## Intro to ML - Homework 4

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Github: https://github.com/QueenSophiaLo/Intro-To-ML/tree/main/Homework%204

## Problem 1

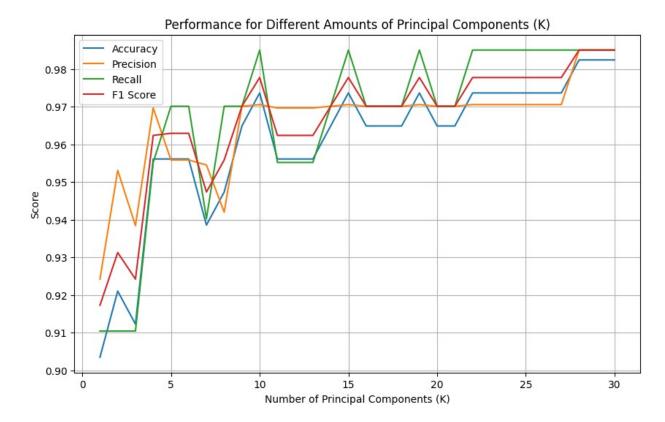
Use the cancer dataset to build an SVM classifier to classify the type of cancer (Malignant vs. benign)

```
import numpy as np
from sklearn.datasets import load_breast_cancer
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
from scipy import stats
from sklearn.svm import SVC
from sklearn.metrics import accuracy score, precision score,
recall score, f1 score
# Load dataset
data = load breast cancer()
X = data.data
v = data.target
# Split into training and test sets (80/20)
X train, X test, y train, y test = train test split(
    X, y, train_size=0.8, test_size=0.2, random_state=0
# Standardize features
scaler = StandardScaler()
X_train = scaler.fit_transform(X train)
X test = scaler.transform(X test)
# PCA (feature extraction)
pca = PCA()
# metrics to plot for different values of K
accuracy_list = []
precision_list = []
recall list = []
f1 list = [] # F1 Score is the weighted average of Precision and
Recall
k value = list(range(1, X train.shape[1] + 1)) # the number of
features, K ranges 1-30
```

```
# Evaluate SVM performance for each number of PCA components
for k in k value:
    # Reduce data dimensionality using PCA
    pca = PCA(n components=k)
    X train pca = pca.fit transform(X train)
    X test pca = pca.transform(X test)
    # Fit SVM with a linear kernel on the reduced data
    svm_model = SVC(kernel='linear', random_state=42)
    svm model.fit(X_train_pca, y_train)
    # Make predictions on the test set
    predictions = svm model.predict(X test pca)
    # Compute evaluation metrics
    acc score = accuracy score(y test, predictions)
    prec score = precision_score(y_test, predictions)
    rec score = recall score(y test, predictions)
    f1 score val = f1 score(y test, predictions)
    # Save metrics for later plotting
    accuracy list.append(acc score)
    precision list.append(prec score)
    recall list.append(rec score)
    f1 list.append(f1 score val)
# Find the optimal number of K components
best k = k value[np.argmax(accuracy list)]
print(f"The optimal number of principal components (K): {best k}")
print(f"Highest accuracy K:{best k}: {np.max(accuracy list)}")
The optimal number of principal components (K): 28
Highest accuracy K:28: 0.9824561403508771
```

Plot your classification accuracy, precision, and recall.

```
# Plot over different values of K
import matplotlib.pyplot as plt
plt.figure(figsize=(10, 6))
plt.plot(k_value, accuracy_list, label='Accuracy')
plt.plot(k_value, precision_list, label='Precision')
plt.plot(k_value, recall_list, label='Recall')
plt.plot(k_value, f1_list, label='F1 Score')
plt.xlabel('Number of Principal Components (K)')
plt.ylabel('Score')
plt.title('Performance for Different Amounts of Principal Components
(K)')
plt.legend()
plt.grid(True)
plt.show()
```

