# Assignment 2: Naive Bayes Classifier

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# Preparatory remarks

To start of naive bayes classification we utilized the methods provided in the assignment description to read the data and split it into training and validation sets. We then trained the model using the training data and then classified the validation data. We then evaluated the classifier using the accuracy, precision, recall, and F1 score.

### Frequency-counting in Python.

```
from collections import Counter
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
def read documents(doc file):
    Reads a document and returns a list of documents and their
corresponding labels.
    docs = []
    labels = []
    categories = []
    with open(doc_file, encoding='utf-8') as f:
        for line in f:
            words = line.strip().split()
            docs.append(words[3:])
            labels.append(words[1])
            categories.append(words[0]) # Used in domain sensitivity
section
    return docs, labels, categories
```

# Reading the review data

```
def split_data(doc_file, train_split=0.8):
```

```
Splits the data into training and validation sets.

all_docs, all_labels, all_categories = read_documents(doc_file)
    split_point = int(train_split * len(all_docs))
    train_docs = all_docs[:split_point]
    train_labels = all_labels[:split_point]
    val_docs = all_docs[split_point:]
    val_labels = all_labels[split_point:]
    return train_docs, train_labels, val_docs, val_labels, all_docs,
all_labels, all_categories
```

## Estimating parameters for the Naive Bayes classifier

```
def train nb(documents, labels):
    Trains a Naive Bayes classifier given the documents and labels.
    Returns a model containing log probabilities.
    # Initialize counters for each class and a counter for all words
    pos counter = Counter()
    neg counter = Counter()
    all words = set()
    # Count word frequencies per class
    for doc, label in zip(documents, labels):
        if label == 'pos':
             pos counter.update(doc)
        elif label == 'neg':
             neg counter.update(doc)
        all words.update(doc)
    # Total count of words in each class
    total pos = sum(pos counter.values())
    total neg = sum(neg counter.values())
    # Vocabulary size
    V = len(all_words)
    # Calculate log probabilities with Laplace smoothing
    log probs = {}
    for word in all words:
        # Apply laplace smoothing
        \log \text{ prob pos} = \text{np.log}((\text{pos counter}[\text{word}] + 1) / (\text{total pos} + 1))
V))
        \log \text{ prob neg} = \text{np.log}((\text{neg counter}[\text{word}] + 1)) / (\text{total neg} + 1)
V))
        log probs[word] = {'pos': log prob pos, 'neg': log prob neg}
    # Calculate the log probabilities of each class
    num pos = sum(1 for l in labels if l == 'pos')
```

```
num neg = sum(1 for l in labels if l == 'neg')
    prob pos = num pos / len(labels)
    prob neg = num neg / len(labels)
    log prob pos class = np.log(prob pos)
    log prob neg class = np.log(prob neg)
    log prob class = {'pos': log prob pos class, 'neg':
log prob neg class}
    return log probs, log prob class
# Splitting the data and then training the model using the training
data
train docs, train labels, val docs, val labels, all docs, all labels,
all categories = split data('reviews.txt')
model, log_prob_class = train_nb(train_docs, train_labels)
# Print the first 5 instances and their log probabilities
print("Model Instances:")
for word, probs in list(model.items())[:5]:
    print(f"Word: '{word}'\n Pos Log Prob: {probs['pos']}\n Neg Log
Prob: {probs['neg']}\n")
print("Class Log Probabilities:")
for class label, class log prob in log prob class.items():
    print(f"Class '{class label}' Log Prob: {class log prob}")
Model Instances:
Word: 'brillant'
  Pos Log Prob: -12.172112471446578
 Neg Log Prob: -13.55064356428109
Word: 'faile'
  Pos Log Prob: -13.558406832566469
 Neg Log Prob: -12.857496383721145
Word: 'recommened'
  Pos Log Prob: -12.172112471446578
 Neg Log Prob: -13.55064356428109
Word: 'evade'
  Pos Log Prob: -12.865259652006523
 Neg Log Prob: -13.55064356428109
Word: '$35.00'
  Pos Log Prob: -13.558406832566469
 Neg Log Prob: -12.857496383721145
Class Log Probabilities:
```

```
Class 'pos' Log Prob: -0.6761896870922498
Class 'neg' Log Prob: -0.7103971982200179
```

