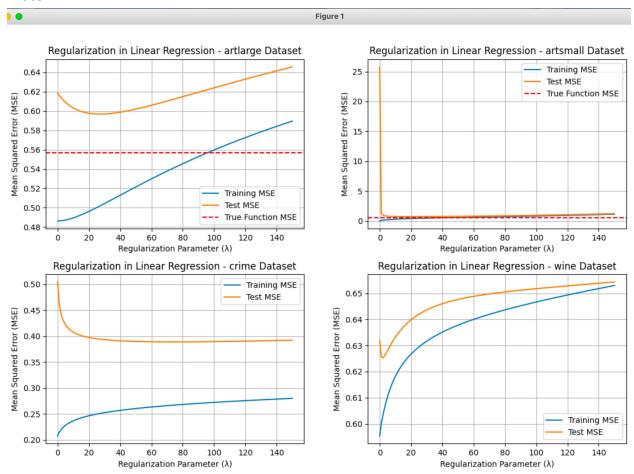
Sunilsakthivel Sakthi Velavan Roni Khardon CSCI-B 555 12 Oct.2023

Programming Project 2

Task 1:

For this experiment, 4 different plots were generated entailing the training and the testing MSE scores with respect to the λ values between 0 and 150. As seen below, the artificial datasets (artsmall and artlarge), we also compare the plotted MSEs with the MSE of the hidden true functions generating the data that give 0.533 (artsmall), and 0.557 (artlarge).

Plots:



Dataset	λ of Lowest MSE	Lowest MSE
---------	-----------------	------------

artlarge	27	0.5967438457327094
artsmall	18	0.7202788056527154
crime	75	0.38902338771344425
wine	2	0.625308842304743

Observations:

For each dataset, one can make on the test MSE curves follow a pattern of a high initial MSE, followed by a quick drop to a minimum point followed by a steady growth as λ values from 0 to 150. Simultaneously, train MSE curves typically start off with a low MSE value and experience a quick increase to a point where the test MSE also experiences its lowest MSE and also followed by a slower increase (much like the tail end of the U-shaped test MSE curve). This actually explains why the training set MSE cannot be trusted to select the right λ values as when you use a small or zero value of λ , the model can fit the training data extremely well, resulting in a very low training set MSE. However, this does not necessarily mean that the model will generalize well to unseen data. The model might likely overfit the training data, capturing noise in the data, which will lead to poor performance on the test data. However, on the test set error, one can observe the U-shaped curve for all 4 datasets where a minimum MSE inflection point with an optimal λ value is reached after which larger λ values only increase the MSE. This suggests that the test MSE will be relatively high for small values of λ , indicating a model that overfits the training data. As λ increases, the test MSE decreases, suggesting better generalization and reduced overfitting. However, beyond a certain point, as λ becomes too large, the test MSE starts to increase, indicating underfitting or a too simple model.

While this fact can be stated in principle, the nuances of λ 's impact on MSE actually vary between the 4 datasets. For instance, the minimal increase in λ for the artsmall dataset showcases how quickly the test MSE drops and converges at its true optimal value at an MSE of 0.533. This is in direct contrast to the artlarge dataset whose significantly larger size makes it more difficult to reach its true optimal λ value and its concurrent minimum MSE.

Task 2:

For this experiment, 10 fold cross validation was implemented on the training set to choose a λ value in the same range. We then retrained it on the entire training set and evaluated its performance on the training set. For parameter selection with a fixed train/test split, the data is first split into 10 disjoint portions, then each value of λ in the range V1, ..., VK, the following steps are performed: For each fold (i in 1...10), we train

the ridge regression model on all portions except i. We then set the model on portion i and record the validation error for that fold. The average performance of λ on the 10 folds is recorded, the λ with the best average performance is selected, the model is then retrained on the entire training set (D) using the chosen λ , and the performance of the model with the selected λ is evaluated on the test set.

Results:

Dataset	Selected λ	Validation MSE	Test MSE	Runtime (seconds)
artlarge	27	0.5991366438451824	0.5967438457327094	3.0185658931732178
artsmall	16	0.703037732817224	0.7210310814686591	1.919619083404541
crime	150	0.344590450459746	0.39233899203438105	2.3310739994049072
wine	1	0.639636967500184	0.6256423273038403	0.5152280330657959

Observations:

For the "artlarge" dataset, Task 2 selected the same λ as in Task 1, and the test MSE remained very close. For the "artsmall" dataset, Task 2 chose a different λ compared to Task 1, and the test MSE was slightly higher in Task 2. For the "wine" dataset, Task 2 selected a different λ , and the test MSE was similar to Task 1. For the "crime" dataset, Task 2 selected a different λ , and the test MSE increased slightly compared to Task 1. However, the crime dataset is less visually actualized as the λ chosen here is 150 which suggests that the minimum point on the curve could be outside the range of [1, 150]. This is a relative possibility as the test MSE curve for crime could be argued to never quite reach its minimum within the same range either. However, overall, Task 2's approach allows for more flexibility in selecting λ based on cross-validation results.

