EE258 HOMEWORK #3

(ONLINE SUBMISSION ONLY – MAKE A PDF FILE INCLUDING ALL THE REQUIRED SUBMISSIONS)

DUE: SEP 26th

- 1. **Read and practice** the codes in Chapter 2 (pages 33-68) of textbook "Hands-On Machine Learning with Scikit-learn and tensorflow" (NOT TO BE SUBMITTED)
- **2. Get the data:** For this homework, use the iris dataset under Files-> Datasets on Canvas. (submit the code)

Hint: Pandas knows to treat rows with 'NA' as missing values

- 3. **Take a look at the data structure:** Write a brief paragraph about the data set related to each of the following observations (submit the code, the result, and the paragraphs):
 - a. Look at the top five rows of the data set
 - b. Get a quick description of the data: Notice if there are any missing values or categorical features
 - c. Get a summary of the numerical features
 - d. Plot the histogram of the numerical features
- 4. **Discover and visualize the data to gain insights:** (submit the plots, code, and your observations of each plot)
 - a. Obtain scatter matrix
 - b. Obtain the correlations among features and comment
- **5. Data Cleaning** (submit the code, results and plots showing the changes in data)
 - a. Drop the data points with NA in it
 - b. Tidy up the data by renaming the "class" data point correctly.
 - c. Remove the outliers: drop the 'Iris-setosa' rows with a sepal width less than 2.5 cm.

- d. One of the data collectors forgot to convert the sepal length for "Iris-versicolor" to cm, instead added the data as meters. Find those and convert them to cm.
- e. Handle the categorical variables
- f. Save the clean data into a new file.
- 6. **Utilizing a perceptron learning algorithm to check if a flower is "iris-setosa" or not ?** (submit plots, results, code explainations)
 - a. Modify the clean dataset such that you have class =+1: for "iris-setosa", and class = -1 for others.
 - b. Is the new data set linearly separable? Will perceptron algorithm work?
 - c. Explain what is the functionality of each line of code in the perceptron.py file.
 - d. Separate the data into test and training data
 - e. Use the below **perceptron.py** to train your perceptron
 - f. Does the algorithm converge? What is niter in the code?
 - g. Obtain a plot of the training accuracy as a function of epocs (number of times you go over the entire training data)
 - h. Obtain the test data accuracy
 - i. Sketch the decision boundary (the line that separates the data; obtained via perceptron algorithm) and the scatter plot of data points. For "iris-setosa" use a different identifier than the rest of the classes.

As performano	ce metric use accuracy:
	Number of data points predicted correctly
accuracy =	
•	Total number of data points

```
# perceptron.py
import numpy as np
class Perceptron(object):
   def init (self, rate = 0.01, niter = 10):
     self.rate = rate
      self.niter = niter
   def fit(self, X, y):
      """Fit training data
     X : Training vectors, X.shape : [#samples, #features]
      y : Target values, y.shape : [#samples]
      # weights
      self.weight = np.zeros(1 + X.shape[1])
      # Number of misclassifications
      self.errors = [] # Number of misclassifications
      for i in range (self.niter):
         err = 0
         for xi, target in zip(X, y):
            delta w = self.rate * (target - self.predict(xi))
            self.weight[1:] += delta w * xi
            self.weight[0] += delta w
            err += int(delta w != 0.0)
         self.errors.append(err)
      return self
   def net input(self, X):
      """Calculate net input"""
      return np.dot(X, self.weight[1:]) + self.weight[0]
   def predict(self, X):
      """Return class label after unit step"""
      return np.where(self.net input(X) \geq 0.0, 1, -1)
```

```
>>> # import Perceptron from perceptron.py
>>> from perceptron import Perceptron
>>> pn = Perceptron(0.1, 10)
>>> pn.fit(X, y)
>>> plt.plot(range(1, len(pn.errors) + 1), pn.errors, marker='o')
>>> plt.xlabel('Epochs')
>>> plt.ylabel('Number of misclassifications')
>>> plt.show()
```

```
from matplotlib.colors import ListedColormap
def plot_decision_regions(X, y, classifier, resolution=0.02):
   # setup marker generator and color map
  markers = ('s', 'x', 'o', '^i, 'v')
   colors = ('red', 'blue', 'lightgreen', 'gray', 'cyan')
   cmap = ListedColormap(colors[:len(np.unique(y))])
   # plot the decision surface
   x1_{\min}, x1_{\max} = X[:, 0].min() - 1, X[:, 0].max() + 1
   x2_{min}, x2_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
   xx1, xx2 = np.meshgrid(np.arange(x1 min, x1 max, resolution),
  np.arange(x2_min, x2_max, resolution))
   Z = classifier.predict(np.array([xx1.ravel(), xx2.ravel()]).T)
   Z = Z.reshape(xx1.shape)
   plt.contourf(xx1, xx2, Z, alpha=0.4, cmap=cmap)
  plt.xlim(xx1.min(), xx1.max())
   plt.ylim(xx2.min(), xx2.max())
   # plot class samples
   for idx, cl in enumerate(np.unique(y)):
      plt.scatter(x=X[y == cl, 0], y=X[y == cl, 1],
      alpha=0.8, c=cmap(idx),
      marker=markers[idx], label=cl)
```

Code is modified version of the code in "Python Machine Learning by Sebastian Raschka"