experiment1

```
# import necessary libraries
import pandas as pd
                                   # pandas: for creating and manipulating the data
import seaborn as sns
                                  # seaborn: for making the plot
import matplotlib.pyplot as plt # matplotlib: for additional plot customization
import numpy as np
                                    # numpy: for numerical operations (e.g., random
number generation)
# create a dataframe with the initial data
data = pd.DataFrame({
'Name': ['Alice', 'Bob', 'Charlie'],
                                   # List of student names
                                       # Math scores for each student
'Math': [85, 90, 95],
                                      # Science scores for each student
'Science': [88, 92, 98]
})
# Add a new subject, "History", with randomly generated scores
data['History'] = np.random.randint(80, 100, len(data))
                                                                      # Generates random
history scores between 80 and 100
# Prepare the data for plotting: we need a tidy format (long format)
combined = pd.DataFrame({
'Name': list(data['Name']) *
3,
                                                                   # Duplicate the names for
the 3 subjects
'Subject': ['Math', 'Science', 'History'] *
                                                     # List of subjects repeated for each
len(data),
individual
'Score': list(data['Math']) + list(data['Science']) + list(data['History'])
                                                                                      #
Combine the scores into one column
})
# Plot the data using seaborn's barplot function
sns.barplot(data=combined, x='Name', y='Score', hue='Subject')
                                                                                          #
Create a bar plot, hue differentiates the subjects
plt.title('All
                                                                                       # Add
Scores')
a title to the plot
plt.tight_layout()
    # Adjust layout to make sure everything fits properly
```

```
plt.show()
   # Display the plot
in r
# Load the necessary libraries
                                      # dplyr: for data manipulation (filtering, reshaping,
library(dplyr)
etc.)
library(ggplot2)
                                        # ggplot2: for creating plots
# Create a data frame with the initial data
data <- tibble(
Name = c("Alice", "Bob", "Charlie"),
                                                      # List of student names
                                                       # Math scores for each student
Math = c(85, 90, 95),
Science = c(88, 92, 98)
                                                         # Science scores for each student
)
# Add a new subject, "History", with randomly generated scores
set.seed(0)
                                                                               # Set a
random seed for reproducibility
data$History <- sample(80:100, nrow(data), replace = TRUE) # Generate random
history scores between 80 and 100
# Combine the data into a long format (tidy format) for plotting
combined <- tibble(
Name = rep(data$Name, 3),
                                                                                  #
Duplicate names for each subject (Math, Science, History)
Subject = rep(c("Math", "Science", "History"), each = nrow(data)),
                                                                          # Repeat the
subject names for each individual
Score = c(data$Math, data$Science, data$History)
                                                                         # Combine the
scores for each subject into one column
)
# Plot the data using ggplot2
ggplot(combined, aes(x = Name, y = Score, fill = Subject)) +
Create a bar plot with Name on x-axis, Score on y-axis, and bars filled by Subject
geom col(position = "dodge")
+
                                                              # Create grouped bars (dodge
makes them side by side)
ggtitle("All Scores") # Add a title to the plot
```

experiment 2

```
import pandas as pd
from sklearn.linear_model import LinearRegression
from sklearn.metrics import r2 score, mean squared error
import matplotlib.pyplot as plt
# Load the CSV data (ensure 'data.csv' is in your working directory)
data = pd.read_csv('data.csv')
# Define the independent (Experience) and dependent (Salary) variables
X = data[['Experience']] # Independent variable (2D array)
y = data['Salary'] # Dependent variable
# Create and fit the linear regression model
model = LinearRegression()
model.fit(X, y)
# Make predictions
y_pred = model.predict(X)
# Plot actual vs predicted data
plt.scatter(X, y, color='blue', label='Actual')
plt.plot(X, y pred, color='red', label='Predicted Line')
plt.xlabel('Experience')
plt.ylabel('Salary')
plt.title('Simple Linear Regression')
plt.legend()
plt.show()
# Calculate and print R<sup>2</sup> and RMSE
r2 = r2\_score(y, y\_pred)
rmse = mean_squared_error(y, y_pred, squared=False)
print("R2:", r2)
```

```
print("RMSE:", rmse)
# Calculate residuals (actual - predicted)
residuals = y - y pred
# Plot the residuals
plt.scatter(X, residuals, color='purple')
plt.axhline(y=0, color='black', linestyle='--')
plt.title('Residual Plot')
plt.xlabel('Experience')
plt.ylabel('Residuals')
plt.show()
in r
# Install required package (if not already installed)
# install.packages("ggplot2")
# Load necessary library for plotting
library(ggplot2)
# Load the CSV data (ensure 'data.csv' is in your working directory)
data <- read.csv("data.csv")</pre>
# Define the independent variable (X) and dependent variable (y)
X <- data$Experience # Independent variable
y <- data$Salary # Dependent variable
# Fit the linear regression model
model <- Im(Salary ~ Experience, data = data)
# Display the model summary (coefficients, R-squared, etc.)
summary(model)
# Make predictions using the model
y_pred <- predict(model, newdata = data)</pre>
# Plot actual vs predicted values
ggplot(data, aes(x = Experience, y = Salary)) +
```

```
geom point(color = "blue") + # Actual data points (scatter plot)
geom smooth(method = "Im", color = "red") + # Predicted line (regression line)
labs(title = "Simple Linear Regression", # Add title
x = "Experience", y = "Salary") # Label axes
# Calculate R-squared and RMSE
r2 <- summary(model)$r.squared
rmse <- sqrt(mean((y - y_pred)^2)) # Calculate Root Mean Squared Error
cat("R<sup>2</sup>:", r2, "\n")
cat("RMSE:", rmse, "\n")
# Calculate residuals (actual - predicted)
residuals <- y - y pred
# Plot the residuals
ggplot(data, aes(x = Experience, y = residuals)) +
geom point(color = "purple") +
geom_hline(yintercept = 0, color = "black", linetype = "dashed") +
labs(title = "Residual Plot", x = "Experience", y = "Residuals")
experiment 3 multiple regression
import pandas as pd # Import pandas for data manipulation
import statsmodels.api as sm # Import statsmodels for regression analysis
import seaborn as sns # Import seaborn for data visualization
import matplotlib.pyplot as plt # Import matplotlib for plotting
# Load data from CSV
data = pd.read csv('employee data.csv') # Read the employee data from
'employee data.csv' file
# Convert categorical Education column to dummy variables
data = pd.get dummies(data, columns=['Education'], drop first=True)
# Create dummy variables for the 'Education' column (e.g., converts 'Education' to
'Education Master' and 'Education PhD')
# Features and target
X = data.drop(columns='Salary') # Define features (independent variables) by dropping the
target column 'Salary'
```

```
y = data['Salary'] # Define the target variable (dependent variable)
# Add constant for intercept
X const = sm.add constant(X) # Add a constant column to the feature matrix (intercept
term)
# Fit regression model
model = sm.OLS(y, X_const).fit() # Fit an OLS (Ordinary Least Squares) regression model
using statsmodels
print(model.summary()) # Print the summary of the regression model, including coefficients,
R-squared, etc.
# Predict for new employee: 6 years experience, PhD
new data = pd.DataFrame({
'Experience': [6], # New employee has 6 years of experience
'Education Master': [0], # The employee doesn't have a Master's degree (i.e., it's a PhD)
'Education PhD': [1] # The employee has a PhD (this is the reference category after dummy
encoding)
})
new data = sm.add constant(new data) # Add constant for the new data (intercept term)