Task 3:

Given the formulation of Bayesian linear regression with the simple prior w $^{\sim}$ N(0,(1/ α)I), we can recall that the evidence function (and evidence approximation) gives a method to pick the parameters α and β . For this experiment, we implemented Bayesian Model Selection within the context of Baysian linear regression. This was done through integrating an iterative algorithm that converges and selects the ideal values of α and β using the training set to arrive at the values for m_n and SN, where mN is utilized to calculate the MSE on the test set for prediction. We initialized both α and β to random values in the range [1, 10] and a convergence threshold of 0.0001 that helps

the algorithm ascertain that we've converged close enough to the actual values of α and β , and indirectly mN .

We first calculate the lambda values (λ_i) through determining the eigenvalues of the matrix $(\beta \Phi^T \Phi)$. These eigenvalues are used in the calculations to update alpha and beta. Then, S_N is computed using equation (3.53), and m_N is calculated based on equation (3.54) which uses the value of S_N. These quantities represent the posterior covariance matrix and the posterior mean of the model coefficients, respectively. Next, using equations (3.91) and (3.92), we generate the values for gamma (using the old α and λ_i) and the new updated α . Then using equation (3.95), we generate the updated β (once again using the gamma dependent on the old α). Once the new α and β values are created, this algorithm reiterates until the difference between the old and new α and β values are less than the preset threshold value of .0001. Once the ideal α and β values have been converged upon, we can visualize the effective λ_i value can be computed as α/β and the m_N that was dependent on ideal β can be used to generate the predicted target value with which to generate the test MSE. Fundamentally, the task aims to find the optimal values of alpha and beta that result in the best predictions and minimal test MSE for Bayesian linear regression. It provides a Bayesian framework for model selection and parameter estimation.

Results:

Dataset	α	β	λ (α/β)	Test MSE	Runtime
artlarge	10.285784709807992	1.8603095111059353	5.52907172080908	0.6083085987853174	0.03248906135559082
artsmall	5.154543569445669	3.154402098963837	1.634079425428623	1.0635277958316793	0.0455479621887207
crime	425.6452824267614	3.250432133920129	130.95036748649758	0.39110230579177896	0.09948515892028809
wine	6.163857650413885	1.6098094296700247	3.828936231089999	0.6267461570340823	0.005588054656982422

Observations:

While for artlarge, artsmall and crime datasets, the λ values in Task 1 and Task 3 are drastically different with only wine having a close λ value to Task 1's. However, artlarge and artsmall both generated a significantly higher MSE compared to Task 1, indicating that the inclusion of α and β aren't particularly helpful. However, Task 3 achieved a lower test MSE compared to Task 1 on the crime dataset, suggesting improvement in predictive performance.

Discussion of Results:

