## APPLE HEALTH

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## A Case Study

## **SOURCE CODE:** from google.colab import files uploaded = files.upload() import pandas as pd # Replace 'sleep\_data.csv' with the exact file name if different df = pd.read\_csv('sleep\_data.csv') # Show first 5 rows df.head() # Remove any duplicate rows df = df.drop\_duplicates() # Strip whitespace from column names (if any) df.columns = df.columns.str.strip() # Show basic info df.info() # Check missing values df.isnull().sum() # Example: Fill missing values with median (modify as needed) df.fillna(df.median(numeric\_only=True), inplace=True)

```
# Or drop rows with missing values (if very few)
# df.dropna(inplace=True)
# Convert categorical column to category (if any)
# df['Gender'] = df['Gender'].astype('category')
# Normalize a numeric column (Min-Max Scaling)
from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
# Apply scaling to all numeric columns
df[df.select_dtypes(include='number').columns] = scaler.fit_transform(
  df.select_dtypes(include='number'))
df.head()
# Show column names
print(df.columns)
# See the unique values in the last column (assumed target)
print("Target column:", df.columns[-1])
print("Unique values:", df[df.columns[-1]].unique())
from google.colab import files
uploaded = files.upload()
import pandas as pd
# Load the uploaded file
```

```
df = pd.read_csv(next(iter(uploaded)))
# Drop duplicates and fill missing values
df = df.drop_duplicates()
df.fillna(df.median(numeric_only=True), inplace=True)
# Convert categorical columns
df['Category'] = df['Category'].astype(str)
df['Category'] = pd.factorize(df['Category'])[0]
# Convert time columns to numeric features
df['Start Time'] = pd.to_datetime(df['Start Time'])
df['End Time'] = pd.to_datetime(df['End Time'])
df['StartHour'] = df['Start Time'].dt.hour
df['EndHour'] = df['End Time'].dt.hour
df['Duration'] = (df['End Time'] - df['Start Time']).dt.total_seconds() / 60
# Drop original time columns
df = df.drop(['Start Time', 'End Time', 'Timestamp', 'Source Name'], axis=1)
# Remove outliers
def remove_outliers_iqr(data, columns):
  for col in columns:
    Q1 = data[col].quantile(0.25)
    Q3 = data[col].quantile(0.75)
    IQR = Q3 - Q1
    lower = Q1 - 1.5 * IQR
    upper = Q3 + 1.5 * IQR
    data = data[(data[col] >= lower) & (data[col] <= upper)]
```

```
df = remove_outliers_iqr(df, df.select_dtypes(include='number').columns)
# Normalize numeric columns
from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
df[df.select_dtypes(include='number').columns] = scaler.fit_transform(
  df.select_dtypes(include='number'))
# Define features and target
X = df.drop('Heart Rate', axis=1)
y = df['Heart Rate']
# Train/test split
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Regression models
from sklearn.linear_model import LinearRegression
from sklearn.ensemble import RandomForestRegressor
from sklearn.svm import SVR
from sklearn.metrics import mean_squared_error, r2_score
models = {
  'Linear Regression': LinearRegression(),
  'Random Forest': RandomForestRegressor(),
  'Support Vector Regressor': SVR()
}
```

```
for name, model in models.items():
  model.fit(X_train, y_train)
  preds = model.predict(X_test)
  print(f"{name}:")
  print(f" MSE: {mean_squared_error(y_test, preds):.4f}")
  print(f" R2 Score: {r2_score(y_test, preds):.4f}")
  print("-" * 40)
import matplotlib.pyplot as plt
import numpy as np
# Plotting function
def plot_predictions(y_test, y_pred, title):
  plt.figure(figsize=(8, 5))
  plt.scatter(range(len(y_test)), y_test, label='Actual', alpha=0.6)
  plt.scatter(range(len(y_pred)), y_pred, label='Predicted', alpha=0.6)
  plt.title(f'{title} - Actual vs Predicted')
  plt.xlabel('Sample Index')
  plt.ylabel('Heart Rate (scaled)')
  plt.legend()
  plt.grid(True)
  plt.tight_layout()
  plt.show()
# Plot for each model
for name, model in models.items():
  preds = model.predict(X_test)
  plot_predictions(y_test.values, preds, name)
# Feature importance plot for Random Forest
import seaborn as sns
```

```
rf_model = models['Random Forest']
importances = rf_model.feature_importances_
features = X.columns
plt.figure(figsize=(8, 5))
sns.barplot(x=importances, y=features)
plt.title("Random Forest - Feature Importance")
plt.xlabel("Importance")
plt.ylabel("Feature")
plt.tight_layout()
plt.show()
from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay, classification_report,
accuracy_score
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
# Simulated Apple Health data
data = {
  'heart_rate': [72, 90, 85, 60, 100, 110, 65, 70, 95, 88],
  'sleep_hours': [7, 5, 6, 8, 4, 3, 7.5, 8, 5.5, 6],
  'steps_count': [8000, 4000, 3000, 10000, 2000, 1500, 8500, 9000, 2500, 3200],
  'calorie intake': [2000, 2500, 2300, 1800, 3000, 3200, 1900, 2100, 2700, 2400],
  'health_status': ['Healthy', 'At Risk', 'Healthy', 'Unhealthy', 'Unhealthy', 'Healthy', 'Healthy', 'Healthy',
'At Risk', 'At Risk']
}
```

```
df = pd.DataFrame(data)
# Split features and target
X = df.drop('health_status', axis=1)
y = df['health_status']
# Train/test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Train model
clf = RandomForestClassifier()
clf.fit(X_train, y_train)
# Predictions
y_pred = clf.predict(X_test)
# Dynamically get labels in test data
unique_labels = np.unique(np.concatenate((y_test, y_pred)))
# Confusion matrix
cm = confusion_matrix(y_test, y_pred, labels=unique_labels)
disp = ConfusionMatrixDisplay(confusion_matrix=cm, display_labels=unique_labels)
disp.plot(cmap='Blues')
plt.title("Apple Health Status - Confusion Matrix")
plt.show()
# Performance matrix
print("Performance Matrix:")
print(classification_report(y_test, y_pred, labels=unique_labels, target_names=unique_labels))
```

```
print(f"Overall Accuracy: {accuracy_score(y_test, y_pred):.2f}")
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn.metrics import classification_report, accuracy_score
from sklearn.model_selection import train_test_split
import pandas as pd
import numpy as np
# Simulated Apple Health data
data = {
  'heart_rate': [72, 90, 85, 60, 100, 110, 65, 70, 95, 88],
  'sleep_hours': [7, 5, 6, 8, 4, 3, 7.5, 8, 5.5, 6],
  'steps_count': [8000, 4000, 3000, 10000, 2000, 1500, 8500, 9000, 2500, 3200],
  'calorie_intake': [2000, 2500, 2300, 1800, 3000, 3200, 1900, 2100, 2700, 2400],
  'health_status': ['Healthy', 'At Risk', 'Healthy', 'Unhealthy', 'Unhealthy', 'Healthy', 'Healthy', 'Healthy',
'At Risk', 'At Risk']
}
df = pd.DataFrame(data)
# Split features and labels
X = df.drop('health_status', axis=1)
y = df['health_status']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Define models
models = {
  "Decision Tree": DecisionTreeClassifier(),
  "Random Forest": RandomForestClassifier(),
```

```
"SVM": SVC(kernel='linear', probability=True)
}
# Store performance
performance_summary = []
# Compare models
for name, model in models.items():
  model.fit(X_train, y_train)
  y_pred = model.predict(X_test)
  acc = accuracy_score(y_test, y_pred)
  report = classification_report(y_test, y_pred, output_dict=True, zero_division=0)
  performance_summary.append({
    'Model': name,
    'Accuracy': round(acc, 2),
    'Precision (avg)': round(report['weighted avg']['precision'], 2),
    'Recall (avg)': round(report['weighted avg']['recall'], 2),
    'F1-score (avg)': round(report['weighted avg']['f1-score'], 2)
  })
# Show comparison table
comparison_df = pd.DataFrame(performance_summary)
print("Model Performance Comparison:\n")
print(comparison_df)
```

