

written3

October 17, 2023

1 STP598 Machine Learning & Deep Learning

1.1 Written Assignment 3

1.1.1 Due 11:59pm Friday Oct. 27, 2023 on Canvas

1.1.2 name, id

```
[1]: import warnings
import numpy as np
import matplotlib.pyplot as plt
```

1.2 Classification

We have learned several classification methods including LogisticRegression, LinearDiscriminantAnalysis, QuadraticDiscriminantAnalysis, RandomForestClassifier and GaussianProcessClassifier. It is of interest to compare them side-by-side. To visualize the results in a 2d picture, it is important to plot their boundaries. In this homework, we will compare the classification boundaries of different methods.

Let's first look at a synthetic example.

```
[2]: import warnings
import numpy as np
from sklearn.datasets import make_blobs
from sklearn.linear_model import LogisticRegression

# make 3-class dataset for classification
centers = [[-5, 0], [0, 1.5], [5, -1]]
X, y = make_blobs(n_samples=1000, centers=centers, random_state=2021)
transformation = [[0.4, 0.2], [-0.4, 1.2]]
X = np.dot(X, transformation)

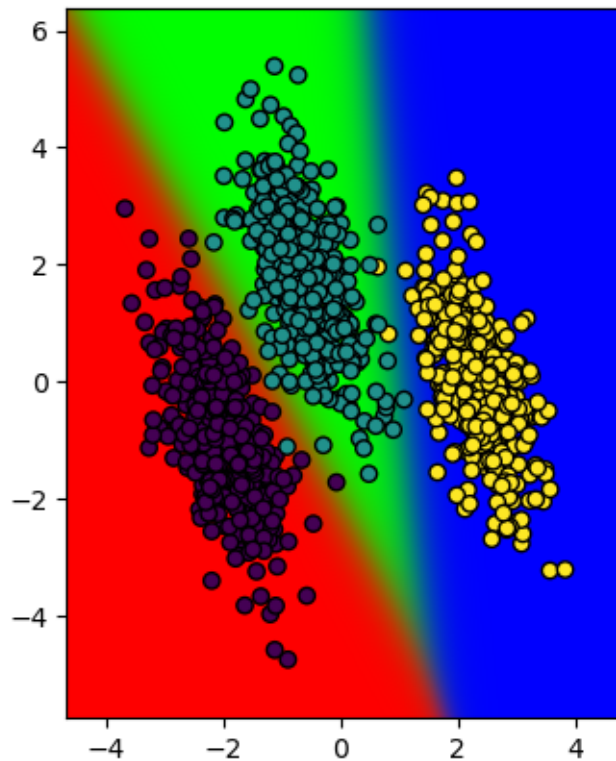
# define the plotting functions
def clf_bdy(clf, X, y):
    K=len(np.unique(y))
    # plot training points
    plt.scatter(X[:, 0], X[:, 1], c=plt.get_cmap('viridis',K).colors[y],
edgecolor='k', cmap=plt.get_cmap('viridis'))
```

```

# create a mesh to plot in
h = .02 # step size in the mesh
x_min, x_max = X[:, 0].min() - 1, X[:, 0].max() + 1
y_min, y_max = X[:, 1].min() - 1, X[:, 1].max() + 1
xx, yy = np.meshgrid(np.arange(x_min, x_max, h),
                     np.arange(y_min, y_max, h))
# Plot the decision boundary. For that, we will assign a color to each
# point in the mesh [x_min, x_max][y_min, y_max].
Z = clf.predict_proba(np.c_[xx.ravel(), yy.ravel()])
# Put the result into a color plot
Z = Z.reshape((xx.shape[0], xx.shape[1], -1))
plt.imshow(Z, extent=(x_min, x_max, y_min, y_max), origin="lower", cmap=plt.
    ↳get_cmap('viridis'))

# classify it and plot the boundary
clf = LogisticRegression(solver='sag', max_iter=100, random_state=2021,
                        multi_class='multinomial').fit(X, y)
clf_bdy(clf, X, y)