prediction = model.predict(new_data) # Predict the salary for the new employee based on
the fitted model
print("Predicted Salary:", prediction.iloc[0]) # Print the predicted salary for the new
employee
# Plot actual vs predicted
sns.scatterplot(x=model.fittedvalues, y=y) # Create a scatter plot of actual vs predicted salary
values
plt.plot([y.min(), y.max()], [y.min(), y.max()], 'r--') # Plot a red dashed line to represent
perfect predictions
plt.xlabel("Predicted Salary") # Label for the x-axis
plt.ylabel("Actual Salary") # Label for the y-axis
plt.title("Predicted vs Actual Salary") # Title of the plot
plt.show() # Show the plot
```

```
# Load necessary libraries
library(readr) # For reading CSV files
library(dplyr) # For data manipulation (if needed)
library(stats) # For fitting linear models (Im function)
library(ggplot2) # For plotting
# Load the data from CSV
data <- read_csv("employee_data.csv") # Assuming the CSV is in the current working
directory
# Convert categorical Education column to factors (automatically handled by Im function)
data$Education <- factor(data$Education)</pre>
# Create the multiple regression model
# Education is converted to dummy variables automatically by the 'lm' function
model <- Im(Salary ~ Experience + Education, data = data)
# Print summary of the regression model
summary(model)
# Predict for a new employee: 6 years experience, PhD
new data <- data.frame(</pre>
Experience = 6,
Education = factor("PhD", levels = levels(data$Education)) # Assuming 'PhD' is one of the
Education categories
)
predicted salary <- predict(model, new data) # Predict using the fitted model
print(paste("Predicted Salary for new employee:", predicted_salary))
# Plot actual vs predicted salary values
ggplot(data, aes(x = fitted(model), y = Salary)) +
geom_point() + # Actual vs predicted values
geom abline(slope = 1, intercept = 0, color = "red") + # Red line for perfect predictions
labs(x = "Predicted Salary", y = "Actual Salary", title = "Predicted vs Actual Salary") +
theme minimal()
```

```
# Load necessary libraries
library(lubridate) # For date manipulation
library(forecast) # For time series forecasting and analysis
library(dplyr) # For data manipulation (e.g., drop na, arrange)
# Load the data
data <- read.csv("your file.csv") # Replace "your file.csv" with your actual file path
# Convert the 'Date' column to Date type
data$Date <- as.Date(data$Date, format="%Y-%m-%d")
# Remove rows with missing values in the 'close' column
data <- data %>% drop na(close)
# Sort data by Date
data <- data %>% arrange(Date)
# Create a time series object
ts_data <- ts(data$close, start=c(year(min(data$Date)), month(min(data$Date))),
frequency=252)
# Plot the time series
plot(ts_data, main="TCS Stock Closing Price Time Series", ylab="Close Price", xlab="Time")
experiment 5 ARIMA MODEL
# load required libraries
library(forecast)
library(ggplot2)
# read the csv data (adjust path as needed)
solar prod input <- read.csv("c:/users/lab204/downloads/solar prod.csv")</pre>
# convert to time series object (assuming monthly data)
solar prod <- ts(solar prod input[, 2], start = c(1), frequency = 12) # adjust 'start' if needed
# plot original time series
plot(solar prod, xlab = "time (months)", ylab = "solar production (kwh)", main = "solar
production time series")
```

```
# fit arima model (auto.arima can also be used)
arima_model <- arima(solar_prod, order = c(0, 1, 0), seasonal = list(order = c(1, 0, 0), period
= 12))
# print model summary
summary(arima model)
# forecast next 12 periods
arima_forecast <- forecast(arima_model, h = 12)</pre>
# plot forecast with confidence intervals
plot(arima_forecast, main = "forecast of solar production")
# optionally extract forecast values and confidence intervals
pred_matrix <- cbind(</pre>
LB = arima forecast$lower[, 2],
Pred = arima forecast$mean,
UB = arima_forecast$upper[, 2]
)
# show prediction matrix
print(round(pred_matrix, 2))
experiment6 -email spam classifier code
import pandas as pd
import string
import nltk
from nltk.corpus import stopwords
from sklearn.model selection import train test split
from sklearn.feature_extraction.text import CountVectorizer
from sklearn.naive_bayes import MultinomialNB
from sklearn.metrics import accuracy score, classification report
```