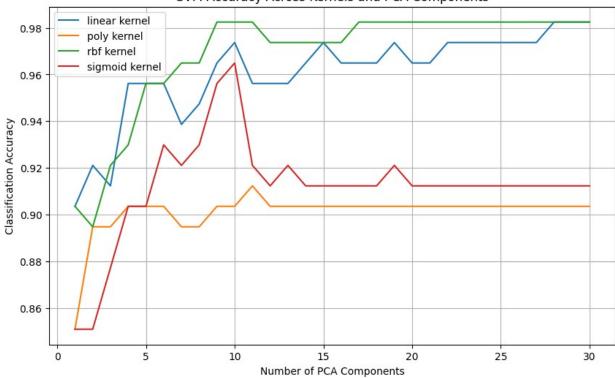


Explore different kernel tricks to capture non-linearities within your data. Plot the results and compare the accuracies for different kernels.

```
# Define the range of PCA components and the SVM kernels to test
component range = list(range(1, X train.shape[1] + 1))
svm kernels = ['linear', 'poly', 'rbf', 'sigmoid']
# Dictionary to save accuracy for each kernel across different PCA
dimensions
results by kernel = {kernel: {'components': [], 'accuracy': []} for
kernel in svm kernels}
# Iterate over each kernel type
for kernel in svm kernels:
    # Keep track of the highest accuracy and the corresponding number
of components
    top accuracy = 0
    optimal k = None
    # Test SVM performance for each number of PCA components
    for k in component range:
        # Apply PCA to reduce feature dimensions
        pca model = PCA(n components=k)
        X train reduced = pca model.fit transform(X train)
        X test reduced = pca model.transform(X test)
```

```
# Train the SVM model with the current kernel
        svm model = SVC(kernel=kernel, random state=0)
        svm model.fit(X train reduced, y train)
        # Make predictions and calculate accuracy
        predictions = svm model.predict(X test reduced)
        acc = accuracy_score(y_test, predictions)
        # Save accuracy for plotting
        results by kernel[kernel]['components'].append(k)
        results by kernel[kernel]['accuracy'].append(acc)
        # Update best accuracy and corresponding K if current is
better
       if acc > top accuracy:
            top accuracy = acc
            optimal k = k
# Visualize how accuracy changes with PCA components for each kernel
plt.figure(figsize=(10, 6))
for kernel in svm kernels:
    plt.plot(results by kernel[kernel]['components'],
             results by kernel[kernel]['accuracy'],
             label=f'{kernel} kernel')
plt.xlabel('Number of PCA Components')
plt.ylabel('Classification Accuracy')
plt.title('SVM Accuracy Across Kernels and PCA Components')
plt.legend()
plt.grid(True)
plt.show()
```





```
import pandas as pd
# List to store kernel summary
kernel summary = []
# Loop over each kernel type
for kernel in svm kernels:
    best accuracy = 0
    best k = None
    for k in k value:
        # Apply PCA
        pca = PCA(n components=k)
        X_train_pca = pca.fit_transform(X_train)
        X test pca = pca.transform(X test)
        # Train SVM
        svm = SVC(kernel=kernel, random state=0)
        svm.fit(X_train_pca, y_train)
        # Predict and calculate accuracy
        Y_pred = svm.predict(X_test_pca)
        accuracy = accuracy_score(y_test, Y_pred)
        # Update best accuracy
        if accuracy > best accuracy:
```

```
best accuracy = accuracy
            best k = k
    # Append results to the list
    kernel summary.append({
        'Kernel': kernel,
        'Best K': best k,
        'Accuracy for Best K': best accuracy
    })
# Convert to DataFrame and display as a table
kernel table = pd.DataFrame(kernel summary)
print(kernel table)
            Best K Accuracy for Best K
    Kernel
0
    linear
                28
                               0.982456
1
                11
      poly
                               0.912281
2
       rbf
                 9
                               0.982456
3 sigmoid
                10
                               0.964912
```

## Problem 2

Develop a SVR regression model that predicts housing price based on the following input variables:

