Import some libraries. Right now, we're about to using PCA from decomposition from sklearn

```
In [1]: import numpy as np
   import pandas as pd
   import seaborn as sb

import matplotlib.pyplot as plt
   from pylab import rcParams

import sklearn
   from sklearn import datasets
   from sklearn import decomposition
   from sklearn.decomposition import PCA
```

Configuring the canvas

```
In [2]: %matplotlib inline
    rcParams['figure.figsize'] = 5,4
    sb.set_style('whitegrid')
```

Read Iris Datasets

```
In [3]: iris = datasets.load_iris()
    x = iris.data
    variable_names = iris.feature_names
```

We're about to calculate the PCA. First thing first, we use decomposition.PCA() function, then fit_transform(x) and we use pca.explained_variance_ratio_ to explained first few component variance

And then, we calculate the sum(). Remember, make sure to retain at least 70% of dataset's

original information

```
In [5]: pca.explained_variance_ratio_.sum()
```

Out[5]: 1.0

In [11]: comp = pd.DataFrame(pca.components_, columns = variable_names)
comp

Out[11]:

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	0.361387	-0.084523	0.856671	0.358289
1	0.656589	0.730161	-0.173373	-0.075481
2	-0.582030	0.597911	0.076236	0.545831
3	-0.315487	0.319723	0.479839	-0.753657

In [8]: sb.heatmap(comp)

Out[8]: <matplotlib.axes._subplots.AxesSubplot at 0xc03c490>

