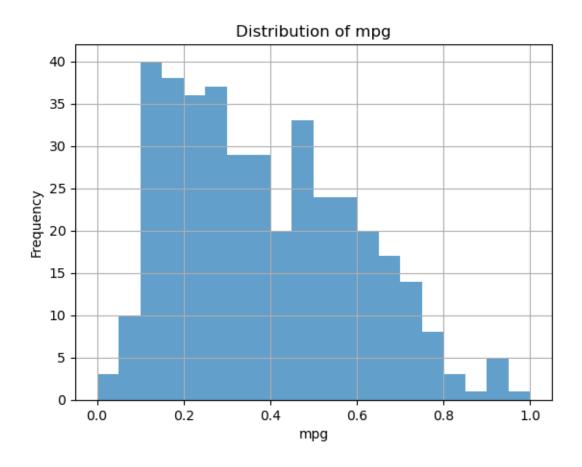
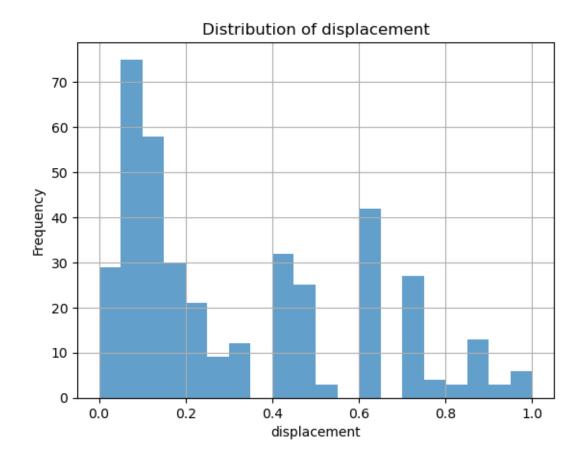
hw2

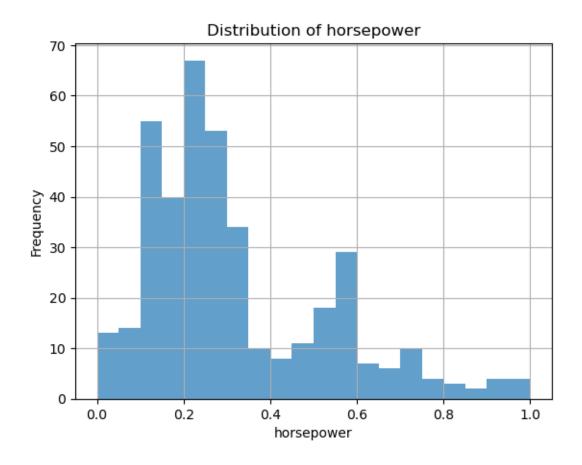
January 30, 2025

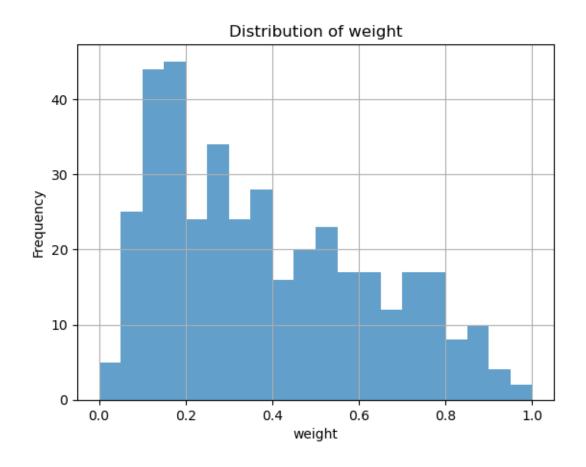
```
import pandas as pd
    from sklearn.preprocessing import MinMaxScaler
    import matplotlib.pyplot as plt
    data = pd.read_csv('auto-mpg.csv')
    # Apply one-hot encoding
    encoded data = pd.get_dummies(data, columns=['origin'], prefix='origin')
    # Normalize each field of the input data using the min-max normalization_
     \hookrightarrow technique
    scaler = MinMaxScaler()
    encoded_data[numerical_columns] = scaler.

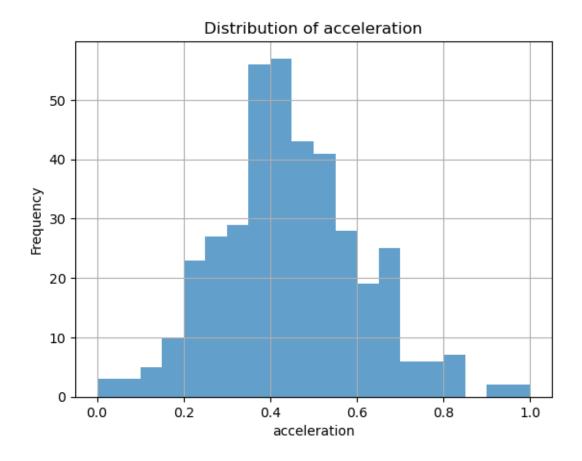
→fit_transform(encoded_data[numerical_columns])
    # Plot the data
    for column in numerical columns:
        encoded_data[column].hist(bins=20, alpha=0.7)
        plt.title(f'Distribution of {column}')
        plt.xlabel(column)
        plt.ylabel('Frequency')
        plt.show()
    # Data Analyzation
    # - mpq: skewed to right
    # - displacement: skewed to right
    # - horsepower: skewed to right
    # - weight: skewed to right
    # - acceleration: symmetric
```