#### Interpretation of results

These results show likelihood scores for words in two classes, 'pos' (positive) and 'neg' (negative). For example, 'blige' is more likely in 'pos' with a log probability of -12.46 compared to -13.55 in 'neg.' 'High-definition' and 'vat' also lean 'pos,' while '...world' and 'akim' have lower 'pos' probabilities. Overall, the model slightly favors 'pos' but not strongly.

# Classifying new documents

```
def score doc label(document, label, model, log prob class):
    Computes logarithm probability of the observed words in a document
given a sentiment label.
    # Start with the log probability of the label
    log_prob = log_prob_class[label]
    # Add the log probability of each word in the document
    for word in document:
        if word in model:
            log prob += model[word][label]
        else:
            # If the word is not in the model we choose to ignore it
            pass
    return log prob
# Sanity Check 1: Testing with the word "great"
log_prob_pos_great = score_doc_label(["great"], "pos", model,
log prob class)
log prob neg great = score doc label(["great"], "neg", model,
log prob class)
prob pos great = np.exp(log prob pos great)
prob_neg_great = np.exp(log_prob_neg_great)
# Sanity Check 2: Testing with the word "bad"
log prob pos bad = score doc label(["bad"], "pos", model,
log prob class)
log prob neg bad = score doc label(["bad"], "neg", model,
log prob class)
prob pos bad = np.exp(log prob pos bad)
prob_neg_bad = np.exp(log_prob_neg_bad)
# Sanity Check 3: Testing with the document ['a', 'top-quality',
'performance']
log prob pos doc = score doc label(['a', 'top-quality',
'performance'], "pos", model, log_prob_class)
```

```
log_prob_neg_doc = score_doc_label(['a', 'top-quality',
'performance'], "neg", model, log prob class)
print("Sanity Check 1:")
print(f"Probability of 'great' given positive class:
{prob pos great}")
print(f"Probability of 'great' given negative class:
{prob neg great}")
print("\nSanity Check 2:")
print(f"Probability of 'bad' given positive class: {prob_pos_bad}")
print(f"Probability of 'bad' given negative class: {prob neg bad}")
print("\nSanity Check 3:")
print(f"Log probability of ['a', 'top-quality', 'performance'] given
positive class: {log prob pos doc}")
print(f"Log probability of ['a', 'top-quality', 'performance'] given
negative class: {log prob neg doc}")
Sanity Check 1:
Probability of 'great' given positive class: 0.0013212141496043825
Probability of 'great' given negative class: 0.0005283997934747295
Sanity Check 2:
Probability of 'bad' given positive class: 0.00017230368700664423
Probability of 'bad' given negative class: 0.0004547440646873432
Sanity Check 3:
Log probability of ['a', 'top-quality', 'performance'] given positive
class: -12.807858361140351
Log probability of ['a', 'top-quality', 'performance'] given negative
class: -13.486891735775352
```

#### Interpretation of results

In our Naive Bayes test, the word 'great' showed up more in positive reviews, and 'bad' was more common in negative ones. It was interesting the classifier pick up on these vibes correctly. Also we can see that when we tested the document ['a', 'top-quality', 'performance'] the log probability of the positive class was higher than the negative class.

```
def classify_nb(document, model, log_prob_class):
    Classify a new document using the Naive Bayes classifier.
    # Compute the log probability for each class
    log_prob_pos = score_doc_label(document, "pos", model,
log_prob_class)
    log_prob_neg = score_doc_label(document, "neg", model,
log_prob_class)
```

```
# Return the class with the higher log probability
    if log prob pos > log prob neg:
        return "pos"
    else:
        return "neg"
# Sanity checks on small test documents
test docs = [["great"], ["bad"], ["amazing"], ["terrible"], ['a',
'top-quality', 'performance']]
# Applying the classify nb function to the test documents
classified docs = []
for doc in test docs:
    classification = classify nb(doc, model, log prob class)
    classified docs.append((doc, classification))
# Printing the results
print("Classified Documents:")
for doc, classification in classified docs:
    formatted_doc = ' '.join(doc)
    print(f"Document: '{formatted doc}' → Classification:
{classification}")
Classified Documents:
Document: 'great' → Classification: pos
Document: 'bad' → Classification: neg
Document: 'amazing' → Classification: pos
Document: 'terrible' → Classification: neg
Document: 'a top-quality performance' → Classification: pos
```