In Task 2, the effective λ was determined as a result of cross-validation, which led to different λ values for each dataset. Also, Task 2 provided test MSE values that varied based on the λ selection through cross-validation. Task 3 calculated the effective λ as a result of the Bayesian model selection algorithm. This process also led to different λ values for each dataset and also provided test MSE values that varied based on the a and β selection through the Bayesian algorithm. The test set MSE values varied between Task 2 and Task 3. Task 3, the Bayesian model selection, tended to produce lower MSE values, indicating that it might lead to models with better predictive performance. However, this improvement in MSE might come at the cost of longer computation time. Comparatively, both Task 2 and Task 3 resulted in different effective λ values, suggesting that the choice of λ depends on the model selection method and dataset parameters. Overall, Task 2 seemed more suitable when computational efficiency is critical, and you want a data-driven approach for λ selection. It can work well when a quick model evaluation is needed, even if it might not yield the lowest possible test MSE. However, Task 3 is preferable when the focus is on achieving the best predictive performance. While it involves slightly longer run times, it leads to lower test MSE, which can be crucial in applications where accuracy is a lot more integral to the validity of the predicted values.

```
Codebase:
import numpy as np
import matplotlib
matplotlib.use('TkAgg')
import matplotlib.pyplot as plt
import pandas as pd
from sklearn.model_selection import KFold
import time
datasets = ['artlarge', 'artsmall', 'crime', 'wine']
# Defines a range of regularization parameters (λ)
lambda values = np.arange(0, 151, 1)
# Create subplots
fig, axs = plt.subplots(2, 2, figsize=(12, 10))
fig.tight layout(pad=5.0) # Add padding between subplots
def ridge_regression(X, y, lambda val):
 N = X.shape[0]
 M = X.shape[1]
 # Calculate \Phi^\mathsf{T}\Phi
 phi phi transpose = np.dot(X.T, X)
 lambda I = lambda val * np.identity(M)
 inv term = np.linalg.inv(np.add(lambda I, phi phi transpose))
 # Calculate \Phi^\mathsf{T}t
 phi transpose y = np.dot(X.T, y)
  # Calculate w using the equation: w = (\lambda I + \Phi^T \Phi)^{-1} \Phi^T t
 w = np.dot(inv term, phi transpose y)
 # Calculate predictions
 y pred = np.dot(X, w)
 # Calculate Mean Squared Error (MSE)
mse = np.sum((y - y pred) ** 2) / N
return w, mse
#Task 1: Regularization
```

print("Task 1:")