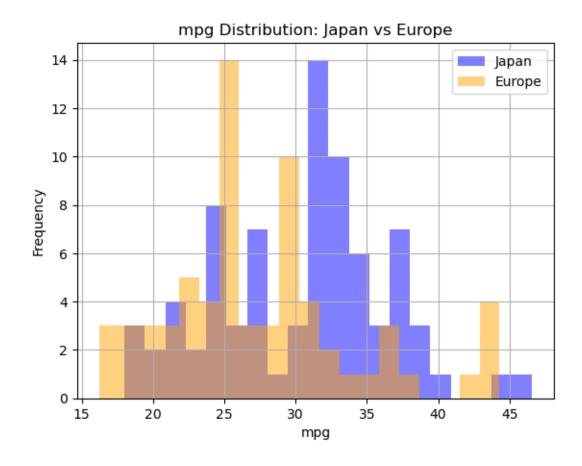


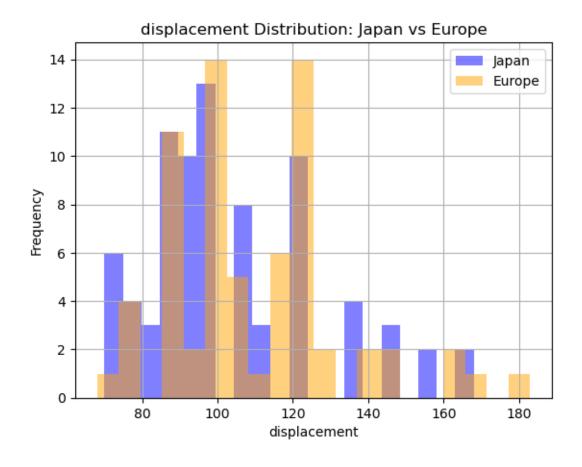


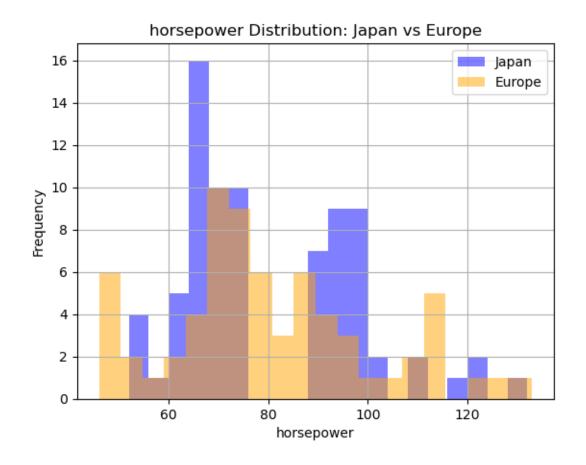


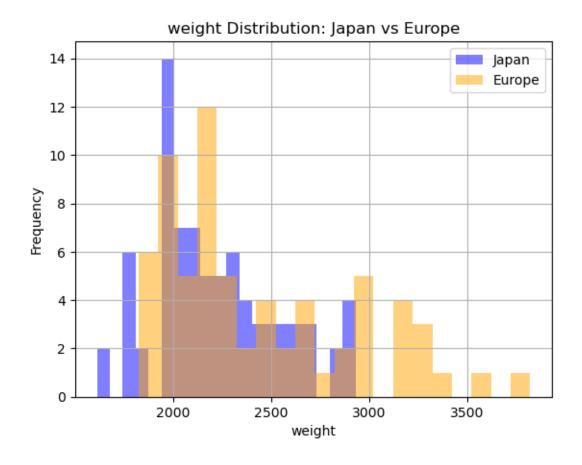
```
X = filtered_data[numerical_columns]
y = filtered_data['origin_USA']
# Split the data
→random_state=42)
# Build a Logistic Regression model
model = LogisticRegression()
model.fit(X_train, y_train)
# Prediction
y_pred = model.predict(X_test)
# Report the precision and recall
report = classification_report(y_test, y_pred, target_names=['Japan', 'USA'])
print(report)
japan_europe_data = data[data['origin'].isin(['Japan', 'Europe'])]
japan_europe_data = pd.get_dummies(japan_europe_data, columns=['origin'],_
 ⇔prefix='origin')
for column in numerical_columns:
   japan_europe_data[japan_europe_data['origin_Japan'] == 1][column].
 ⇔hist(bins=20, alpha=0.5, label='Japan', color='blue')
   japan_europe_data[japan_europe_data['origin_Japan'] == 0][column].
 ⇔hist(bins=20, alpha=0.5, label='Europe', color='orange')
   plt.title(f'{column} Distribution: Japan vs Europe')
   plt.xlabel(column)
   plt.ylabel('Frequency')
   plt.legend()
   plt.show()
# From the histograms, the horsepower, mpg, and acceleration have clearly_
⇒different distributions, which is highly predictive of the accuracy of a⊔
 ⇔model using them.
# On the other hand, the other features have a lot of overlap in their
 →distributions, which could potentially have negative effect on model
 ⇒performance.
```

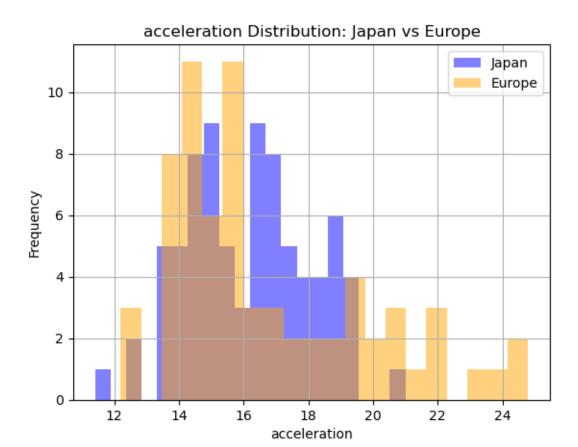
	precision	recall	f1-score	support
Japan	0.76	0.94	0.84	17
USA	0.98	0.90	0.93	48
accuracy			0.91	65
macro avg	0.87	0.92	0.89	65











```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import PolynomialFeatures
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error, r2_score

