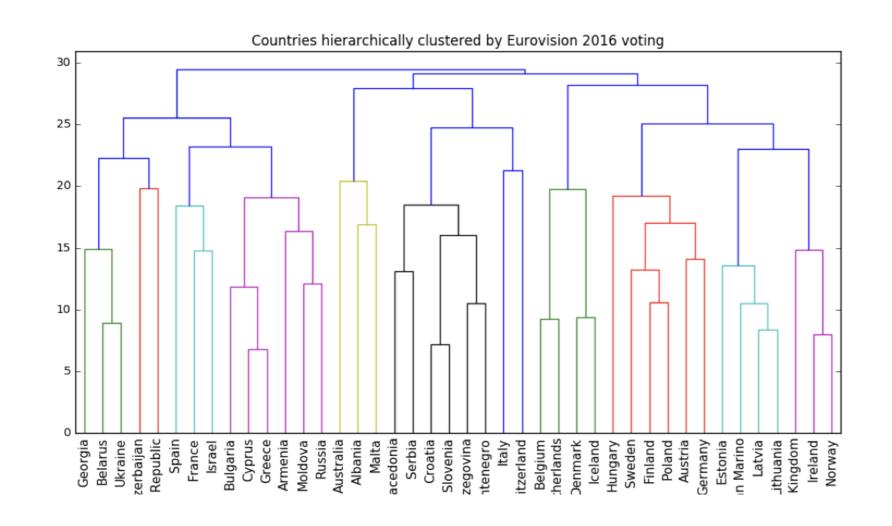
Hierarchical clustering

- Every country begins in a separate cluster
- At each step, the two closest clusters are merged
- Continue until all countries in a single cluster
- This is "agglomerative" hierarchical clustering



The dendrogram of a hierarchical clustering

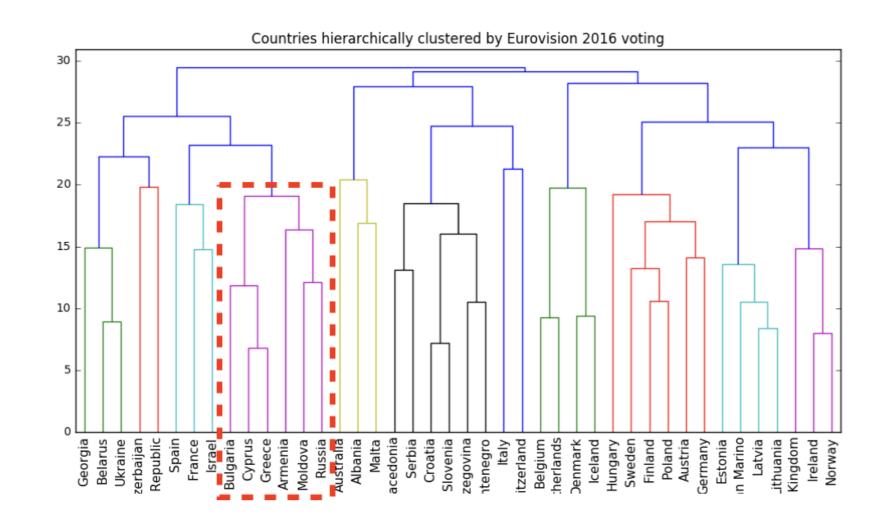
- Read from the bottom up
- Vertical lines represent clusters



