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**Emotional State Analysis in
Consumer Behavior**

INTRODUCTION

This project delves into the dataset "Study 4," focusing on the impact of anxiety on consumer product evaluations among 83 healthy subjects. Through data mining and analytics, we aim to predict emotional states based on psychophysiological responses. The script employs feature selection, PCA, and classification, showcasing a systematic approach from data loading to model evaluation. This analysis aims to be a predictive tool, offering insights into the relationship between anxiety and consumer choices. The constructed model holds implications for understanding and predicting emotional responses in consumer behavior.

CODE EXPLANATION

1.Import necessary libraries

This segment involves importing essential libraries for data analysis and machine learning. These include pandas for data manipulation, matplotlib and seaborn for data visualization, and scikit-learn for machine learning tasks such as model building and evaluation.

```
✓ [1] import pandas as pd
3s    import matplotlib.pyplot as plt
      import seaborn as sns
      from sklearn.model_selection import train_test_split, cross_val_score
      from sklearn.preprocessing import StandardScaler
      from sklearn.feature_selection import SelectKBest, f_classif
      from sklearn.decomposition import PCA
      from sklearn.cluster import KMeans
      from sklearn.svm import SVC
      from sklearn.ensemble import RandomForestClassifier
      from sklearn.metrics import classification_report, accuracy_score, confusion_matrix
```

2. Load the dataset

Here, the code reads the dataset from a CSV file into a Pandas DataFrame.

```
✓ [2] file_path = "Dataset_Study4.csv"
0s dataset = pd.read_csv(file_path)
```

3. Display first few rows of the dataset

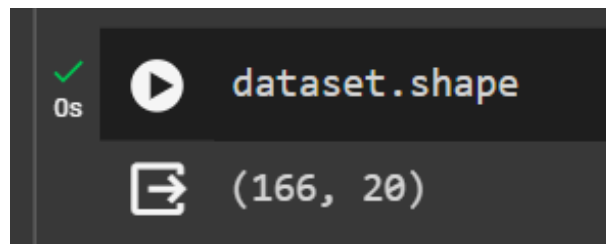
The `dataset.head()` function is used to display the first few rows of the loaded dataset. It's a quick way to get an overview of the structure and content of the data.

```
✓ [3] dataset.head()
```

	HRV_MeanNN	HRV_SDNN	HRV_RMSSD	HRV_Prc20NN	HRV_Prc80NN	HRV_pNN50	HRV_HTI	HRV_VLF	HRV_LF	HRV_HF	HRV_TP	HRV_LFHF	HRV_SD1	HRV_SD2
0	833.314763	106.424590	88.217436	733.8	932.0	50.277778	24.000000	62.654127	713.021695	4228.126423	5161.003600	0.168638	62.465836	137.0405
1	766.966581	92.330308	76.581684	682.8	856.0	49.487179	18.571429	73.007851	1245.114713	3010.533164	4423.639565	0.413586	54.220273	118.6511
2	896.408408	61.685886	51.577499	851.2	948.0	36.526946	14.521739	36.950885	1201.268110	1179.137264	2458.917474	1.018769	36.525411	79.2823
3	833.256267	71.637093	46.592095	789.8	889.0	26.666667	12.857143	156.099084	1241.711995	1299.760891	2731.734549	0.955339	32.991403	95.8961
4	782.793194	45.483921	21.992054	753.4	820.0	1.566580	9.820513	28.439521	396.440159	156.126782	591.933499	2.539219	15.571013	62.4761

4. Get the shape of the dataset

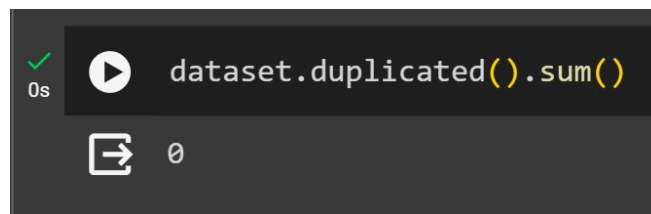
When you run `dataset.shape`, it will return a tuple where the first element is the number of rows, and the second element is the number of columns.



A screenshot of a Jupyter Notebook cell. The cell contains the code `dataset.shape`. To the left of the code is a green checkmark and the text "0s". Below the code is a play button icon. Below the play button is a square icon with a right-pointing arrow, followed by the output `(166, 20)`.

5. Check for duplicated rows

The `dataset.duplicated().sum()` code is checking for and counting the number of duplicated rows in your dataset.



A screenshot of a Jupyter Notebook cell. The cell contains the code `dataset.duplicated().sum()`. To the left of the code is a green checkmark and the text "0s". Below the code is a play button icon. Below the play button is a square icon with a right-pointing arrow, followed by the output `0`.

6. Check for missing values

This part checks for missing values in the dataset using `isnull().sum()` and prints the count of missing values for each column.

```
[3] print(dataset.isnull().sum())
```

HRV_MeanNN	0
HRV_SDNN	0
HRV_RMSSD	0
HRV_Prc20NN	0
HRV_Prc80NN	0
HRV_pNN50	0
HRV_HTI	0
HRV_VLF	0
HRV_LF	0
HRV_HF	0
HRV_TP	0
HRV_LFHF	0
HRV_SD1	0
HRV_SD2	0
HRV_SD1SD2	0
HRV_DFA_alpha1	0
HRV_DFA_alpha2	0
HRV_ApEn	0
HRV_SampEn	0
Label	0

<> dtype: int64

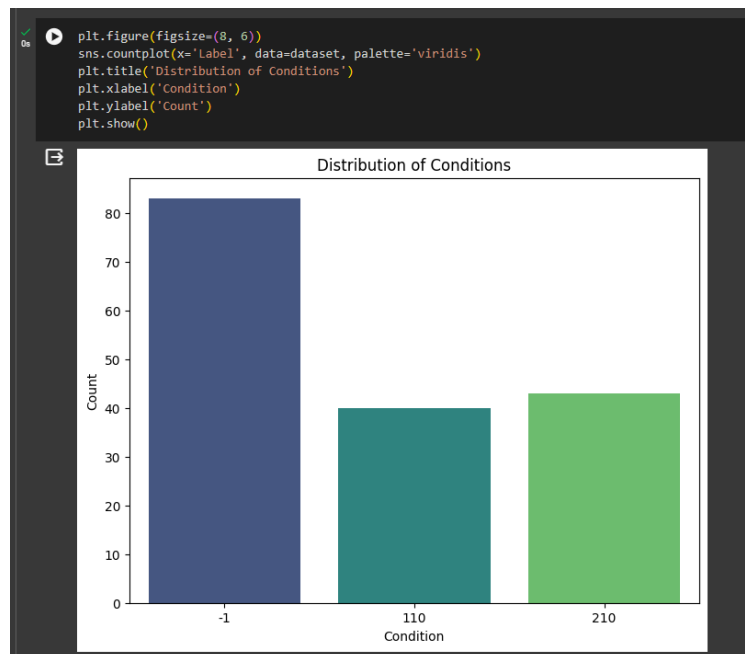
7. Explore the dataset

`dataset.describe()` provides a concise summary of the dataset, including the number of non-null values and data types for each column and total memory usage of the DataFrame. It helps you understand the distribution and variability of your numerical features.

dataset.info() provides summary statistics for numerical columns in your dataset, such as mean, standard deviation, minimum, 25th percentile, 50th percentile(median), 75th percentile, and maximum values. It helps you understand the distribution and variability of your numerical features.

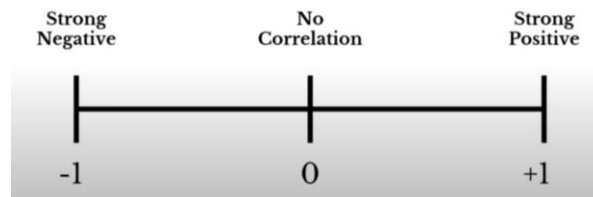
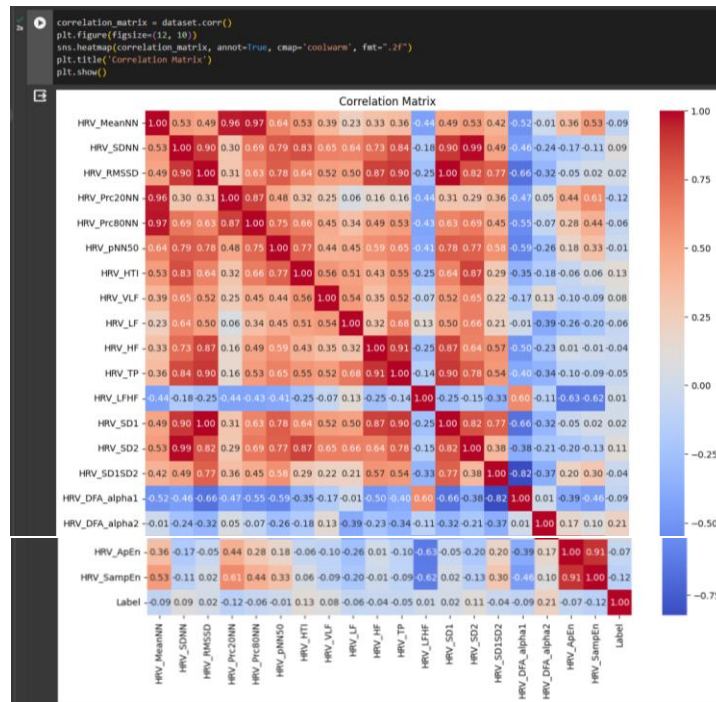
8. Visualize the distribution of the 'Label' column

This code snippet is using the Seaborn library to create a histogram plot to visualize the distribution of emotional states in your dataset.



9. Explore correlations between features

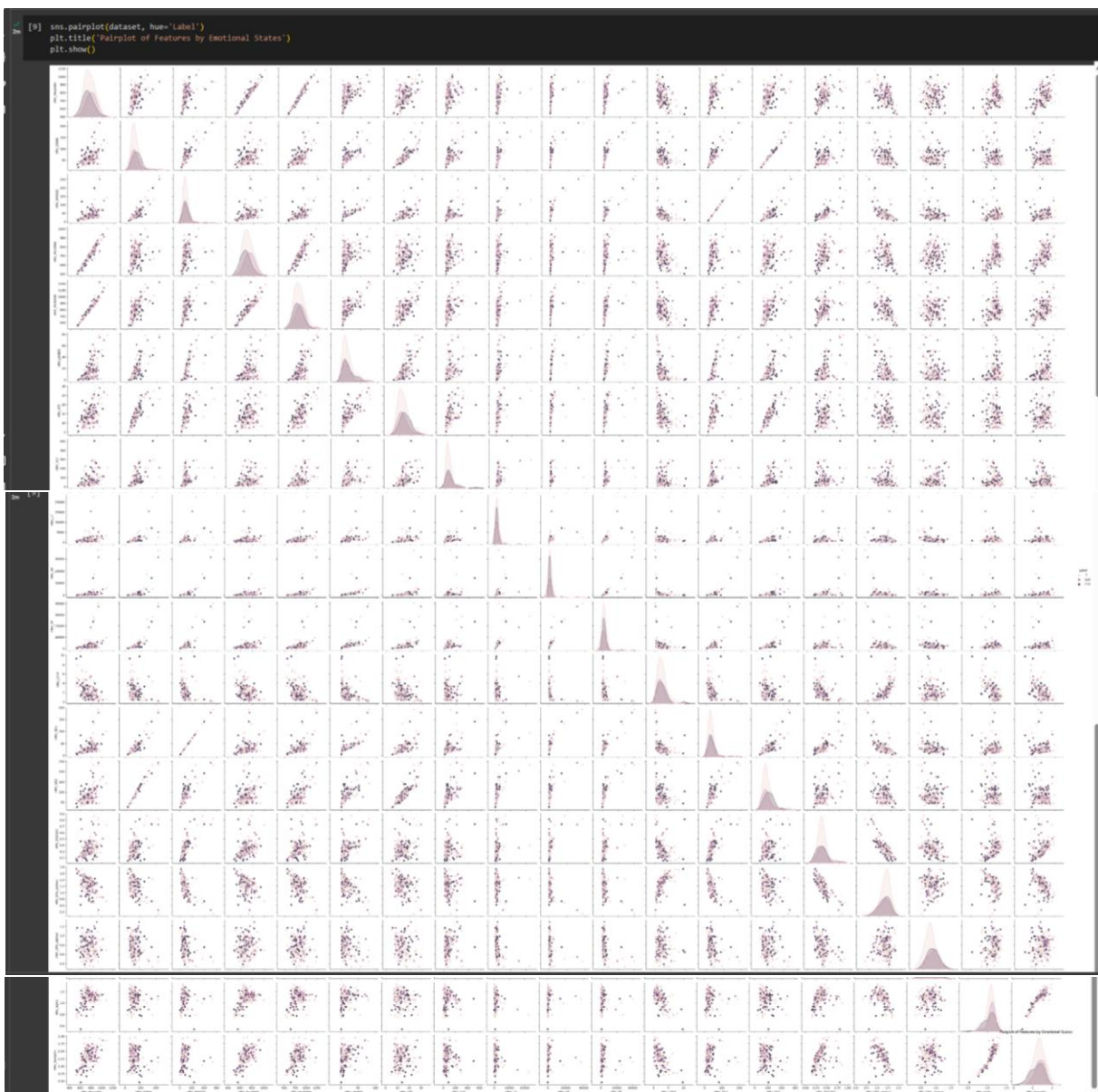
This section generates a heatmap to visualize the correlation matrix of features in the dataset.



Positive correlations suggest that the variables tend to move together, while negative correlations suggest an inverse relationship between the variables.

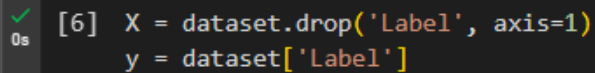
10. Visualize relationships between features and emotional states

Here, a pair plot is generated using seaborn to visualize relationships between features in the dataset, with different emotional states distinguished by color.



11. Separate features and labels

This part separates features (X) and labels (y) from the dataset. The target variable is assumed to be named 'Label.'

A screenshot of a Jupyter Notebook code cell. On the left, there is a green checkmark icon and the text '0s'. The code cell contains two lines of Python code: `X = dataset.drop('Label', axis=1)` and `y = dataset['Label']`.

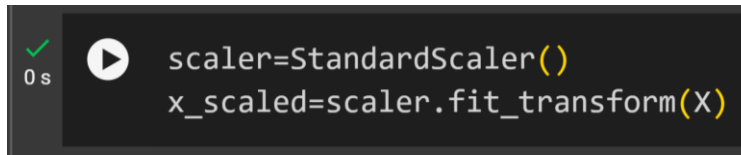
```
[6] X = dataset.drop('Label', axis=1)
     y = dataset['Label']
```

- X: a DataFrame containing only the feature columns (all columns except 'Label').
- y: a Series containing the labels (the 'Label' column).

This separation is essential for training machine learning models, as the features (X) are used as inputs to the model, while the labels (y) are the target outputs that the model will learn to predict.

12. Standardization

The `StandardScaler` in `scikit-learn` is used for standardizing features by removing the mean and scaling them to unit variance.

A code execution snippet showing the initialization of a StandardScaler and its application to a dataset X. The snippet includes a green checkmark, a play button icon, and a timer showing 0s.

```
scaler=StandardScaler()  
x_scaled=scaler.fit_transform(X)
```

13. Feature selection

select the top 4 features based on their importance scores obtained from the ANOVA F-value test (`f_classif`). These selected features are considered the most informative for predicting the target variable (emotional states) in the classification task.

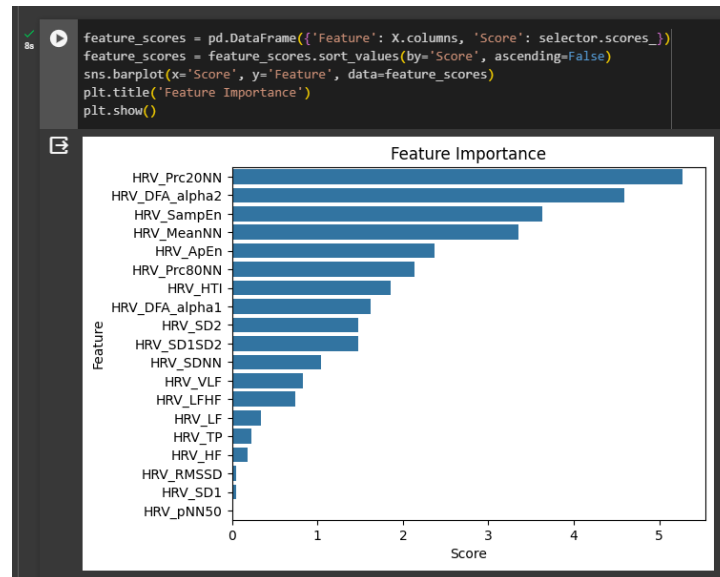
and returns a new array `X_selected` containing only these selected features.

A code execution snippet showing the use of SelectKBest to select the top 4 features based on f_classif scores. The snippet includes a green checkmark, a play button icon, and a timer showing 0s.

```
selector = SelectKBest(f_classif, k=4)  
X_selected = selector.fit_transform(x_scaled, y)
```

14. Visualize feature importance

This section creates a bar plot to visualize the importance scores of features obtained from the ANOVA F-statistic.



15. Standardize features

This part standardizes the selected features using StandardScaler from scikit-learn.

```
[10] scaler = StandardScaler()
      X_standardized = scaler.fit_transform(X_selected)
```

16. Visualize the selected features

The code segment retrieves the names of the selected features after applying the SelectKBest feature selection method.

```
0 s selected_features = X.columns[selector.get_support()]\nprint(selected_features)\n\nIndex(['HRV_MeanNN', 'HRV_Prc20NN', 'HRV_DFA_alpha2', 'HRV_SampEn'], dtype='object')
```

17. Apply PCA to reduce dimensionality.

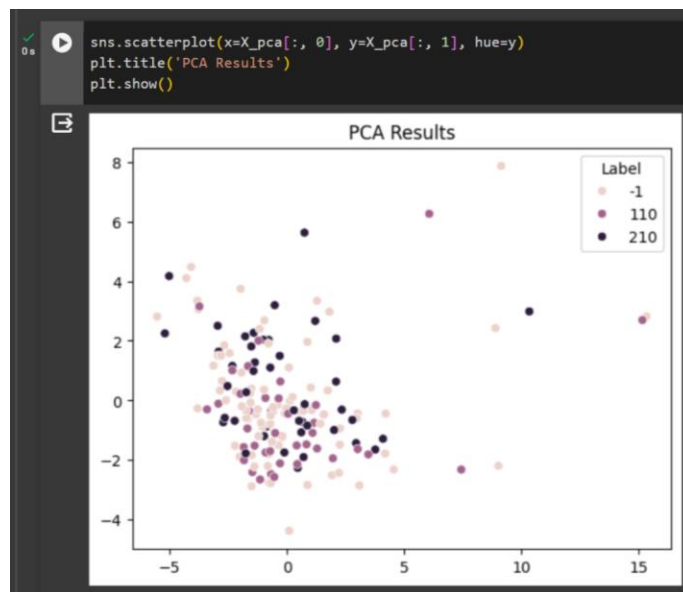
This line of code performs Principal Component Analysis (PCA) on the standardized data `X_scaled` to reduce its dimensionality to 3 principal components.

```
0 s pca = PCA(n_components=3)\nX_pca = pca.fit_transform(X_scaled)
```

`X_pca` will contain the transformed data with reduced dimensionality, where each sample is represented by three principal components.