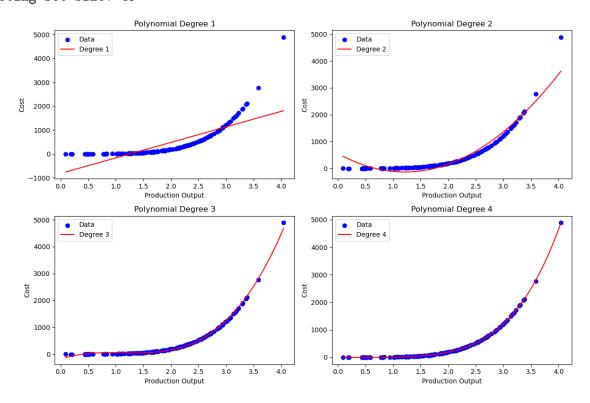
data = pd.read_csv('cost.csv')

# Define features and targets
X_cost = data[['production_output']]
y_cost = data['cost']

# Split the data
X_train_cost, X_test_cost, y_train_cost, y_test_cost = train_test_split(X_cost,u_oy_cost, test_size=0.3, random_state=42)
```

```
print("Training set size:", X_train_cost.shape[0])
print("Testing set size:", X_test_cost.shape[0])
degrees = [1, 2, 3, 4]
results = []
plt.figure(figsize=(12, 8))
for degree in degrees:
   poly = PolynomialFeatures(degree)
   X_train_poly = poly.fit_transform(X_train_cost)
   X_test_poly = poly.fit_transform(X_test_cost)
   model = LinearRegression()
   model.fit(X_train_poly, y_train_cost)
   y_train_pred = model.predict(X_train_poly)
   y_test_pred = model.predict(X_test_poly)
   # Calculate the RMSE and R^2
   rmse_train = np.sqrt(mean_squared_error(y_train_cost, y_train_pred))
   rmse_test = np.sqrt(mean_squared_error(y_test_cost, y_test_pred))
   r2_train = r2_score(y_train_cost, y_train_pred)
   r2_test = r2_score(y_test_cost, y_test_pred)
    # Store the results
   results.append({
        'degree': degree,
        'rmse_train': rmse_train,
        'rmse_test': rmse_test,
        'r2_train': r2_train,
       'r2_test': r2_test
   })
    # Plotting and fitting line
   plt.subplot(2, 2, degree)
   plt.scatter(X_cost, y_cost, color='blue', label='Data')
   X range = np.linspace(X cost.min(), X cost.max(), 100).reshape(-1, 1)
   X_range_poly = poly.fit_transform(X_range)
   y_range_pred = model.predict(X_range_poly)
   plt.plot(X_range, y_range_pred, color='red', label=f'Degree {degree}')
   plt.title(f'Polynomial Degree {degree}')
   plt.xlabel('Production Output')
   plt.ylabel('Cost')
   plt.legend()
```