# Evaluating the classifier

```
def classify_documents(docs, model, log_prob_class):
    Classifies documents in the provided collection. Handling both
textual and numerical data.

predictions = []
for doc in docs:
    prediction = classify_nb(doc, model, log_prob_class)
    predictions.append(prediction)
    return predictions

def accuracy(true_labels, guessed_labels):
    Computes the accuracy of the classifier.
    """
    correct_count = 0
    for t, g in zip(true_labels, guessed_labels):
```

```
if t == q:
            correct count += 1
    acc = correct count / len(true labels)
    return acc
def precision_recall_f1(true_labels, predicted_labels):
    # Modified to handle multi-class labels
    unique labels = set(true_labels)
    precision, recall, f1 = \{\}, \{\}, \{\}
    for label in unique labels:
        true positives, false positives, false negatives = 0, 0, 0
        # Counting TP, FP, FN for each label
        for true, pred in zip(true labels, predicted labels):
            if true == label and pred == label:
                true positives += 1
            elif true != label and pred == label:
                false positives += 1
            elif true == label and pred != label:
                false negatives += 1
        # Calculate precision and recall for each label
        if true positives + false positives > 0:
            label precision = true positives / (true positives +
false positives)
        else:
            label precision = 0
        if true positives + false negatives > 0:
            label recall = true positives / (true positives +
false negatives)
        else:
            label recall = 0
        # Calculate F1 score for each label
        if label_precision + label_recall > 0:
            label f1 = 2 * (label precision * label recall) /
(label precision + label recall)
        else:
            label f1 = 00
        precision[label] = label precision
        recall[label] = label recall
        f1[label] = label f1
    return precision, recall, f1
```

```
# Calculating the accuracy, precision, recall, and F1 score
predicted_labels = classify_documents(val_docs, model, log_prob_class)
accuracy_result = accuracy(val_labels, predicted_labels)
precision, recall, f1_score = precision_recall_f1(val_labels,
predicted_labels)

print(f"Accuracy: {accuracy_result} \nPrecision: {precision['pos']}\nRecall: {recall['pos']}\nF1 score: {f1_score['pos']}")

Accuracy: 0.8153587914393622
Precision: 0.8237965485921889
Recall: 0.7866435385949696
F1 score: 0.8047914818101153
```

### What is the difference between F1 score and accuracy?

**Accuracy** Accuracy is a measures the proportion of correctly predicted instances out of all the predictions made. In short, it is the number of correct predictions divided by the total number of predictions. Accuracy gives us a general idea of how well the model is performing.

**Precision** F1 score is a slightly more nuanced metric. It is the harmonic mean of precision and recall. Precision is the ratio of true positives to all positive predictions. Recall is the ratio of true positives to all actual positive instances. F1 score balances precision and recall, providing a more holistic view of how well the model is performing especially when the classes are imbalanced.