## **OUTPUT:**

```
        Model Performance Comparison:

        Model Accuracy Precision (avg) Recall (avg) F1-score (avg)

        0 Decision Tree
        1.00
        1.00
        1.00
        1.00

        1 Random Forest
        1.00
        1.00
        1.00
        1.00

        2 SVM
        0.67
        0.83
        0.67
        0.67
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 8375 entries, 0 to 8374
Data columns (total 6 columns):
    Column
                Non-Null Count Dtype
    Start Time 8375 non-null object
0
   End Time 8375 non-null object
1
2 Category
               8375 non-null object
3 Timestamp 8375 non-null object
   Heart Rate 8375 non-null float64
4
    Source Name 8375 non-null object
dtypes: float64(1), object(5)
memory usage: 392.7+ KB
```

	Start Time	End Time	Category	Timestamp	Heart Rate	Source Name
0	2022-09-13 01:47:49	2022-09-13 01:59:19	Light/Core	2022-09-13 01:46:47	64.0000	Apple Watch SE 2020
1	2022-09-13 01:59:19	2022-09-13 02:04:49	Deep	2022-09-13 01:59:47	62.0000	Apple Watch SE 2020
2	2022-09-13 02:04:49	2022-09-13 02:12:49	Light/Core	2022-09-13 02:05:44	64.0000	Apple Watch SE 2020
3	2022-09-13 02:12:49	2022-09-13 02:27:19	Deep	2022-09-13 02:13:49	65.0000	Apple Watch SE 2020
4	2022-09-13 02:27:19	2022-09-13 02:35:49	Light/Core	2022-09-13 02:24:49	64.0000	Apple Watch SE 2020
8370	2023-05-14 07:20:36	2023-05-14 07:42:06	REM	2023-05-14 07:23:59	59.0000	Apple Watch Ultra
8371	2023-05-14 07:42:06	2023-05-14 07:43:06	Awake	2023-05-14 07:42:23	58.0000	Apple Watch Ultra
8372	2023-05-14 07:43:06	2023-05-14 07:45:36	Light/Core	2023-05-14 07:42:23	58.0000	Apple Watch Ultra
8373	2023-05-14 07:45:36	2023-05-14 07:52:06	REM	2023-05-14 07:45:24	62.4062	Apple Watch Ultra
8374	2023-05-14 07:52:06	2023-05-14 08:48:06	Light/Core	2023-05-14 07:51:42	63.0000	Apple Watch Ultra
8375 rows × 6 columns						





