211112238-ml-lab02

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0.1 Gradient Descent

```
[]: import numpy as np
     import pandas as pd
     import matplotlib.pyplot as plt
     # Sigmoid function
     def sigmoid(z):
         return 1 / (1 + np.exp(-z))
     # Logistic regression function with gradient descent
     def gd(X, y):
         X = np.insert(X, 0, 1, axis=1)
         weights = np.ones(X.shape[1])
         lr = 0.5
         for _ in range(5000):
             y_hat = sigmoid(np.dot(X, weights))
             weights = weights + lr * (np.dot((y - y_hat), X) / X.shape[0])
         return weights[1:], weights[0]
     # Function to plot points
```

```
plt.figure(figsize=(8, 6))
         class_0 = X[y == 0]
         class_1 = X[y == 1]
         plt.scatter(class_0, [0] * len(class_0), c='blue', label='Class 0', u

marker='o')
         plt.scatter(class_1, [0] * len(class_1), c='red', label='Class 1', __

marker='x')
         plt.title(title)
         plt.xlabel('X')
         plt.legend()
         plt.grid(True)
     # Convert data to DataFrame
     →1]})
     # Plot original points
     plot_points(data['X'], data['y'], title="Scatter Plot of Points with Binaryu
       ⇔Class Labels")
     # Train logistic regression using gradient descent
     weights, bias = gd(data['X'].values.reshape(-1, 1), data['y'].values)
     # Plot logistic regression line
     x_range = np.linspace(min(data['X']), max(data['X']), 100)
     log_reg_line = sigmoid(weights * x_range + bias)
     plt.plot(x_range, log_reg_line, color='green', label='Logistic Regression Line')
     plt.legend()
     plt.show()
[17]: import numpy as np
     import matplotlib.pyplot as plt
     from sklearn.linear_model import LinearRegression
     # Generate example data
     X = \text{np.array}([1, 2, 3, 4, 5, 6, 7, 8]).\text{reshape}(-1, 1) \# \text{Reshape to make it } a_{\perp}
      →2D array(Column wise)
     y = np.array([0, 0, 0, 0, 1, 1, 1, 1])
     # Create a linear regression model
     model = LinearRegression()
     # Train the model
     model.fit(X, y)
```

def plot_points(X, y, title="Scatter Plot"):

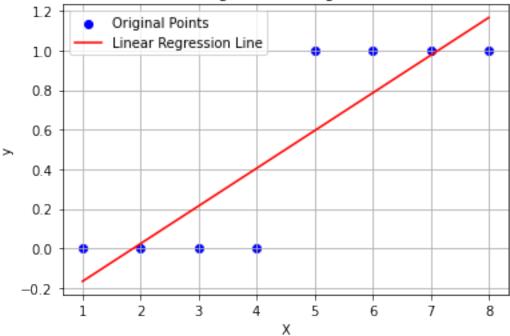
```
# Get the slope (coefficient) and intercept
slope = model.coef_[0]
intercept = model.intercept_

# Plot the original points
plt.scatter(X, y, color='blue', marker='o', label='Original Points')

# Plot the linear regression line
x_range = np.linspace(min(X), max(X), 100)
y_pred = model.predict(x_range.reshape(-1, 1))
plt.plot(x_range, y_pred, color='red', label='Linear Regression Line')

plt.title('Linear Regression using scikit-learn')
plt.xlabel('X')
plt.ylabel('y')
plt.legend()
plt.grid(True)
plt.show()
```





0.2 Support Vector Machine

```
[19]: %matplotlib inline
import numpy as np
import matplotlib.pyplot as plt
from scipy import stats
```

```
[]: import glob
     import pandas as pd
     from sklearn.svm import SVR
     from sklearn.metrics import mean_squared_error, mean_absolute_error, r2_score
     from sklearn.preprocessing import StandardScaler
     from sklearn.model_selection import GridSearchCV
     # Function to perform SVM Regression with RBF kernel
     def SVRRBFKernelMetrics(x_train, y_train, x_test, y_test, C, gamma):
         scaler = StandardScaler()
         x_train_scaled = scaler.fit_transform(x_train)
         x_test_scaled = scaler.transform(x_test)
         svr = SVR(kernel='rbf', C=C, gamma=gamma)
         svr.fit(x_train_scaled, y_train)
         y_pred = svr.predict(x_test_scaled)
         rmse = mean_squared_error(y_test, y_pred, squared=False)
         mae = mean_absolute_error(y_test, y_pred)
         r2 = r2_score(y_test, y_pred)
         return rmse, mae, r2
     folders = ["diabetes-5-fold", "machineCPU-5-fold", "mortgage-5-fold"]
     for folder in folders:
         print(folder)
         # Define the file pattern
         file_pattern = "*tra.dat"
         training_files = glob.glob("./" + folder + "/" + file_pattern)
         file_pattern = "*tst.dat"
         testing_files = glob.glob("./" + folder + "/" + file_pattern)
         alpha_values = [2 ** i for i in range(0, 2, 2)]
         degree = [2, 3]
         for d in degree:
             svm_map = {}
```

```
for C in alpha_values:
                 for gamma in alpha_values:
                     trmse = 0
                     tmae = 0
                     tr2 = 0
                     for train_file, test_file in zip(training_files, testing_files):
                         df = pd.read_csv(train_file, delimiter=',', header=None,__

comment='@')

                         df_test = pd.read_csv(test_file, delimiter=',',u
      ⇔header=None, comment='@')
                         x_train = df.iloc[:, :-1]
                         y_train = df.iloc[:, -1]
                         x_test = df_test.iloc[:, :-1]
                         y_test = df_test.iloc[:, -1]
                         rmse, mae, r2 = SVRRBFKernelMetrics(x_train, y_train, __
      →x_test, y_test, C, gamma)
                         trmse += rmse
                         tmae += mae
                         tr2 += r2
                     trmse /= 5
                     tmae /= 5
                     tr2 /= 5
                     svm_map[(folder, d, C, gamma)] = (trmse, tmae, tr2)
             # Print or store the results as needed
             # for key, values in svm_map.items():
                   print(f"{key}: RMSE={values[0]}, MAE={values[1]}, R2={values[2]}")
[]: from sklearn.metrics import confusion_matrix
     import matplotlib.pyplot as plt
     import seaborn as sns
     # Example true labels and predicted labels
     y_true = [1, 2, 3, 4, 5, 6, 7, 8]
     y_pred = [0, 0, 0, 0, 1, 1, 1, 1]
     # Compute confusion matrix
     cm = confusion_matrix(y_true, y_pred)
     # Print the confusion matrix
```

[]: