## Exercise 10: Hierarchical clustering of the grain data

In the video, you learnt that the SciPy <code>linkage()</code> function performs hierarchical clustering on an array of samples. Use the <code>linkage()</code> function to obtain a hierarchical clustering of the grain samples, and use <code>dendrogram()</code> to visualize the result. A sample of the grain measurements is provided in the array <code>samples</code>, while the variety of each grain sample is given by the list <code>varieties</code>.

From the course *Transition to Data Science*. Buy the entire course for just \$10 for many more exercises and helpful video lectures.

**Step 1:** Load the dataset *(done for you)*.

```
In [1]: import pandas as pd

seeds_df = pd.read_csv('../datasets/seeds-less-rows.csv')

# remove the grain species from the DataFrame, save for later
varieties = list(seeds_df.pop('grain_variety'))

# extract the measurements as a NumPy array
samples = seeds_df.values
```

## Step 2: Import:

- linkage and dendrogram from scipy.cluster.hierarchy.
- matplotlib.pyplot **as** plt.

```
In [2]: from scipy.cluster.hierarchy import linkage, dendrogram
    import matplotlib.pyplot as plt
```

**Step 3:** Perform hierarchical clustering on samples using the linkage() function with the method='complete' keyword argument. Assign the result to mergings.

```
In [3]: mergings = linkage(samples, method='complete')
```

**Step 4:** Plot a dendrogram using the <code>dendrogram()</code> function on <code>mergings</code>, <code>specifying</code> the keyword arguments <code>labels=varieties</code>, <code>leaf\_rotation=90</code>, and <code>leaf\_font\_size=6</code>. Remember to call <code>plt.show()</code> afterwards, to display your plot.