```

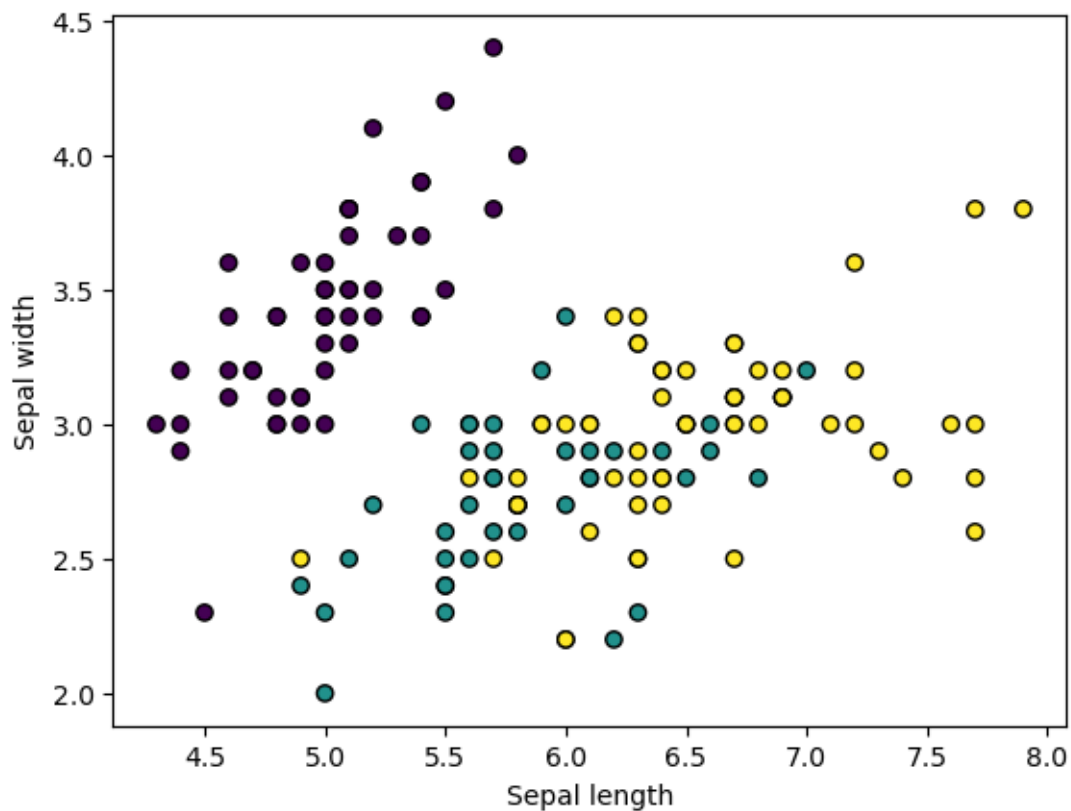


1.2.1 iris data

Now consider iris data and classify them with the first two features using different methods. Let's load the data.

```
[3]: from sklearn import datasets

iris = datasets.load_iris()
X = iris.data[:, :2]
y = iris.target
K = len(np.unique(y))
plt.scatter(X[:, 0], X[:, 1], c=plt.get_cmap('viridis', K).colors[y],
            edgecolor='k', cmap=plt.get_cmap('viridis'))
plt.xlabel('Sepal length')
plt.ylabel('Sepal width')
plt.show()
```



1.2.2 Question 1

Fit LDA and QDA models to the iris data and save the fitted models as **lda** and **qda** respectively.

```
[ ]: # Fit LDA and QDA
```

Now compare their decision boundaries in the following plot.

```
[ ]: # plot decision boundairs of LDA and QDA
```

1.2.3 Question 2

Fit RandomForest and GaussianProcess models to the iris data. For RandomForest, use **50 random trees** (`n_estimators`). For GaussianProcess, use anisotropic **Radial-basis function** kernel (aka squared-exponential kernel, RBF). Save the fitted models as **rfc** and **gpc** respectively.

Note! Plesae set `random_state=2023` in RandomForest

```
[ ]: # Fit RandomForest and GaussianProcess
```

Now compare their decision boundaries in the following plot.

```
[ ]: # plot decision boundairs of RandomForest and GaussianProcess
```