Download stopwords

```
nltk.download('stopwords')
stop words = set(stopwords.words('english'))
# Load dataset (replace with your actual dataset URL or file path)
url = 'https://raw.githubusercontent.com/justmarkham/pycon-2016-
tutorial/master/data/sms.tsv'
df = pd.read csv(url, sep='\t', header=None, names=['label', 'message'])
# Preprocess text
def clean_text(msg):
msg = msg.lower()
msg = ".join([char for char in msg if char not in string.punctuation])
msg = ''.join([word for word in msg.split() if word not in stop words])
return msg
df['cleaned'] = df['message'].apply(clean_text)
# Vectorization
vectorizer = CountVectorizer()
X = vectorizer.fit transform(df['cleaned'])
# Label encoding
y = df['label'].map({'ham': 0, 'spam': 1})
# Train-test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Train the model
model = MultinomialNB()
model.fit(X_train, y_train)
# Evaluate model
y pred = model.predict(X test)
print("Accuracy:", accuracy_score(y_test, y_pred))
print(classification report(y test, y pred))
# Function to predict if a message is spam or not
def predict spam(message):
# Clean the input message
cleaned message = clean text(message)
```

```
# Convert the message into a feature vector
message_vector = vectorizer.transform([cleaned_message])
# Predict the category (0 = not spam, 1 = spam)
prediction = model.predict(message_vector)
# Return the result
if prediction == 1:
return "This message is SPAM"
else:
return "This message is NOT SPAM"
# Test the function with an example
test_message = input("Enter the message to classify (spam or not): ")
result = predict_spam(test_message)
print(result)
SENTIMENT ANALYSIS
import nltk
from nltk.corpus import movie_reviews
from nltk.classify import NaiveBayesClassifier
from nltk.classify.util import accuracy
# Download necessary datasets
nltk.download('movie reviews')
nltk.download('punkt')
# Function to extract features from a document (review)
def document_features(doc):
words = set(doc)
features = {}
for word in all words:
features[word] = (word in words)
return features
# Load the movie reviews dataset
positive_reviews = movie_reviews.categories('pos')
negative_reviews = movie_reviews.categories('neg')
```

```
# Create a list of (document, category) pairs for training data
documents = []
for category in positive reviews:
for fileid in movie_reviews.fileids(category):
documents.append((movie reviews.words(fileid), 'pos'))
for category in negative reviews:
for fileid in movie_reviews.fileids(category):
documents.append((movie reviews.words(fileid), 'neg'))
# Shufle the documents to randomize the dataset
import random
random.shufle(documents)
# Get all words in the corpus for feature extraction
all words = nltk.FreqDist(w.lower() for w in movie reviews.words())
# Select the most common 2000 words as features
word features = list(all words.keys())[:2000]
# Extract features for each document
featuresets = [(document features(doc), category) for (doc, category) in documents]
# Split the data into training and testing sets (80% training, 20% testing)
train set, test set = featuresets[:int(len(featuresets) * 0.8)], featuresets[int(len(featuresets)
* 0.8):]
# Train a Naive Bayes classifier
classifier = NaiveBayesClassifier.train(train set)
# Evaluate the classifier on the test set
accuracy_result = accuracy(classifier, test_set)
print(f"Accuracy: {accuracy result * 100:.2f}%")
# Print the most informative features
classifier.show most informative features(10)
# Function to classify a new review
def classify review(review):
words_in_review = nltk.word_tokenize(review)
features = document features(words in review)
```

```
return classifier.classify(features)
# Test the sentiment analysis with a custom review
test review = input("Enter a movie review for sentiment analysis: ")
result = classify_review(test_review)
print(f"The sentiment of the review is: {result}")
EXPERIMENT 7
DIFFRENT VISUALTION IN R
# Install ggplot2 if not already installed
# install.packages("ggplot2")
library(ggplot2)
#1. Line Plot
x < - seq(0, 10, by=0.1)
y <- \sin(x)
plot(x, y, type="l", col="blue", main="Line Plot of Sine Wave", xlab="X-axis", ylab="Y-axis")
grid()
# 2. Bar Chart
categories <- c('A', 'B', 'C', 'D')
values <- c(5, 7, 3, 9)
barplot(values, names.arg=categories, col='green', main="Bar Chart Example",
xlab="Categories", ylab="Values")
#3. Histogram
data <- rnorm(1000)
hist(data, breaks=30, col='purple', border='black', main="Histogram Example", xlab="Data
Values", ylab="Frequency")
# 4. Scatter Plot
```