### **Error Analysis**

We analyzed the first 10 documents and their corresponding prediction. Below you will see three instances which were harder to predict:

- 1. Document about an Iron (Predicted: Negative):
- Content: The document starts with a warning not to buy the iron, mentions it's fabulous when it works, but then highlights that it broke in less than a week and had other issues.
- Analysis: The prediction as negative seems correct. The initial positive remark "it's fabulous" probably was outweighed by the predominantly negative context.
- 1. Document about Madame Bovary (Predicted: Positive):
- Content: The review seems to be more about the characters in the story, with a mix of critique and appreciation for the adaptation and performances.
- Analysis: This document was tricky to label as it wrote polarizing thoughts about the book. We assume that the classifier picked up on more positive aspects than negative aspects of the book, leading to a positive prediction.
- 1. Document about Microsoft Office (Predicted: Negative):
- Content: The reviewer criticizes the new interface of Microsoft Office, finding it infuriating and less efficient.
- Analysis: The negative prediction aligns with the strong criticism expressed in the document.

Overall the classifier seemed to predict well the sentiment expressed in the first 10 documents. The more challenging cases for the classifier could be the ones with mixed sentiments or when

the sentiment is expressed in a not so straightforward way such as the Madame Bovary document.

#### Cross-validation

```
def cross validation 10 fold(all docs, all labels, N=10):
    Performs 10-fold cross-validation.
    accuracy_scores = []
    for fold nbr in range(N):
        split point 1 = int(float(fold nbr)/N*len(all docs))
        split point 2 = int(float(fold nbr+1)/N*len(all docs))
        train docs fold = all docs[:split point 1] +
all docs[split point_2:]
        train labels fold = all labels[:split point 1] +
all_labels[split_point_2:]
        val docs fold = all docs[split point 1:split point 2]
        val labels fold = all labels[split point 1:split point 2]
        model, log prob class = train nb(train docs fold,
train labels fold)
        predicted labels = classify documents(val docs fold, model,
log prob class)
        accuracy scores.append(accuracy(val labels fold,
predicted labels))
    return sum(accuracy scores) / N
def cross_validation_loocv(all docs, all labels, max iterations=100):
    Performs leave-one-out cross-validation.
    accuracy scores = []
    for i in range(min(max iterations, len(all docs))):
        train docs fold = all docs[:i] + all docs[i+1:]
        train labels fold = all_labels[:i] + all_labels[i+1:]
        val docs fold = [all docs[i]]
        val labels fold = [all labels[i]]
        model, log prob class = train nb(train docs fold,
train labels fold)
        predicted labels = classify documents(val docs fold, model,
log prob class)
        accuracy scores.append(accuracy(val labels fold,
predicted labels)
    return sum(accuracy scores) / len(accuracy scores)
```

```
# 10-Fold Cross-Validation
avg_accuracy_10_fold = cross_validation_10_fold(all_docs, all_labels,
N=10)
print("Average Accuracy (10-Fold Cross-Validation):",
avg_accuracy_10_fold)

# LOOCV
avg_accuracy_loocv = cross_validation_loocv(all_docs, all_labels,
max_iterations=100)
print("Average Accuracy (LOOCV, first 100 iterations):",
avg_accuracy_loocv)

Average Accuracy (10-Fold Cross-Validation): 0.8076216900805256
Average Accuracy (LOOCV, first 100 iterations): 0.83
```

#### Cross-validation results

- **10-Fold Cross-Validation**: Achieved an average accuracy of about 0.808. This method offers a balanced evaluation, as it trains and validates across different subsets of the data resulting in a consistent classifier performance.
- LOOCV (100 Iterations): Showed a slightly higher average accuracy of 0.83. By training on nearly all available data for each test case, this method offers a more accurate evaluation of the classifier performance. However, it is computationally expensive and can be slow to run.

The 10-fold and LOOCV results indicate a reliable classifier, with 10-fold offering a broader accuracy measure and LOOCV providing more detailed validation, however with higher computational demands.