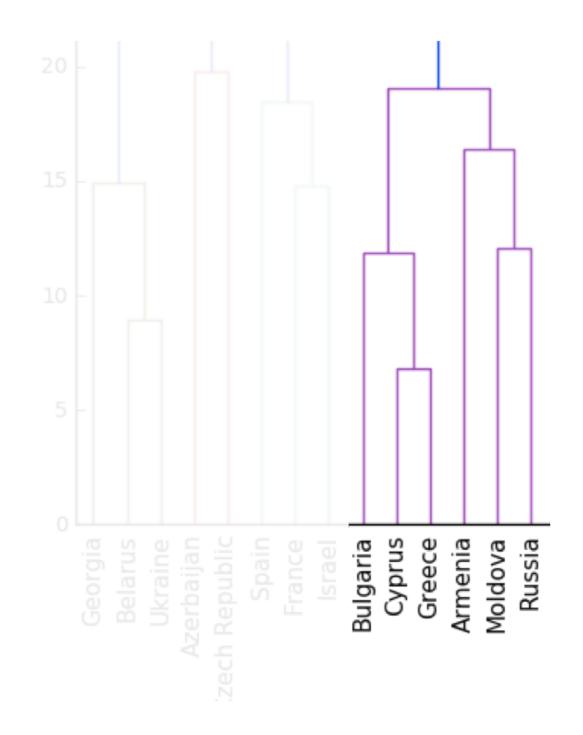


The dendrogram of a hierarchical clustering

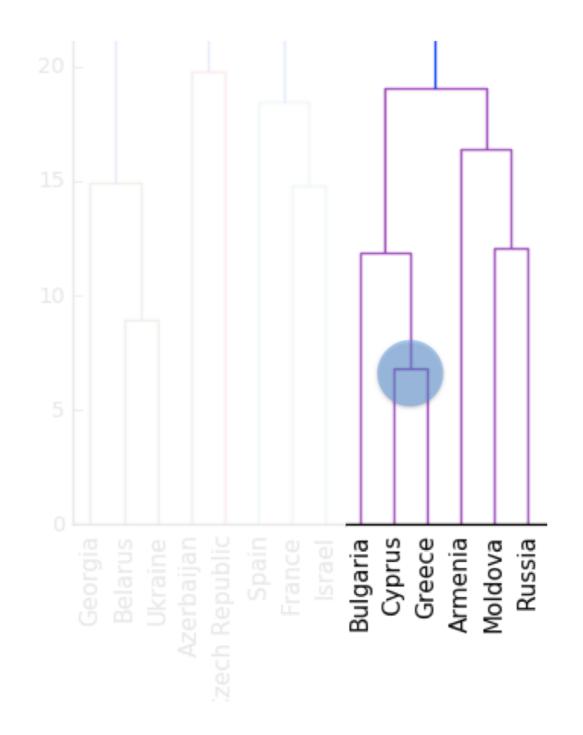
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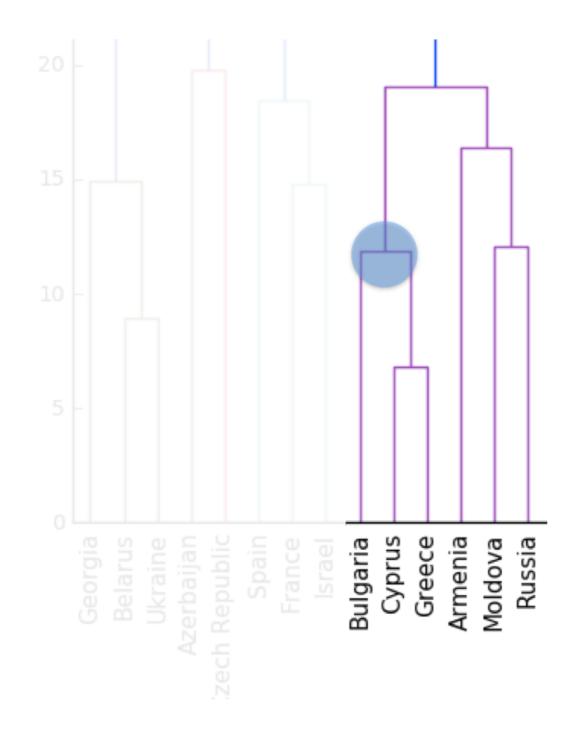