plot(x_scatter, y_scatter, col="red", main="Scatter Plot Example", xlab="X-axis", ylab="Y-

x_scatter <- runif(50)
y scatter <- runif(50)</pre>

```
axis")
# 5. Pie Chart
labels <- c('Python', 'C++', 'Java', 'JavaScript')</pre>
sizes <- c(40, 30, 20, 10)
pie(sizes, labels=labels, col=c('blue', 'green', 'orange', 'red'), main="Pie Chart Example")
EXPERIMENT 8 DIFFRENT VISUALIZATION IN PYTHON
import matplotlib.pyplot as plt
import numpy as np
# 1. Line Plot
x = np.arange(0, 10, 0.1)
y = np.sin(x)
plt.plot(x, y, label='Sine Wave', color='blue')
plt.title('Line Plot of Sine Wave')
plt.xlabel('X-axis')
plt.ylabel('Y-axis')
plt.legend()
plt.grid(True)
plt.show()
# 2. Bar Chart
categories = ['A', 'B', 'C', 'D']
values = [5, 7, 3, 9]
plt.bar(categories, values, color='green')
plt.title('Bar Chart Example')
plt.xlabel('Categories')
plt.ylabel('Values')
plt.show()
```

#3. Histogram

data = np.random.randn(1000)

plt.title('Histogram Example')

plt.xlabel('Data Values')

plt.hist(data, bins=30, color='purple', edgecolor='black')

```
plt.ylabel('Frequency')
plt.show()
#4. Scatter Plot
x_scatter = np.random.rand(50)
y_scatter = np.random.rand(50)
plt.scatter(x scatter, y scatter, color='red')
plt.title('Scatter Plot Example')
plt.xlabel('X-axis')
plt.ylabel('Y-axis')
plt.show()
#5. Pie Chart
labels = ['Python', 'C++', 'Java', 'JavaScript']
sizes = [40, 30, 20, 10]
plt.pie(sizes, labels=labels, autopct='%1.1f%%', startangle=140, colors=['blue', 'green',
'orange', 'red'])
plt.title('Pie Chart Example')
plt.show()
### EXP 8
import matplotlib.pyplot as plt
x=[10,20,30,40]
y=[20,25,35,55]
plt.plot(x,y)
plt.title("Line Chart")
plt.ylabel('Y-axis')
plt.xlabel('X-axis')
plt.show()
import matplotlib.pyplot as plt
import pandas as pd
data =
pd.read csv('/content/tips.csv')
x=data['day']
y=data['total bill']
plt.bar(x,y)
plt.title("Tips Dataset")
plt.ylabel('Total Bill')
plt.xlabel('Day')
plt.show()
```

```
import matplotlib.pyplot as plt
import pandas as pd
data =
pd.read_csv('/content/tips.csv')
x=data['day']
y=data['total_bill']
plt.scatter(x,y)
plt.title("Tips Dataset")
plt.ylabel('Total Bill')
plt.xlabel('Day')
plt.show()
import matplotlib.pyplot as plt
import pandas as pd
data =
pd.read_csv('/content/tips.csv')
cars=['AUDI','BMW','FORD','TESLA'
,'JAGUAR']
data = [23,10,35,15,12]
plt.pie(data,labels=cars)
plt.title("Cars Dataset")
plt.show()
import matplotlib.pyplot as plt
import pandas as pd
np.random.seed(10)
data
=[np.random.normal(0,std,100)
for std in range(1,4)]
plt.boxplot(data,vert=True,patch_
artist=True,
boxprops=dict(facecolor='blue'),
medianprops=dict(color='red'))
plt.xlabel('Data Set')
plt.ylabel('Value')
plt.title('Box Plot')
plt.show()
```

```
data = np.random.randn(1000)
plt.hist(data, bins=30, color='purple', edgecolor='black')
plt.title('Histogram Example')
plt.xlabel('Data Values'
plt.ylabel('Frequency')
plt.show()
```