(PR)
$$\underset{\omega \in \mathbb{R}^{4}}{\text{min}} \frac{1}{N} \underset{\varepsilon \in \mathbb{R}^{4}}{\overset{\mathcal{L}}{\longrightarrow}} (\underset{\varepsilon \in \mathbb{R}^{4}}{\text{min}} \underset{\varepsilon \in \mathbb{R}^{4}}{\text{min}} \frac{1}{N} \underset{\varepsilon \in \mathbb{R}^{4}}{\overset{\mathcal{L}}{\longrightarrow}} (\underset{\varepsilon \in \mathbb{R}^{4}}{\text{min}} \frac{1}{N} \underset{\varepsilon \in \mathbb{R}^{4}}{\overset{\mathcal{L}}{\longrightarrow}} (\underset{\varepsilon \in \mathbb{R}^{4}}{\text{min}} \frac{1}{N} \underset{\varepsilon \in \mathbb{R}^{4}}{\text{min}} \frac{1}{N} \underset{\varepsilon$$

= (Pv) min $\frac{1}{N}$ $\frac{N}{N}$ \frac

i. optimal ω^* obtained by (Pv) is the same as the optimal solution (= $\omega_{REG} = (X^TX + \lambda x) X^T y$) obtained by solving (PR)

$$\widetilde{E}_{aug}(\underline{\omega}) = \widetilde{E}_{in}(\underline{\omega}) + \frac{\lambda}{N} ||\underline{\omega}||^{2}$$

$$= E_{in}(\underline{\omega}^{*}) + \frac{1}{2}(\underline{\omega} - \underline{\omega}^{*})^{T} \underline{H}(\underline{\omega} - \underline{\omega}^{*}) + \frac{\lambda}{N} ||\underline{\omega}||^{2}$$

$$= E_{in}(\underline{\omega}^{*}) + \frac{1}{2}(\underline{\omega}^{T} \underline{H} \underline{\omega} - \underline{\omega}^{T} \underline{H} \underline{\omega}^{*} - \underline{\omega}^{*} \underline{H} \underline{\omega}^{*}) + \frac{\lambda}{N} \underline{\omega}^{T} \underline{\omega}$$

$$\nabla \widetilde{E}_{aug}(\underline{\omega}) = \nabla E_{in}(\underline{\omega}^{*}) + \frac{1}{2}(2\underline{H} \underline{\omega} - \underline{H} \underline{\omega}^{*} - \underline{H} \underline{\omega}^{*}) + \frac{2\lambda}{N} \underline{\omega}$$

$$= \underbrace{H(\omega - \underline{\omega}^*)}_{N} + \underbrace{\frac{2\lambda}{N}}_{N} \underline{\omega}$$

Let want is the minimizer of \widetilde{E} ang(ω)

$$\nabla \widehat{E}_{aug}(\underline{W}_{aug}) = \underline{O}$$

$$\Rightarrow \underline{H}(\underline{\omega}_{aug} - \underline{\omega}^{*}) + \frac{2\lambda}{N} \underline{\omega}_{aug} = \underline{0}$$

$$\Rightarrow (\underline{H} + \frac{2\lambda}{N} \underline{I}) \underline{\omega_{aug}} = \underline{H} \underline{\omega}^* - \underline{H} + \frac{2\lambda}{N} \underline{I} = 15 \text{ positive definite}$$

$$\Rightarrow \underline{\omega_{aug}} = (\underline{H} + \frac{2\lambda}{N} \underline{I}) \underline{H} \underline{\omega}^* = \underline{H} + \frac{2\lambda}{N} \underline{I} = 15 \text{ positive definite}$$

$$E\left(\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}-0)^{2}\right) = 6^{2}$$

$$\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}-y_{n}^{2}) = 6^{2}$$

$$\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}^{2}-2y_{n}y_{n}+y_{n}^{2})$$

$$=\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}^{2}-2y_{n}y_{n}+y_{n}^{2})$$

$$=\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}^{2}-2y_{n}y_{n}+y_{n}^{2})$$

$$=\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}^{2}+y_{n}^{2}+y_{n}^{2}+y_{n}^{2})$$

$$=\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}^{2}+y_{n}^{2}+y_{n}^{2}+y_{n}^{2}+y_{n}^{2}+y_{n}^{2})$$

$$=\frac{1}{K},\frac{N}{N^{2}N^{2}K^{2}},(y_{n}^{2}+y_{n}$$

 $= () + \frac{2N - K}{K(N - K)}) 6^{2}$

$$\begin{aligned} \left(\text{Var} \left(\overrightarrow{y} \right) &= \text{Var} \left(\frac{1}{N-K} \frac{N-K}{N-K} y_n \right) \\ &= \left(\frac{1}{N-K} \right)^2 \text{Var} \left(\frac{\Sigma}{N-K} y_n \right) \\ &= \left(\frac{1}{N-K} \right)^2 \frac{N-K}{N-K} \text{Var} \left(y_n \right) \\ &= \left(\frac{1}{N-K} \right)^2 \left(N-K \right) 6^2 \\ &= \frac{1}{N-K} 6^2 \end{aligned}$$

 $\sum_{n=1}^{N} y_n = 0$

> 27 yn + 27 yn = 0

 $\Rightarrow \frac{N}{57} yn = - \sum_{h=1}^{N-K} yn$

=-(N-K)7

no collaborators

 $Ein(\omega^*) = \frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} \sum_{n=1}^{N} (n - y_n)^{\frac{1}{N}})$ $= (\frac{1}{N}) \left[\frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} - Ny_n)^{\frac{1}{N}} \right]$ $= \frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} - Ny_n)^{\frac{1}{N}}$ $= \frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} - Ny_n)^{\frac{1}{N}}$ $= \frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} - \frac{1}{N} y_i - y_n)^{\frac{1}{N}}$ $= (\frac{1}{N}) \left[\frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} y_i - y_n) - y_n \right]$ $= (\frac{1}{N}) \left[\frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} y_i - y_n) - y_n \right]$ $= (\frac{1}{N}) \left[\frac{1}{N} \sum_{n=1}^{N} (\frac{1}{N} y_i - Ny_n) \right]$ $= (\frac{1}{N}) Ein(\omega^*) \quad \text{for } N \ge 2$ Q. E. V.

P17 of lecture 15

no collaborators East (g) = Eout(gc) $\Rightarrow pe_{-} + (1-p)e_{+} = p$ $\Rightarrow p(e_{-} - e_{+} - 1) = -e_{+}$ $\Rightarrow p = \frac{e_{+}}{1+e_{+} - e_{-}}$



Histogram of $E_{out}(g)$ Histogram of Non-Zero Components in g 200 175 175 150 150 ਨੂੰ 125 100 100 75 50 50 25 25 520 Eout (%) Number of Non-Zero Components

collaborators:

B11201009 黄勤元 B11901073 林禹融

```
loupy ...
import os
import requests
import bz2
import numpy as np
import scipy.sparse
from liblinear.liblinearutil import *
import matplotlib.pyplot as plt
# Step 1: Download and decompress the data
> def download_and_extract(url, dest_path):
> def decompress bz2(file path, output path):-
   train_url = "https://www.csie.ntu.edu.tw/-cjlin/libsymtools/datasets/multiclass/mnist.scale.bz2"
test_url = "https://www.csie.ntu.edu.tw/-cjlin/libsymtools/datasets/multiclass/mnist.scale.t.bz2"
train file compressed = "mnist.scale.bz2"
train file = "mnist.scale.t.bz2"
train file = "mnist.scale.t"
    download_and_extract(train_url, train_file_compressed) download_and_extract(test_url, test_file_compressed) decompress_bz2(train_file_compressed, train_file) decompress_bz2(test_file_compressed, test_file)
    # Step 2: Load and preprocess the data
y_train, X_train = svm_read_problem(train_file, return_scipy=True)
y_test, X_test = svm_read_problem(test_file, return_scipy=True)
  # Filter for classes 2 and 6, and relabel them for binary classification
train mask = np.isin(y_train, [2, 6])
test_mask = np.isin(y_test, [2, 6])
y_train, X_train = y_train[train_mask], X_train[train_mask]
y_test, X_test = y_test[test_mask], X_test[test_mask]
y_train = np.where(y_train == 2, 1, -1)
y_test = np.where(y_test == 2, 1, -1)
  \label{eq:continuous} \begin{tabular}{ll} \# \ Align feature dimensions between training and test sets $$X_train, $X_test$ = align_features($X_train, $X_test$)$ \\ \end{tabular}
   # Scale the data using csr_scale
scale_param = csr_find_scale_param(X_train, lower=0)
X_train = csr_scale(X_train, scale_param)
X_test = csr_scale(X_test, scale_param)
  # Step 3: Define helper
def lambda_to_C(lmbda):
  lambdas = [0.01, 0.1, 1, 10, 100, 1000]
N = len(y_train)
best_lambda, min_error = None, float('inf')
errors_in = []
   \# Step 4: Find the best lambda and repeat experiments with 1126 seeds random_seeds = range(1126) errors_out, non_zero_counts = [], []
  for seed in random_seeds:
    for lmbda in lambdas:
        C = lambda to C(lmbda)
        model = train(y_train, X_train, f'·s 6 -c {C} -B 1 -q')
        _, p_acc, _ = predictly_train, X_train, model)
        E_in = 100 - p_acc[0]
        errors_in.append(E_in)
        if E_in < min_error_or (E_in == min_error_and lmbda > best_lambda):
            best_lambda, min_error = lmbda, E_in
               L = (amboa_to_(lest_lamboa)
np.random.seed(seed)
model = train(y_train, X_train, f'-s 6 -c {C} -B 1 -q')
_, p_acc, _ = predict(y_test, X_test, model)
E_out = 100 - p_acc(0) # accuracy to error
errors_out.append(E_out)
                # Access weights and bias terms
weights, bias = model.get_decfun()
non_zero_counts.append(np.sum(np.abs(weights) > 1e-6))
  # Step 6: Plot histograms
plt.figure(figsize=(10, 5))
plt.subplot(1, 2, 1)
plt.hist(errors_out, bins=30, color="blue", edgecolor="black", alpha=0.7)
plt.title("Histogram of $E {out}(g)$")
plt.xlabel("$E {out}$' ($\$")
plt.ylabel("Frequency")
  plt.subplot(1, 2, 2)
plt.hist(non_zero_counts, bins=30, color="green", edgecolor="black", alpha=0.7)
plt.title("Histogram of Non-Zero Components in $g$")
plt.xlabel("Mumber of Non-Zero Components")
plt.ylabel("Frequency")
```