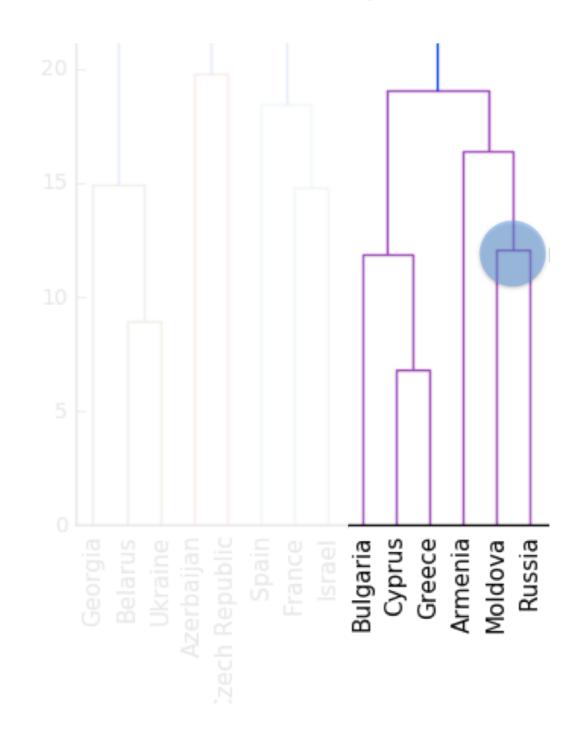




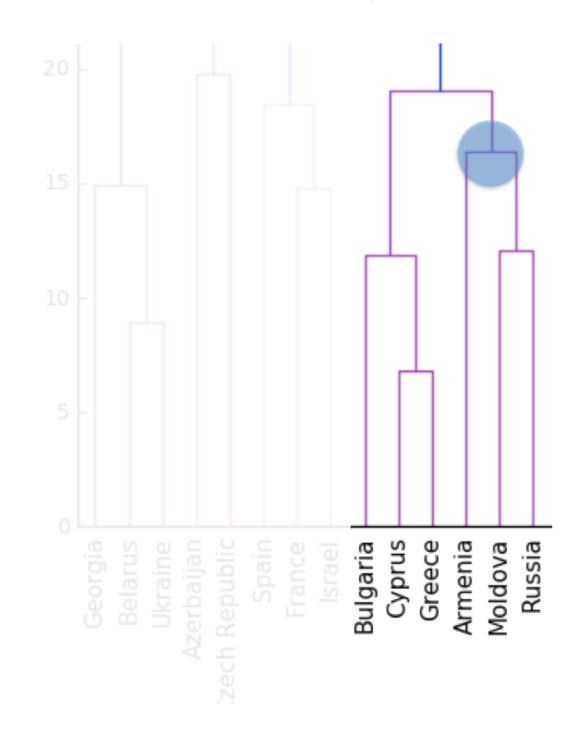




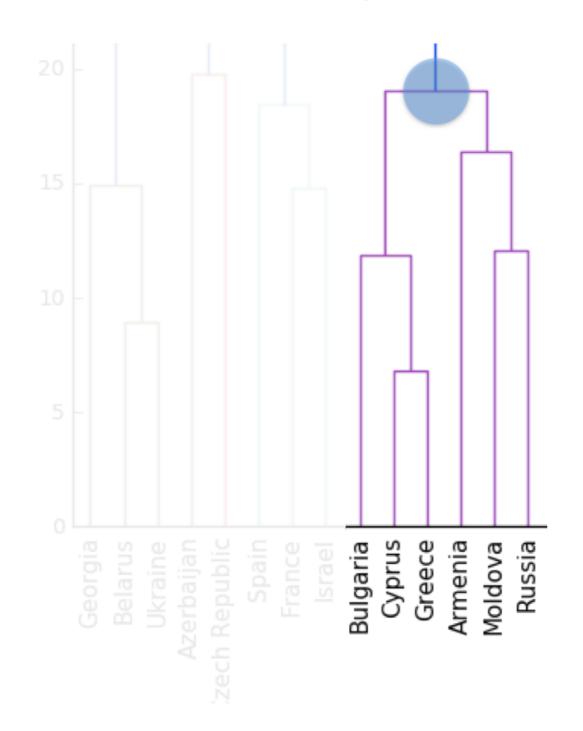




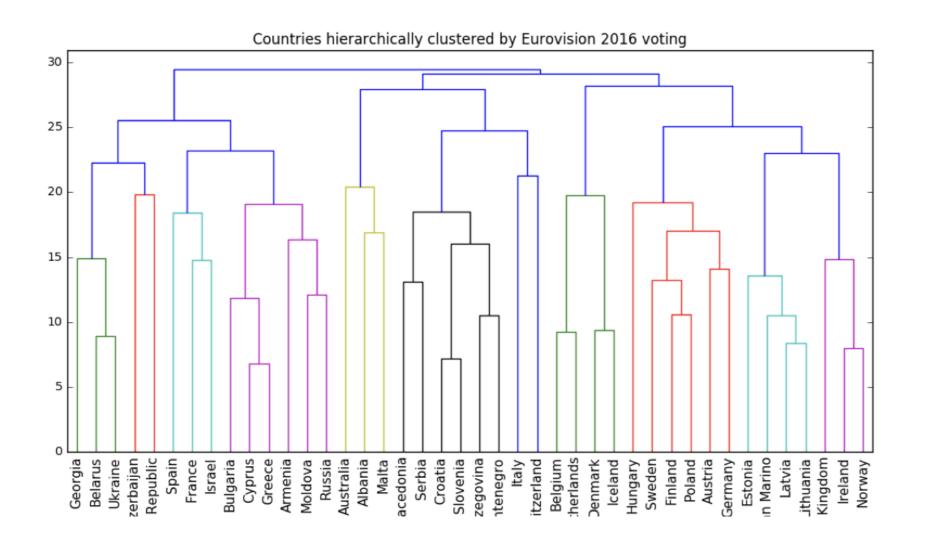














Hierarchical clustering with SciPy

• Given samples (the array of scores), and country_names

Let's practice!

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Cluster labels in hierarchical clustering

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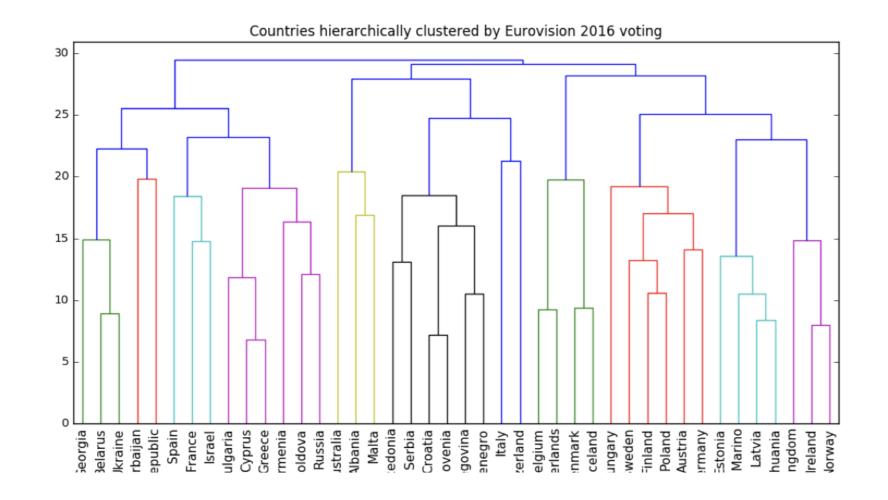


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Cluster labels in hierarchical clustering