18. Visualize PCA results

This segment generates a scatter plot to visualize the results of PCA, where each point is colored based on the corresponding emotional state.



19. Split the dataset into training and testing sets

The dataset is split into training and testing sets using `train_test_split` from `scikit-learn`.

```
✓ [16] X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
```

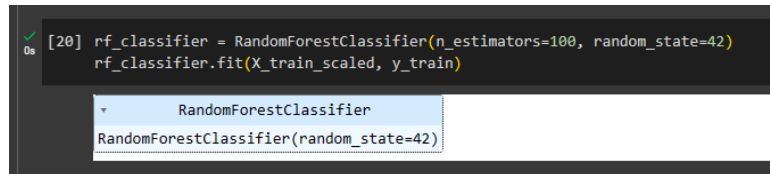
20. Standardize the features

Features in the training and testing sets are standardized using `StandardScaler`.

```
✓ [19] scaler = StandardScaler()  
      X_train_scaled = scaler.fit_transform(X_train)  
      X_test_scaled = scaler.transform(X_test)
```

21. Build a Random Forest Classifier

A Random Forest Classifier is instantiated, trained on the scaled training data.



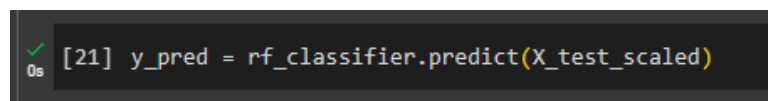
```
[20] rf_classifier = RandomForestClassifier(n_estimators=100, random_state=42)
      rf_classifier.fit(X_train_scaled, y_train)
```

The screenshot shows a Jupyter Notebook cell with the above code. Below the code, the variable `rf_classifier` is expanded, showing its type as `RandomForestClassifier` and its parameters as `RandomForestClassifier(random_state=42)`.

- `X_train_scaled` is the feature matrix of the training data, where the features have been scaled using `StandardScaler`.
- `y_train` is the target variable (labels) corresponding to the training data.

22. Predict on the test set

This line of code predicts the labels for the test set using the trained Random Forest Classifier model.



```
[21] y_pred = rf_classifier.predict(X_test_scaled)
```

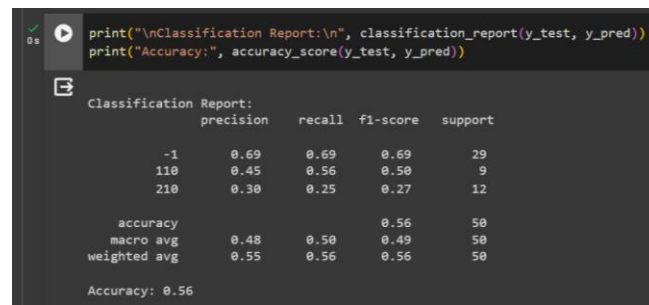
The screenshot shows a Jupyter Notebook cell with the above code. A green checkmark and '0s' are visible to the left of the code line.

`y_pred` will contain the predicted labels for the test set based on the input features `X_test_scaled`

23. Evaluate the model

This segment prints the classification report and accuracy score to evaluate the performance of the Random Forest Classifier on the test set. The classification report provides metrics such as precision, recall, and F1-score for each class.

It compares the true labels (`y_test`) to the predicted labels (`y_pred`) and computes these metrics for each class.



```
print("\nClassification Report:\n", classification_report(y_test, y_pred))
print("Accuracy:", accuracy_score(y_test, y_pred))
```

Classification Report:

	precision	recall	f1-score	support
-1	0.69	0.69	0.69	29
110	0.45	0.56	0.50	9
210	0.30	0.25	0.27	12
accuracy			0.56	50
macro avg	0.48	0.50	0.49	50
weighted avg	0.55	0.56	0.56	50

Accuracy: 0.56

24. Confusion Matrix

This segment computes the confusion matrix using `confusion_matrix` from scikit-learn and visualizes it as a heatmap using seaborn. The confusion matrix helps assess the model's performance in terms of true positives, true negatives, false positives, and false negatives for each class.