Area, bedrooms, bathrooms, stories, mainroad, guestroom, basement, hotwaterheating, airconditioning, parking, prefarea

```
from sklearn.svm import SVR
from sklearn.metrics import mean squared error, r2 score
from sklearn.linear model import Ridge
from sklearn.impute import SimpleImputer
# Mount Google Drive
from google.colab import drive
drive.mount('/content/drive')
# load housing dataset
filePath = '/content/drive/MyDrive/Colab Notebooks/Housing.csv'
df = pd.read csv(filePath)
# Display first 5 rows
df.head()
Drive already mounted at /content/drive; to attempt to forcibly
remount, call drive.mount("/content/drive", force remount=True).
{"summary":"{\n \"name\": \"df\",\n \"rows\": 545,\n \"fields\": [\
             \"column\": \"price\",\n \"properties\": {\n
    {\n
\"dtype\": \"number\",\n \"std\": 1870439,\n
                                                         \"min\":
                \"max\": 13300000,\n
1750000,\n
                                          \"num unique values\":
```

```
5285000,\n
{\n \"dtype\": \"number\",\n \"std\": 0,\n
\"min\": 0,\n \"max\": 3,\n \"num_unique_values\": 4,\n
```

```
\"samples\": [\n
                         3,\n
                                        1\n
\"semantic type\": \"\",\n
                                 \"description\": \"\"\n
                                                               }\
n },\n {\n \"column\": \"prefarea\",\n \"properties\":
           \"dtype\": \"category\",\n \"num_unique_values\":
{\n
                                   \"no\",\n
2,\n
          \"samples\": [\n
                                                        \"ves\"\n
          \"semantic_type\": \"\",\n \"description\": \"\"\n
],\n
}\n },\n {\n \"column\": \"furnishingstatus\",\n \"dtype\": \"category\",\n
\"num_unique_values\": 3,\n
                                  \"samples\": [\n
\"furnished\",\n\\"semi-furnished\"\n
                                                      ],\n
\"semantic type\": \"\",\n \"description\": \"\"\n
                                                              }\
     }\n ]\n}","type":"dataframe","variable_name":"df"}
# Impute numeric columns if there are NaNs
numeric cols = ['area', 'bedrooms', 'bathrooms', 'stories', 'parking']
imputer = SimpleImputer(strategy='median')
df[numeric cols] = imputer.fit transform(df[numeric cols])
# Convert binary categorical columns to 0/1
binary cols =
['mainroad','guestroom','basement','hotwaterheating','airconditioning'
,'prefarea']
for col in binary cols:
    df[col] = df[\overline{col}].map(\{'yes': 1, 'no': 0\})
# Remove any constant columns (zero variance)
for col in numeric_cols + binary_cols:
    if df[col].nunique() <= 1:</pre>
        print(f"Removing constant column: {col}")
        df.drop(columns=[col], inplace=True)
# Define features and target
feature cols = ['area', 'bedrooms', 'bathrooms', 'stories',
                'mainroad', 'guestroom', 'basement',
'hotwaterheating',
                'airconditioning', 'parking', 'prefarea']
X = df[feature cols].values
y = df['price'].values
# Remove any samples with missing target
mask = \sim np.isnan(y)
X = X[mask]
y = y[mask]
# Split into training and test sets (80/20)
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=8)
# Standardize features
scaler = StandardScaler()
```

```
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Ensure no NaNs or Infs
X_{\text{train\_scaled}} = \text{np.nan\_to\_num}(X_{\text{train\_scaled}}, \text{nan=0.0}, \text{posinf=0.0},
neginf=0.0)
X test scaled = np.nan to num(X test scaled, nan=0.0, posinf=0.0,
neginf=0.0)
# Dictionary to store results
results = []
predictions = {}
# Train SVR models with different kernels
svr kernels = {
    'RBF': SVR(kernel='rbf', C=1000, gamma=0.1),
    'Linear': SVR(kernel='linear', C=1000),
    'Polynomial': SVR(kernel='poly', C=1000, degree=2)
}
for name, model in svr kernels.items():
    model.fit(X_train_scaled, y_train)
    y_pred = model.predict(X_test_scaled)
    predictions[name] = y pred
    mse = mean squared error(y test, y pred)
    r2 = r2_score(y_test, y_pred)
    results.append({
        'Model': f'{name} SVR',
        'MSE': round(mse, 2),
        'R2': round(r2, 3)
    })
# Ridge Regression
ridge = Ridge(alpha=1.0)
ridge.fit(X train scaled, y train)
y_ridge_pred = ridge.predict(X test scaled)
ridge mse = mean squared error(y test, y ridge pred)
ridge r2 = r2 score(y test, y ridge pred)
results.append({
    'Model': 'Ridge Regression',
    'MSE': round(ridge mse, 2),
    'R2': round(ridge r2, 3)
})
# Convert results to a DataFrame and display
results df = pd.DataFrame(results)
print(results df)
```

```
Model
                              MSE
0
            RBF SVR 4.066866e+12 -0.060
1
         Linear SVR 2.483255e+12 0.353
2
     Polynomial SVR 4.115708e+12 -0.072
3
  Ridge Regression 1.303046e+12 0.661
# Plot predictions
plt.figure(figsize=(10, 6))
plt.scatter(y test, y test, color='black', label='True prices',
alpha=0.6)
colors = ['navy', 'c', 'cornflowerblue']
for color, name in zip(colors, svr_kernels.keys()):
    plt.scatter(y test, predictions[name], color=color, label=f'{name}
SVR')
plt.scatter(y test, y ridge pred, color='red', label='Ridge
Regression', alpha=0.6)
plt.xlabel("Actual Prices")
plt.ylabel("Predicted Prices")
plt.title("SVR and Ridge Regression Predictions vs Actual Prices")
plt.legend()
plt.grid(True)
plt.show()
```