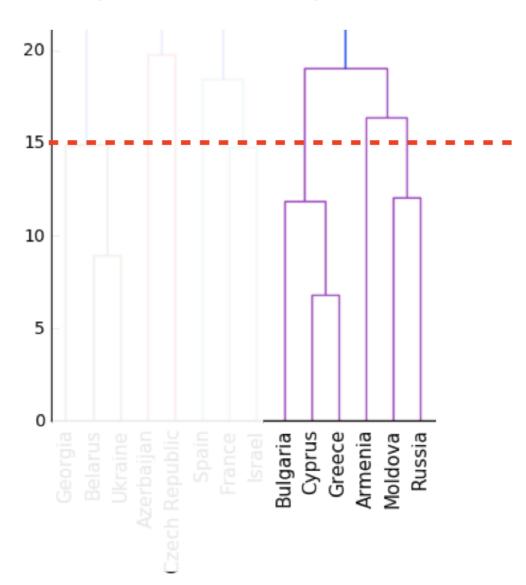
- Not only a visualization tool!
- Cluster labels at any intermediate stage can be recovered
- For use in e.g. cross-tabulations





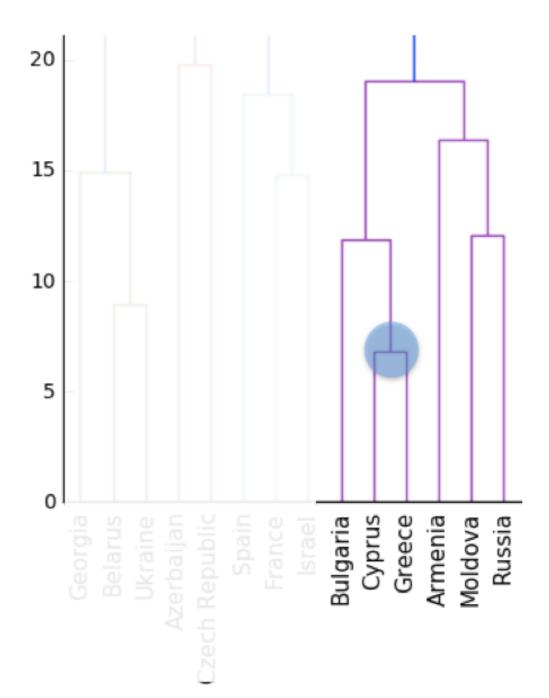
Intermediate clusterings & height on dendrogram

- E.g. at height 15:
 - Bulgaria, Cyprus, Greece are one cluster
 - Russia and Moldova are another
 - Armenia in a cluster on its own



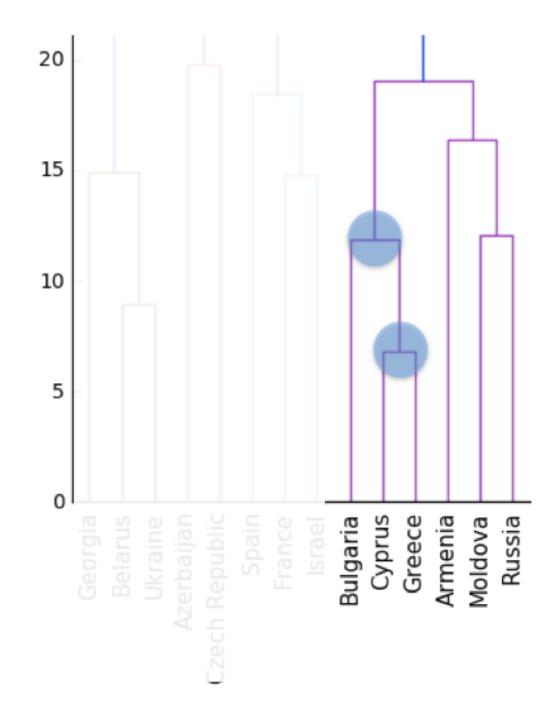
Dendrograms show cluster distances

- Height on dendrogram = distance between merging clusters
- E.g. clusters with only
 Cyprus and Greece had distance approx. 6