25. Use SVM with cross-validation

Initialize a Support Vector Machine (SVM) classifier with a linear kernel

```
✓ 0 s [28] svm_classifier = SVC(kernel='linear', random_state=42)
```


26. Perform cross-validation using X_standardized

`cross_val_score` is a function from scikit-learn that performs cross-validation by splitting the dataset into 'cv' (here, 5) consecutive folds. Each fold is then used once as a validation while the $k - 1$ remaining folds form the training set.

```
✓ 0 s ▶ cv_scores = cross_val_score(svm_classifier, x_scaled, y, cv=5)
print("Cross-Validation Scores:", cv_scores)
print("Mean Accuracy:", cv_scores.mean())

📄 Cross-Validation Scores: [0.55882353 0.48484848 0.57575758 0.54545455 0.51515152]
Mean Accuracy: 0.5360071301247772
```


27. Fit SVM model on the entire dataset

```
0s ✓  svm_classifier.fit(x_scaled, y)
```

SVC

SVC(kernel='linear', random_state=42)

28. Split data into training and testing sets

```
0s ✓  X_train, X_test, y_train, y_test = train_test_split(x_scaled, y, test_size=0.1, random_state=42)
```

29. Predict using the trained model

```
0s ✓ [34] y_pred_svm = svm_classifier.predict(X_test)

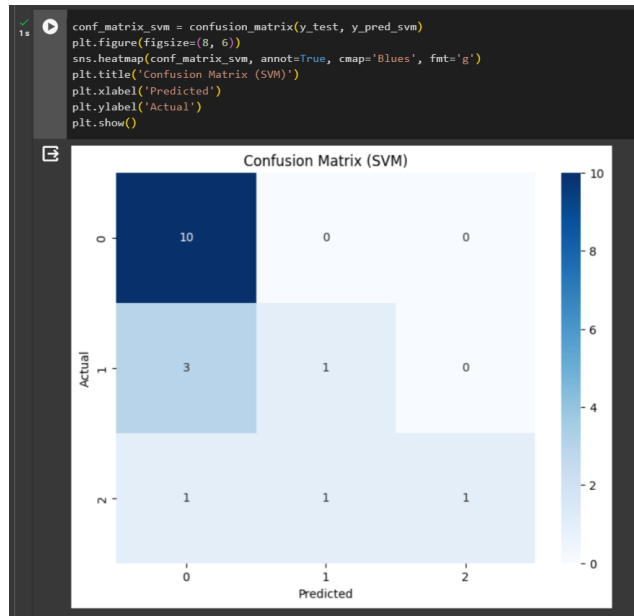
print("\nClassification Report (SVM):\n", classification_report(y_test, y_pred_svm))
print("Accuracy (SVM):", accuracy_score(y_test, y_pred_svm))
```

Classification Report (SVM):

	precision	recall	f1-score	support
-1	0.71	1.00	0.83	10
110	0.50	0.25	0.33	4
210	1.00	0.33	0.50	3
accuracy			0.71	17
macro avg	0.74	0.53	0.56	17
weighted avg	0.71	0.71	0.66	17

Accuracy (SVM): 0.7058823529411765

30. Confusion Matrix



CONCLUSION

In summary, the Random Forest classifier achieved an accuracy of 56%, with relatively lower precision, recall, and F1-score values for the emotional states other than the baseline (-1).

The SVM classifier performed slightly better with an accuracy of 71%, showing higher precision, recall, and F1-score values across all classes. However, it still faces challenges in accurately predicting certain emotional states.

Overall, while both models show promise, there's room for improvement. Fine-tuning parameters and exploring additional features could enhance their predictive power.

REFERENCES

1. Scikit-learn Documentation:

- Link: <https://scikit-learn.org/stable/documentation.html>

2. Seaborn Documentation:

- Link: <https://seaborn.pydata.org/tutorial.html>

3. Matplotlib Documentation:

- Link: <https://matplotlib.org/stable/contents.html>

4. Pandas Documentation:

- Link: <https://pandas.pydata.org/docs/>

