hw2_svm_impl

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[]: !pip install ucimlrepo
[]: from ucimlrepo import fetch_ucirepo
     import pandas as pd
     import numpy as np
     import cvxpy as cp
     from matplotlib import pyplot as plt
[ ]: # !! DO NOT MODIFY THIS CELL !!
     # Download and preprocess the dataset.
     # fetch dataset
     heart_disease = fetch_ucirepo(id=45)
     X = heart_disease.data.features
     # Convert categorical features into one-hot encode
     categorical_features = ['cp', 'thal', 'slope', 'restecg']
     X = pd.get_dummies(X, columns=categorical_features)
     y = heart_disease.data.targets
     print(f"Number of samples in all full dataset is: {len(X)}.")
     # Check if our train set has missing value
     na_in_features = X.isna().any(axis=1).sum()
     na_in_trainY = y.isna().sum()
     print(f"Number of rows with missing values in features: {na_in_features}")
     # Drop the rows with missing values.
     indices_with_nan = X.index[X.isna().any(axis=1)]
     X = X.drop(indices_with_nan)
     y = y.drop(indices_with_nan)
     # Divide train/test
     np.random.seed(6464)
     msk = np.random.rand(len(X)) < 0.75</pre>
     X train = X[msk]
     X_{\text{test}} = X[\text{-msk}]
     y_train = y[msk]
     y_{test} = y[~msk]
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# Convert problem to binary problem
     X_train = np.array(X_train,dtype='float')
     X_test = np.array(X_test,dtype='float')
     y_train = np.array([-1 if i==0 else 1 for i in y_train.values],dtype='float')
     y_test = np.array([-1 if i==0 else 1 for i in y_test.values],dtype='float')
     print(f"Shapes: X_train: {X_train.shape}, y_train: {y_train.shape}, X_test:
      →{X_test.shape}, y_test: {y_test.shape}")
[]: # Normalize X train and X test using the statistics of X train.
     \# 1. Compute the mean and standard deviation for each feature in X_{\_} train
     # 2. Subtract the mean from each feature and divide by the standard deviation
         for both X_t train and X_t test.
     YOUR CODE HERE!
[]: # Print the mean and standard deviation of the first and last feature.
     YOUR CODE HERE!
\# Complete the `trainSVM` function to find the optimal w and b that minimize
     # the primal SVM objective given in the write-up.
     # The function takes three inputs:
     # - trainX: the normalized train features with shape (#train samples, #features)
     # - trainY: train labels with shape (#train_samples,)
     # - C: C parameter of the minimization problem
     # The function should return a three-tuple with:
     # - w: the weight vector with shape (#features,)
     # - b: the bias. A scalar with shape (1,)
     # - xi: the slack variables with shape (#train_samples,)
     # You can use cvxpy that we imported as cp
     # You may find cp. Variable, cp. Minimize, cp. Problem useful
     # For the problem solver, prefer the default, cp.CLARABEL
     def trainSVM(trainX, trainY, C):
        YOUR CODE HERE!
[]: \# Solve SVM with C = 1 and print the first three weights, b and the first
     # three slack variables as instructed in the write-up
     YOUR CODE HERE!
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[]: \# Solve SVM with C = 0 and print the first three weights, b and the first
     # three slack variables as instructed in the write-up
     YOUR CODE HERE!
[ ]:  # Eval SVM
     \# Write a function to evaluate the SVM model given its `w` and `b` parameters
     # on evaluation data `X_eval` and true labels `y_eval`.
     # 1. Estimate the labels of `X_eval`.
     # 2. Return the ratio of accurately estimated labels by comparing with_
     \Rightarrow 'y_eval'.
     def evalSVM(X_eval, y_eval, w, b):
         YOUR CODE HERE!
[]: train_accuracies = []
     test accuracies = []
     C_values = []
     # For each C value given in the homework, find optimal w, b
     # values using the normalized train set. calculate the accuracy
     # on train and test sets using found w and b.
     # Save those values as we will plot them
     YOUR CODE HERE!
[]: # Plotting and reporting the desired values
     YOUR CODE HERE!
```