Lab 3_3

3. An experiment with PCA

For this problem, we'll be using the animals with attributes data set. Go to

http://attributes.kyb.tuebingen.mpg.de

and, under "Downloads", choose the "base package" (the very first file in the list). Unzip it and look over the various text files.

```
In [1]:
         %matplotlib inline
         import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         from pandas import read fwf
         from matplotlib.pyplot import figure
         from sklearn.model_selection import cross_val_score
         from sklearn.tree import DecisionTreeClassifier
         import graphviz
         from sklearn import tree
         from sklearn.ensemble import AdaBoostClassifier
         from sklearn.model selection import cross val predict
         from sklearn.metrics import confusion matrix
         from sklearn.ensemble import RandomForestClassifier
         from gensim.models import Word2Vec
         from sklearn.decomposition import PCA
         from matplotlib import pyplot
         from sklearn.preprocessing import StandardScaler
         from sklearn.decomposition import PCA
         from pylab import rcParams
         import sys
         if not sys.warnoptions:
             import warnings
             warnings.simplefilter("ignore")
```

(a) This is a small data set that has information about 50 animals. The animals are listed in classes.txt. For each animal, the information consists of values for 85 features: does the animal have a tail, is it slow, does it have tusks, etc. The details of the features are in predicates.txt. The full data consists of a 50×85 matrix of real values, in predicate-matrix-continuous.txt. Load this real-valued array.

```
In [2]: data = read_fwf('predicate-matrix-continuous.txt', header = None, sep = '\t')
```

```
classes = read_fwf('classes.txt', header = None, sep = '\t')
```

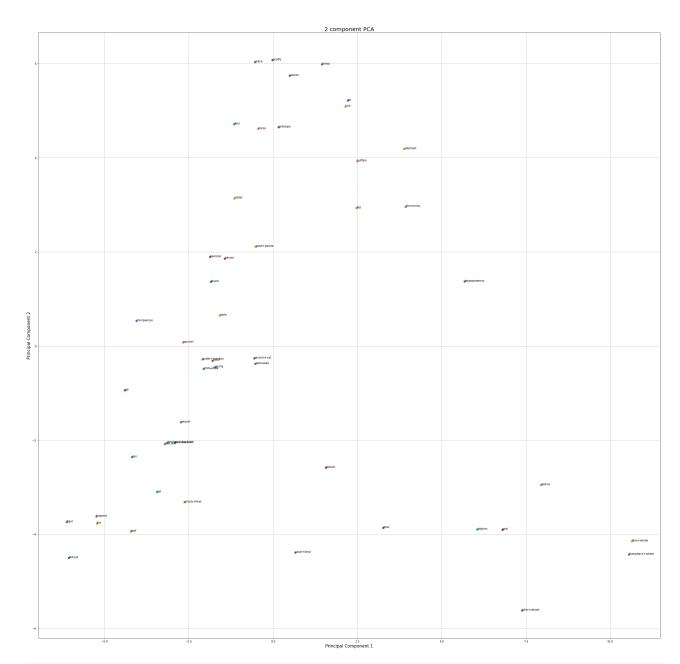
(b) We would like to visualize these animals in 2-d. Show how to do this with a PCA projection from R85 to R2. Show the position of each animal, and label them with their names.

Python notes: You will need to make the plot larger by prefacing your code with

```
from pylab import rcParams
rcParams['figure.figsize'] = 10, 10
```

(or try a different size if this doesn't seem right).

```
In [3]:
         x = StandardScaler().fit_transform(data)
In [4]:
         pca = PCA(n_components=2)
         principalComponents = pca.fit_transform(x)
         principalDf = pd.DataFrame(data = principalComponents
                       , columns = ['principal component 1', 'principal component 2'])
In [5]:
         finalDf = pd.concat([principalDf, classes], axis = 1)
In [7]:
         fig = plt.figure()
         rcParams['figure.figsize'] = 40, 40
         ax = fig.add subplot(1,1,1)
         ax.set xlabel('Principal Component 1', fontsize = 15)
         ax.set_ylabel('Principal Component 2', fontsize = 15)
         ax.set_title('2 component PCA', fontsize = 20)
         targets = finalDf.values[:,3]
         for target in zip(targets):
             indicesToKeep = finalDf.values[:,3] == target
             ax.scatter(finalDf.loc[indicesToKeep, 'principal component 1']
                         , finalDf.loc[indicesToKeep, 'principal component 2']
                         , s = 50)
         #ax.legend(targets)
         ax.grid()
         words = finalDf.values[:,3]
         for i, word in enumerate(words):
             pyplot.annotate(word, xy=(finalDf.loc[i, 'principal component 1']
                         , finalDf.loc[i, 'principal component 2']))
         pyplot.show()
```



In []: