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# File: project 1.py
import numpy as np
import pandas as pd
from tqdm import tqdm
import time
import joblib
from sklearn.datasets import fetch openml
from sklearn.model selection import train test split
from sklearn.utils import check random state
from sklearn.metrics import accuracy score
from sklearn.preprocessing import StandardScaler
from sklearn.cluster import DBSCAN
from sklearn.linear model import LogisticRegression
from sklearn.neighbors import KNeighborsClassifier
SEED = 42
# Load or fetch the MNIST dataset with a progress bar
def load mnist():
    try:
        # Try to load the dataset from the local file if it exists
        mnist = joblib.load('mnist dataset.joblib')
    except FileNotFoundError:
        # If the file doesn't exist, fetch the dataset from OpenML
        print('Downloading MNIST...')
        mnist = fetch_openml(name='mnist_784', version=1, cache=True,
parser='auto')
        # Save the dataset locally
        joblib.dump(mnist, 'mnist_dataset.joblib')
    return mnist
# Load the MNIST dataset
mnist = load mnist()
# Split the data into features and labels
X, y = mnist.data, mnist.target.astype(int)
# Standardize the features
scaler = StandardScaler()
X standardized = scaler.fit transform(X)
# Split the dataset into training and test sets
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X_train, X_test, y_train, y_test = train_test_split(X_standardized, y,
test size=0.2, random state=SEED)
# Prototype Selection Methods
def random sampling(X train, y train, M):
    Randomly select M samples from the training set.
    Parameters:
    - X train: Features of the training set (NumPy array or Pandas DataFrame)
    - y train: Labels of the training set (NumPy array or Pandas DataFrame)
    - M: Number of prototypes to select
    Returns:
    selected_prototypes: Randomly selected prototypes
    # Convert Pandas DataFrame to NumPy array if needed
    X_train = X_train.values if isinstance(X_train, pd.DataFrame) else X_train
    y_train = y_train.values if isinstance(y_train, pd.Series) else y_train
    # Convert y train to a NumPy array
    y train np = np.array(y train)
    # Get the indices of randomly selected prototypes
    selected_indices = np.random.choice(len(X_train), size=M, replace=False)
    # Extract the corresponding samples
    selected prototypes = X train[selected indices]
    selected_labels = y_train_np[selected_indices]
    return selected prototypes, selected labels
def dbscan_prototype_selection(X_train, y_train, M, eps=20, min_samples=10,
random state=None):
    Use DBSCAN for prototype selection.
   Parameters:
    - X train: Features of the training set (NumPy array or Pandas DataFrame)
    - y_train: Labels of the training set (NumPy array or Pandas DataFrame)
    - M: Number of prototypes to select
    - eps: The maximum distance between two samples for one to be considered in
the neighborhood of the other
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- min samples: The number of samples (or total weight) in a neighborhood for
a point to be considered as a core point
    - random state: Seed for reproducibility
   Returns:
    - selected prototypes: Prototypes selected using DBSCAN
    # Convert Pandas DataFrame to NumPy array if needed
   X train = X train.values if isinstance(X train, pd.DataFrame) else X train
   y_train = y_train.values if isinstance(y_train, pd.Series) else y_train
   # Apply DBSCAN for clustering
   dbscan = DBSCAN(eps=eps, min_samples=min_samples)
    labels = dbscan.fit predict(X train)
    # Identify core points (excluding outliers)
    core points indices = np.where(labels != -1)[0]
    if len(core points indices) == 0:
        raise ValueError("No core points found. Adjust parameters (eps,
min samples) or provide more data.")
    # Randomly select M prototypes from core points using random state
    random state = check random state(random state)
    selected indices = random state.choice(core points indices, size=min(M,
len(core points indices)), replace=False)
    # Extract the corresponding samples using NumPy array indexing
    selected prototypes = X train[selected indices, :]
    selected_labels = y_train[selected_indices]
    return selected prototypes, selected labels
def active learning prototype selection(X train, y train, M, max iter=1000,
random state=None):
   Use Active Learning for prototype selection.