for i, dataset in enumerate(datasets):
 # Divide subplots into rows and columns

```
row = i // 2
 col = i % 2
ax = axs[row, col]
train data = pd.read csv('pp2data/train-'+dataset+'.csv', header=None)
 train_labels = pd.read csv('pp2data/trainR-'+dataset+'.csv', header=None)
test data = pd.read csv('pp2data/test-'+dataset+'.csv', header=None)
test labels = pd.read csv('pp2data/testR-'+dataset+'.csv', header=None)
X train = train data.values
 y train = train labels.values.ravel()
X test = test data.values
y test = test labels.values.ravel()
# Lists to store MSE values
train mse = []
test mse = []
# Iterate through lambda values and fit Ridge regression models
for lambda val in lambda values:
    # Fit the Ridge regression model
 w, train mse val = ridge regression(X train, y train, lambda val)
     # Predict on training and test data
     y test pred = np.dot(X test, w)
  # Calculate test set MSE
  test mse val = np.sum((y test pred - y test) ** 2) / len(y test)
train mse.append(train mse val)
    test mse.append(test mse val)
 # Plot the results on the current subplot
 ax.plot(lambda_values, train_mse, label="Training MSE")
ax.plot(lambda values, test mse, label="Test MSE")
 if dataset == 'artsmall':
 ax.axhline(y=0.533, color='r', linestyle='--', label="True Function
elif dataset == 'artlarge':
     ax.axhline(y=0.557, color='r', linestyle='--', label="True Function
 ax.set xlabel("Regularization Parameter (λ)")
 ax.set ylabel("Mean Squared Error (MSE)")
ax.set title("Regularization in Linear Regression - "+dataset+" Dataset")
ax.legend()
ax.grid(True)
```

```
print(f"Dataset: {dataset}")
  minMSElambda = test_mse.index(min(test_mse))
  print("Lambda:"+str(minMSElambda))
  print("Min MSE:"+str(min(test mse)))
  print("\n")
# Show the entire figure with subplots
plt.show()
#Task 2
# Initialize variables to store results
best lambda = {} # Dictionary to store the best λ for each dataset
best avg validation mse = {} # Dictionary to store the best average
validation MSE for each dataset
test mse = {} # Dictionary to store the test MSE for each dataset
runtime = {}
for i, dataset in enumerate(datasets):
 train data = pd.read csv('pp2data/train-'+dataset+'.csv', header=None)
 train labels = pd.read csv('pp2data/trainR-'+dataset+'.csv', header=None)
 test data = pd.read csv('pp2data/test-'+dataset+'.csv', header=None)
 test labels = pd.read csv('pp2data/testR-'+dataset+'.csv', header=None)
 X train = train data.values
  y_train = train_labels.values.ravel()
  X test = test data.values
 y test = test labels.values.ravel()
  # Perform 10-fold cross-validation for \lambda selection
 kf = KFold(n splits=10, shuffle=True, random state=42)
  best lambda[dataset] = None
  best avg validation mse[dataset] = float('inf')
 test mse[dataset] = None
 start time = time.time()
  for lambda val in lambda values:
     validation mses = []
      for train index, val index in kf.split(X train):
          X fold train, X fold val = X train[train index], X train[val index]
   y fold train, y fold val = y train[train_index], y_train[val_index]
         w, train mse_val = ridge_regression(X fold_train, y fold_train,
lambda val)
```

```
# Predict on training and test data
y_val_pred = np.dot(X_fold_val, w)
          # Calculate test set MSE
          validation mse = np.sum((y fold val - y val pred) ** 2) /
len(y_fold_val)
         validation_mses.append(validation_mse)
  avg validation mse = np.mean(validation mses)
      \# Check if this \lambda has the lowest average validation MSE
       if avg validation mse < best avg validation mse[dataset]:</pre>
           best avg validation mse[dataset] = avg_validation_mse
      best lambda[dataset] = lambda val
          w, test mse val = ridge regression(X train, y train,
best lambda[dataset])
        # Evaluate the model on the test set
         y test pred = np.dot(X test, w)
      # Calculate test set MSE
         validation mse = np.sum((y test - y test pred) ** 2) / len(y test)
          test mse[dataset] = validation mse
 end time = time.time()
  runtime[dataset] = end_time - start_time
print("Task 2")
# Report results for all datasets
for dataset in datasets:
 print(f"Dataset: {dataset}")
  print(f"Selected \( \lambda \): {best lambda[dataset]}")
  print(f"Validation MSE with selected λ:
{best avg validation mse[dataset]}")
 print(f"Test MSE with selected \( \lambda: \) { test mse[dataset] }")
 print(f"Runtime: {runtime[dataset]} seconds")
print("\n")
print("\nTask 3")
# Task 3: Bayesian Model Selection
for i, dataset in enumerate(datasets):
  train_data = pd.read_csv('pp2data/train-'+dataset+'.csv', header=None)
 train labels = pd.read csv('pp2data/trainR-'+dataset+'.csv', header=None)
 test data = pd.read csv('pp2data/test-'+dataset+'.csv', header=None)
test_labels = pd.read_csv('pp2data/testR-'+dataset+'.csv', header=None)
 X train = train data.values
```

```
y_train = train_labels.values.ravel()
  X test = test data.values
 y_test = test labels.values.ravel()
  # Initialize alpha (\alpha) and beta (\beta) to random values in the range [1, 10]
  alpha = np.random.uniform(1, 10)
beta = np.random.uniform(1, 10)
 # Define convergence threshold
  threshold = 0.0001
  # Initialize variables to store results
  effective lambda = None
 mse = None
start time = time.time()
 N = X train.shape[0]
 M = X train.shape[1]
 # Iterative algorithm to select \alpha and \beta
 while True:
      lambda values = np.linalg.eigvals(beta * np.dot(X train.T, X train))
      # Calculate SN and mN using equations (3.53) and (3.54)
      SN = np.linalg.inv((alpha * np.identity(M)) + (beta * np.dot(X_train.T,
X train)))
      mN = beta * np.dot(np.dot(SN, X train.T), y train)
  # Update alpha and beta using equations (3.91) and (3.92)
  gamma = np.sum(lambda values / (alpha + lambda values))
 alpha_new = gamma / np.dot(mN.T, mN)
     y_pred = np.dot(X_train, mN)
      beta new = (N - gamma) / (np.sum((y train - y pred) ** 2))
     # Check for convergence
     if abs(alpha new - alpha) < threshold and abs(beta new - beta) <</pre>
threshold:
   break
      alpha = alpha new
    beta = beta new
 # Calculate effective \lambda as \alpha/\beta
effective_lambda = alpha / beta
 mN = beta * np.dot(np.dot(SN, X_train.T), y_train)
```

```
# Use mN for prediction on the test set
y_test_pred = np.dot(X_test, mN)

# Calculate test set MSE
mse = np.sum((y_test - y_test_pred) ** 2) / len(y_test)

end_time = time.time()

# Report results
print(f"Dataset: {dataset}")
print(f"Selected α: {alpha}")
print(f"Selected β: {beta}")
print(f"Effective λ (α/β): {effective_lambda}")
print(f"Test MSE with selected α and β: {mse}")
print(f"Runtime: {end_time - start_time} seconds")
print('\n')
```

README

Just run the PP2.py file to get the 4 plots for the 4 dataset as per Task 1. For further drilldown,

I also generated a table in the terminal with the lambda that generates the lowest MSE for each dataset

For Task 2 and 3 to run, just close the window with the plots for the code to go to the Task 2/3 section.

These sections will also print out the necessary information in the terminal as per the assignment's instructions.