Histogram of E_{out}

200

150

50

1.2

1.3

1.4

1.5

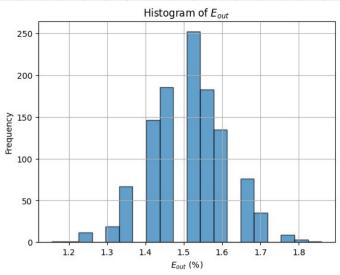
E_{out} (%)

11. collaborator: B[1901073 林禹融

Both histograms are center around East ≈ 1.5%, suggesting that performance is similar between both methods. While the distribution in Problem 11 is slightly wider than that in Problem 10, reflecting that cross-validation approach introdues additional variability because of the random splits into sub-training and validation sets.

```
ilpy > ...
import os
import requests
import bz2
import numpy as np
from liblinear.liblinearutil import *
from sklearn.model_selection import train_test_split
import maplotlib.pyplot as plt
from scipy.sparse import csr_matrix, hstack
  # Constants
LAMBDA VALUES = [0.01, 0.1, 1, 10, 100, 1000]
N EXPERIMENTS = 1126
TRAIN_SIZE = 8000
  # Step 1: Download and decompress the dat
def download_and_extract(url, dest_path):
 def decompress bz2(file path, output path):
train_url = "https://www.csie.ntu.edu.tw/-cjlin/libsvmtools/datasets/multiclass/mnist.scale.bz2"
test_url = "https://www.csie.ntu.edu.tw/-cjlin/libsvmtools/datasets/multiclass/mnist.scale.t.bz2"
train_file_compressed = "mnist.scale.bz2"
test_file_compressed = "mnist.scale.t.bz2"
train_file = "mnist.scale"
test_file = "mnist.scale."
  download_and_extract(train_url, train_file_compressed)
download and extract(test url, test file compressed)
  decompress bz2(train_file_compressed, train_file)
decompress_bz2(test_file_compressed, test_file)
  # Step 2: Load and preprocess the data
y_train, X_train = svm_read_problem(train_file, return_scipy=True)
y_test, X_test = svm_read_problem(test_file, return_scipy=True)
 # Filter for classes 2 and 6, and relabel them for binary classification
train mask = np.isin(y_train, [2, 6])
test_mask = np.isin(y_test, [2, 6])
y_train, X_train = y_train[train mask], X_train[train mask]
y_test, X_test = y_test[test_mask], X_test[test_mask]
y_train = np.where(y_train == 2, 1, -1)
y_test = np.where(y_test == 2, 1, -1)
def align features(train, test): --
  # Align feature dimensions between training and test sets
X_train, X_test = align_features(X_train, X_test)
  # Scale the data using csr_scale
scale_param = csr_find_scale_param(X_train, lower=0)
  X train = csr scale(X train, scale param)
X test = csr scale(X test, scale param)
  def calculate_error(y_true, y_pred):
return np.mean(y_true != y_pred) * 100
 # Prepare for histogram
E_out_values = []
  for experiment in range(N_EXPERIMENTS):
    np.random.seed(experiment)
    # Sten 1: Solit data into sub-train
             np.random.seed(experiment)
# Step 1: Split data into sub-training and validation sets
X_subtrain, X_val, y_subtrain, y_val = train_test_split(
X_train, y_train, train_size=TRAIN_SIZE, random_state=experiment
            # Step 2: Evaluate each lambda
for lambda_value in LAMBOA_VALUES:
    C = 1 / lambda_value
    model = train(y_subtrain, X_subtrain, f'-s 6 -c {C} -B 1 -q')
    p_val, __ = predict(y_val, X_val, model, '-q')
    E_val = calculate_error(y_val, p_val)
    if E_val < best_E_val or (E_val == best_E_val and lambda_value > best_lambda):
        best_E_val = lambda_value
             # Step 3: Re-train with the best lambda on the full training set
C = 1 / best_lambda
final_model = train(y_train, X_train, f'-s 6 -c {C} -B 1 -q')
            # Step 4: Evaluate E_out on the test set
p_test, _, _ = predict(y_test, X_test, final_model, '-q')
E_out = calculate error(y_test, p_test)
E_out_values.append(E_out)
         step 5: Plot histogram of E out
t.hist(E_out_values, bins=20, edgecolor='black', alpha=0.7)
t.title('Histogram of $E_(out)$')
t.xlabel('$E_(out)$ (%)')
t.xlabel('Frequency')
```