   Parameters:
    - X train: Features of the training set (NumPy array or Pandas DataFrame)
   - y train: Labels of the training set (NumPy array or Pandas DataFrame)
    - M: Number of prototypes to select
    - max iter: Maximum number of iterations for Logistic Regression
    - model_filename: File name for saving/loading the model
    - random state: Seed for reproducibility
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Returns:
    - selected prototypes: Prototypes selected using Active Learning
    # Convert Pandas DataFrame to NumPy array if needed
    X_train = X_train.values if isinstance(X_train, pd.DataFrame) else X_train
    y train = y train.values if isinstance(y train, pd.Series) else y train
    # Train a new classifier (don't save the classifier because we need
    # to record the training time for the report)
    classifier = LogisticRegression(max iter=max iter,
random state=check_random_state(random_state))
    classifier.fit(X_train, y_train)
    # Get predicted probabilities for each class
    predicted probs = classifier.predict proba(X train)
    # Calculate uncertainty as the maximum probability across classes
    uncertainty = 1 - np.max(predicted probs, axis=1)
    # Select M prototypes with the highest uncertainty
    selected_indices = np.argsort(uncertainty)[-M:]
    # Extract the corresponding samples
    selected_prototypes = X_train[selected_indices]
    selected_labels = y_train[selected_indices]
    return selected prototypes, selected labels
# Code to run experiments
# Define the range of M values to experiment with
M values = [100, 500, 1000, 5000, 10000]
# Define the prototype selection methods
prototype_methods = [
    # ("Random Sampling", random_sampling),
    # ("DBSCAN", dbscan_prototype_selection),
    ("Active Learning", active_learning_prototype_selection)
# Create a 1-KNN classifier
classifier = KNeighborsClassifier(n_neighbors=1)
# Initialize the results dictionary with empty lists for each metric
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results = {
    "Method": [],
    "M": [],
    "Mean Accuracy": [],
    "Std Accuracy": [],
    "Mean Training Time (s)": [],
    "Std Training Time (s)": [],
    "Mean Test Time (s)": [],
    "Std Test Time (s)": []
# Number of experiments to run for each method and M value
num_experiments = 10
# Create a dictionary to store results for each experiment
experiment results = []
# Run experiments
for method name, method func in prototype methods:
    print(f"\nRunning experiments for {method_name}...\n")
    for M in M values:
        print(f"Number of Prototypes (M): {M}")
        # Lists to store results for each experiment
        accuracies = []
        training_times = []
        test_times = []
        for _ in range(num_experiments):
            # Timer for training time
            start_time = time.time()
            # Prototype selection
            prototypes, labels = method func(X train, y train, M)
            # Train KNN classifier on the selected prototypes
            classifier.fit(prototypes, labels)
            # Record training time
            training_times.append(time.time() - start_time)
            # Timer for test time
            start time = time.time()
            # Evaluate on the test set
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y pred = classifier.predict(X test)
            # Record test time
            test times.append(time.time() - start time)
            # Compute accuracy
            accuracies.append(accuracy score(y test, y pred))
        # Compute statistics across experiments
        mean_accuracy = np.mean(accuracies)
        std accuracy = np.std(accuracies)
        mean training time = np.mean(training times)
        std_training_time = np.std(training_times)
        mean test time = np.mean(test times)
        std_test_time = np.std(test_times)
        # Record results
        results["Method"].append(method_name)
        results["M"].append(M)
        results["Mean Accuracy"].append(mean_accuracy)
        results["Std Accuracy"].append(std_accuracy)
        results["Mean Training Time (s)"].append(mean_training_time)
        results["Std Training Time (s)"].append(std_training_time)
        results["Mean Test Time (s)"].append(mean_test_time)
        results["Std Test Time (s)"].append(std_test_time)
        print(f"Mean Accuracy: {mean accuracy:.4f} (±{std accuracy:.4f})")
        print(f"Mean Training Time: {mean_training_time:.4f} seconds
(±{std training time:.4f})")
        print(f"Mean Test Time: {mean_test_time:.4f} seconds
(±{std test time:.4f})\n")
# Convert results to a DataFrame for easier analysis and visualization
results df = pd.DataFrame(results)
# Save results to a CSV file
results_df.to_csv("prototype_selection_results.csv", index=False)
# Display the results
print(results df)
print("Results saved to prototype selection results.csv")
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# File: visualization.py
import matplotlib.pyplot as plt
import seaborn as sns
import pandas as pd
# Load the results from the CSV file
results df = pd.read csv("results/prototype selection results.csv")
# Visualize Accuracy vs. Number of Prototypes (M)
plt.figure(figsize=(10, 6))
for method in results df['Method'].unique():
    method data = results df[results df['Method'] == method]
    plt.plot(method_data["M"], method_data["Mean Accuracy"], marker="o",
label=method)
    plt.fill_between(method_data["M"],
                     method data["Mean Accuracy"] - method data["Std Accuracy"],
                     method_data["Mean Accuracy"] + method_data["Std Accuracy"],
                     alpha=0.2)
plt.title("Accuracy vs. Number of Prototypes")
plt.xlabel("Number of Prototypes (M)")
plt.ylabel("Mean Accuracy")
plt.grid(True)
plt.legend()
plt.savefig("accuracy vs prototypes.png") # Save the figure
plt.show()
# Visualize Training Time vs. Number of Prototypes (M)
plt.figure(figsize=(10, 6))
for method in results df['Method'].unique():
    method data = results df[results df['Method'] == method]
    plt.plot(method_data["M"], method_data["Mean Training Time (s)"], marker="o",
label=method)
    plt.fill between(method data["M"],
                     method data["Mean Training Time (s)"] - method data["Std
Training Time (s)"],
                     method data["Mean Training Time (s)"] + method data["Std
Training Time (s)"],
                     alpha=0.2)
plt.title("Training Time vs. Number of Prototypes")
plt.xlabel("Number of Prototypes (M)")
plt.ylabel("Mean Training Time (s)")
plt.grid(True)
plt.legend()
plt.savefig("training_time_vs_prototypes.png") # Save the figure
plt.show()
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# Visualize Test Time vs. Number of Prototypes (M)
plt.figure(figsize=(10, 6))
for method in results df['Method'].unique():
    method_data = results_df[results_df['Method'] == method]
    plt.plot(method_data["M"], method_data["Mean Test Time (s)"], marker="o",
label=method)
    plt.fill_between(method_data["M"],
                     method data["Mean Test Time (s)"] - method data["Std Test
Time (s)"],
                     method_data["Mean Test Time (s)"] + method_data["Std Test
Time (s)"],
                     alpha=0.2)
plt.title("Test Time vs. Number of Prototypes")
plt.xlabel("Number of Prototypes (M)")
plt.ylabel("Mean Test Time (s)")
plt.grid(True)
plt.legend()
plt.savefig("test time vs prototypes.png") # Save the figure
plt.show()
# Table of Mean Accuracy and Mean Training Time for each Method and M
summary_table = results_df.pivot_table(index="M", columns="Method", values=["Mean
Accuracy", "Mean Training Time (s)", "Mean Test Time (s)"])
summary_table.to_csv("summary_table.csv")
print("Summary Table:")
print(summary table)
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