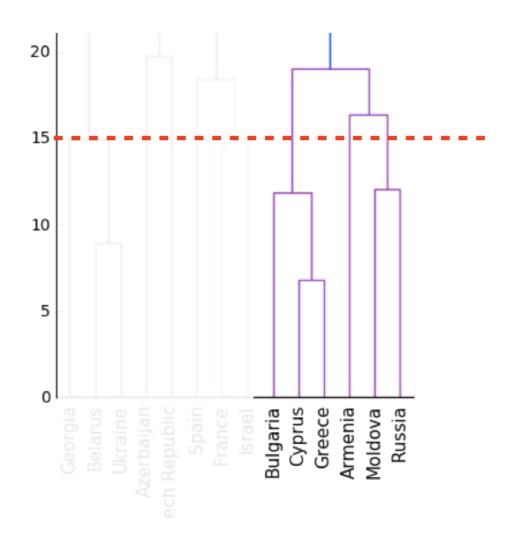
Dendrograms show cluster distances

- Height on dendrogram = distance between merging clusters
- E.g. clusters with only
 Cyprus and Greece had distance approx. 6
- This new cluster distance approx. 12 from cluster with only Bulgaria



Intermediate clusterings & height on dendrogram

- Height on dendrogram specifies max. distance between merging clusters
- Don't merge clusters further apart than this (e.g. 15)

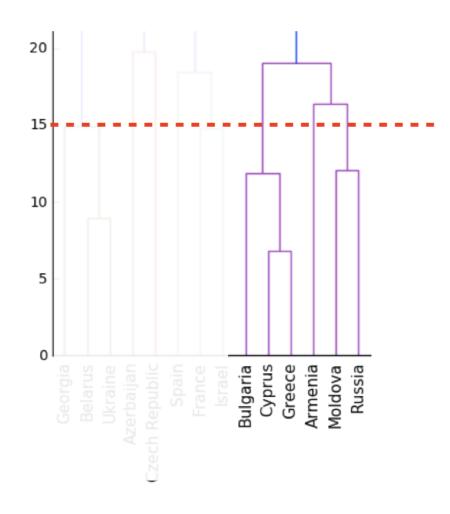


Distance between clusters

- Defined by a "linkage method"
- In "complete" linkage: distance between clusters is max.
 distance between their samples
- Specified via method parameter, e.g. linkage(samples, method="complete")
- Different linkage method, different hierarchical clustering!

Extracting cluster labels

- Use the fcluster() function
- Returns a NumPy array of cluster labels



Extracting cluster labels using fcluster

```
from scipy.cluster.hierarchy import linkage
mergings = linkage(samples, method='complete')
from scipy.cluster.hierarchy import fcluster
labels = fcluster(mergings, 15, criterion='distance')
print(labels)
```

```
[ 9 8 11 20 2 1 17 14 ... ]
```

Aligning cluster labels with country names

Given a list of strings country_names:

```
import pandas as pd
pairs = pd.DataFrame({'labels': labels, 'countries': country_names})
print(pairs.sort_values('labels'))
```

```
countries labels
5
                 Belarus
                 Ukraine
40
                    Spain
                                5
36
                Bulgaria
8
                                6
19
                   Greece
                                6
                  Cyprus
10
                                6
28
                 Moldova
```



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t-SNE for 2dimensional maps

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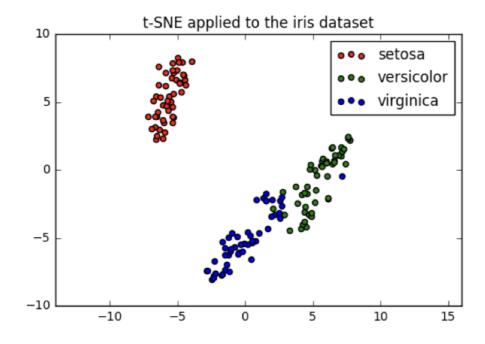


t-SNE for 2-dimensional maps

- t-SNE = "t-distributed stochastic neighbor embedding"
- Maps samples to 2D space (or 3D)
- Map approximately preserves nearness of samples
- Great for inspecting datasets