12. Collaborator: B11901073 林禹就 Both histograms are center



Both histograms are center around East ≈ 1.5%, suggesting that performance is similar between both methods. Additionally, histogram in Problem 12 is more concentrated around the mean. The 3-fold cross validation may be generally a better approach to minize variance in model selection process

```
import numpy as np
from liblinear.liblinearutil import *
from sklearn.model_selection import KFold
import matplotlib.pyplot as plt
from scipy.sparse import csr_matrix, hstack
 # Constants
LAMBDA_VALUES = [0.01, 0.1, 1, 10, 100, 1000]
 N_EXPERIMENTS = 1126
K_FOLDS = 3 # 3-fold cross-validation
 # Step 1: Download and decompress the data
def download_and_extract(url, dest_path):-
 train_url = "https://www.csie.ntu.edu.tw/-cjlin/libsvmtools/datasets/multiclass/mnist.scale.bz2"
test_url = "https://www.csie.ntu.edu.tw/-cjlin/libsvmtools/datasets/multiclass/mnist.scale.t.bz2"
train_file_compressed = "mnist.scale.bz2"
test_file_compressed = "mnist.scale.t.bz2"
train_file = "mnist.scale"
test_file = "mnist.scale.t"
 download_and_extract(train_url, train_file_compressed)
download and extract(test_url, test_file_compressed)
decompress_bz2(train_file_compressed, train_file)
decompress_bz2(test_file_compressed, train_file)
 # Step 2: Load and preprocess the data
y_train, X_train = svm_read_problem(train_file, return_scipy=True)
y_test, X_test = svm_read_problem(test_file, return_scipy=True)
 # Filter for classes 2 and 6, and relabel them for binary classification train_mask = np.isin(y_train, [2, 6])
test_mask = np.isin(y_test, [2, 6])
y_train, X_train = y_train[train_mask], X_train[train_mask]
y_train = y_train[train_mask], X_test[test_mask]
y_test, X_test = y_test[test_mask], X_test[test_mask]
y_train = np.where(y_train = 2, 1, -1)
y_test = np.where(y_test == 2, 1, -1)
 # Scale the data using csr_scale
scale_param = csr_find_scale_param(X_train, lower=0)
X_train = csr_scale(X_train, scale_param)
X_test = csr_scale(X_test, scale_param)
# Function to calculate error
def calculate_error(y_true, y_pred):-
# Function for 3-fold cross-validation
def cross_validate(X, y, lambda_value):
    kf = KFold(n_splits=K_FOLDS, shuffle=True, random_state=None)
    total_error = 0
           for train_index, val_index in kf.split(X):
    X_train_fold, X_val_fold = X[train_index], X[val_index]
    y_train_fold, y_val_fold = y[train_index], y[val_index]
                      C = 1 / lambda_value
model = train(y_train_fold, X_train_fold, f'-s 6 -c {C} -B 1 -q')
                      p_val, _, _ = predict(y_val_fold, X_val_fold, model, '-q')
total_error += calculate_error(y_val_fold, p_val)
          # Return average cross-validation error
return total_error / K_FOLDS
 # Main loop for 1126 experiments
for experiment in range(N_EXPERIMENTS):
    np.random.seed(experiment)
           best lambda, best E CV = None, float('inf')
            # Step 1: Perform 3-fold cross-validation for each lambda
for lambda_value in LAMBDA_VALUES:
    E_CV = cross_validate(X_train, y_train, lambda_value)
                     # Update best lambda
if E_CV < best_E_CV or (E_CV == best_E_CV and lambda_value > best_lambda):
    best_E_CV = E_CV
    best_lambda = lambda_value
            # Step 2: Re-train with the best lambda on the full training set
C = 1 / best_lambda
final_model = train(y_train, X_train, f'-s 6 -c {C} -B 1 -q')
```