Training set size: 105 Testing set size: 45



```
degree
           rmse_train
                        rmse_test
                                    r2_train
                                               r2_test
0
        1
           438.842780
                       281.084793
                                    0.607279
                                              0.529909
1
           181.973847
                        115.351446
                                    0.932472
                                              0.920831
2
        3
            39.467718
                         26.623867
                                    0.996823
                                              0.995783
3
        4
             5.995418
                          5.998611
                                    0.999927
                                              0.999786
```

```
[25]: # Exercise 4 import pandas as pd
```

```
import numpy as np
import matplotlib.pyplot as plt
data = pd.DataFrame({
    'Weight': [7.0, 6.0, 8.0],
    'Length': [50, 55, 56],
     'Actual_Height': [5.80, 5.70, 6.00]
})
# define the parameter of the model
coefficient weight = 0.1
coefficient_length = 0.1
# Assume distribution to be 0.1
sigma_squared = 0.1
# Calculate the predicted height
data['Predicted_Height'] = coefficient_weight * data['Weight'] +__
 →coefficient_length * data['Length']
# Calculate the Squared Residual
data['Squared_Residual'] = (data['Actual_Height'] - data['Predicted_Height'])__
 →** 2
# Function to calculate the Log-Likelihood
def log_likelihood(y, y_pred, sigma_squared):
    residual_squared = (y - y_pred) ** 2
    return -0.5 * np.log10(2 * np.pi * sigma_squared) - (residual_squared / (2_{L}

sigma_squared * np.log(10)))

# Calculate the Log-Likelihood for each data point
data['Log_Likelihood'] = data.apply(
    lambda row: log_likelihood(row['Actual_Height'], row['Predicted Height'], u
 ⇒sigma_squared), axis=1
# Sum the Log-Likelihood
log_likelihood_sum = data['Log_Likelihood'].sum()
print(data[['Weight', 'Length', 'Actual_Height', 'Predicted_Height', u

¬'Squared_Residual', 'Log_Likelihood']])
print(f"\nTotal Log-Likelihood: {log_likelihood_sum:.4f}")
  Weight Length Actual_Height Predicted_Height Squared_Residual \
0
      7.0
               50
                             5.8
                                               5.7
                                                                 0.01
      6.0
                             5.7
                                               6.1
1
               55
                                                                 0.16
2
      8.0
               56
                             6.0
                                               6.4
                                                                 0.16
```

Log_Likelihood 0 0.079195 1 -0.246526 2 -0.246526

Total Log-Likelihood: -0.4139