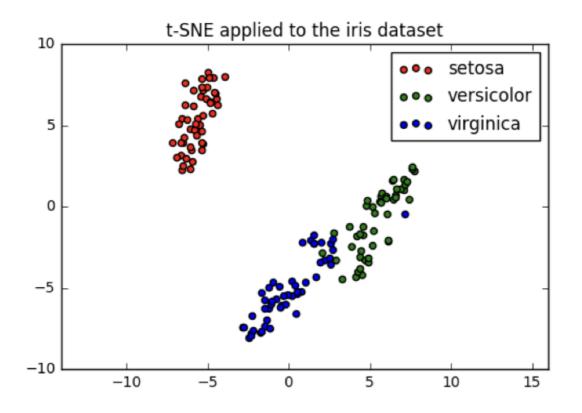
t-SNE on the iris dataset

- Iris dataset has 4 measurements, so samples are 4dimensional
- t-SNE maps samples to 2D space
- t-SNE didn't know that there were different species
- ... yet kept the species mostly separate



Interpreting t-SNE scatter plots

- "versicolor" and "virginica" harder to distinguish from one another
- Consistent with k-means inertia plot: could argue for 2 clusters, or for 3



t-SNE in sklearn

• 2D NumPy array samples

```
print(samples)
```

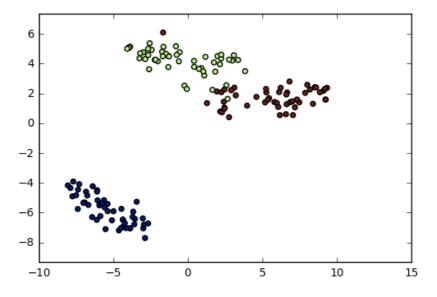
• List species giving species of labels as number (0, 1, or 2)

```
print(species)
```

```
[0, 0, 1, 2, ..., 0]
```

t-SNE in sklearn

```
import matplotlib.pyplot as plt
from sklearn.manifold import TSNE
model = TSNE(learning_rate=100)
transformed = model.fit_transform(samples)
xs = transformed[:,0]
ys = transformed[:,1]
plt.scatter(xs, ys, c=species)
plt.show()
```



t-SNE has only fit_transform()

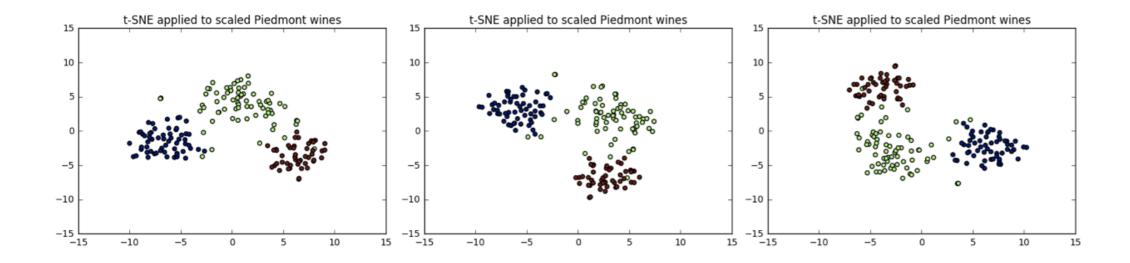
- Has a fit_transform() method
- Simultaneously fits the model and transforms the data
- Has no separate fit() or transform() methods
- Can't extend the map to include new data samples
- Must start over each time!

t-SNE learning rate

- Choose learning rate for the dataset
- Wrong choice: points bunch together
- Try values between 50 and 200

Different every time

- t-SNE features are different every time
- Piedmont wines, 3 runs, 3 different scatter plots!
- ... however: The wine varieties (=colors) have same position relative to one another



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