## Domain sensitivity

```
def filter_by_category(docs, labels, categories, category_name):
    """"
    Filters the reviews by a specific category.
    filtered_docs = []
    filtered_labels = []

    for doc, label, cat in zip(docs, labels, categories):
        if cat == category_name:
            filtered_docs.append(doc)
            filtered_labels.append(label)

    return filtered_docs, filtered_labels

# Filter documents by category
camera_docs, camera_labels = filter_by_category(all_docs, all_labels, all_categories, 'camera')
book_docs, book_labels = filter_by_category(all_docs, all_labels, all_categories, 'books')
```

#### Split the data into training and validation sets

#### Train the classifier on the camera and book reviews.

```
# Train classifier for camera reviews
camera_model, camera_log_prob_class = train_nb(camera_train_docs,
camera_train_labels)
# Evaluate classifier on camera validation set
camera predicted labels = classify documents(camera val docs,
camera model, camera log prob class)
camera accuracy = accuracy(camera val labels, camera predicted labels)
# Train classifier for book reviews
book model, book log prob class = train nb(book train docs,
book train labels)
# Evaluate classifier on book validation set
book_predicted_labels = classify_documents(book_val_docs, book_model,
book log prob class)
book accuracy = accuracy(book val labels, book predicted labels)
# Evaluate the classifiers on the opposite domain
camera on book accuracy = accuracy(book val labels,
classify documents(book val docs, camera model,
camera log prob class))
book on camera accuracy = accuracy(camera val labels,
classify_documents(camera_val_docs, book_model, book log prob class))
# Print the results
print(f"Camera Classifier on Camera Reviews Accuracy:
{camera accuracy}")
print(f"Book Classifier on Book Reviews Accuracy: {book accuracy}")
```

```
print(f"Camera Classifier on Book Reviews Accuracy:
{camera_on_book_accuracy}")
print(f"Book Classifier on Camera Reviews Accuracy:
{book_on_camera_accuracy}")

Camera Classifier on Camera Reviews Accuracy: 0.8675
Book Classifier on Book Reviews Accuracy: 0.7925
Camera Classifier on Book Reviews Accuracy: 0.71
Book Classifier on Camera Reviews Accuracy: 0.65
```

## Classifier Performance Analysis

#### Within-Domain Accuracy

- Camera Classifier on Camera Reviews: Achieved an accuracy of 0.8675. The high accuracy indicates that the classifier is well-tuned to the specific features and patterns present in camera reviews.
- **Book Classifier on Book Reviews**: Accomplished an accuracy of 0.7925. While slightly lower than the camera classifier, it still shows a strong performance suggesting a good model fit to book review data.

#### Cross-Domain Accuracy

- Camera Classifier on Book Reviews: The accuracy dropped to 0.71 when the camera classifier was applied to book reviews. This significant drop in accuracy highlights the domain sensitivity. In turn indicating that features learned from camera reviews are not as effective in classifying book reviews.
- **Book Classifier on Camera Reviews**: Saw an even further drop in accuracy to 0.65. Similarly to the camera classifier, the book classifier was not able to generalize well to the camera reviews.

#### Conclusion

The results demonstrate a clear importance of domain sensitivity. Both classifiers perform well within their respective domains showing that they have learned domain-specific features effectively. However, when applied to a different domain their accuracy drops significantly demonstrating the importance of domain-specific language and features in text classification tasks.

# Naive Bayes for numerical data

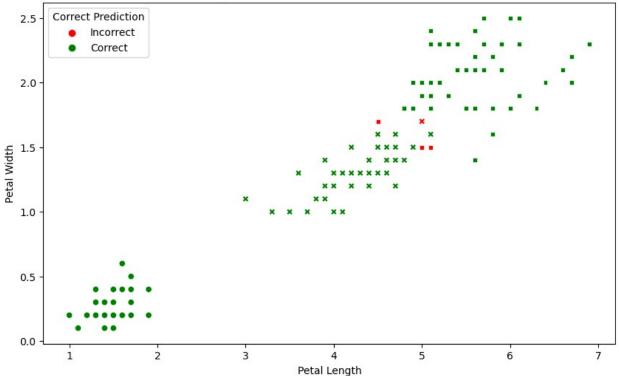
```
# Read the data
df = pd.read_csv('iris.csv')

def train_nb_numerical(dataframe, labels):
    Trains a Gaussian Naive Bayes classifier and returns model and log
probabilities of classes.
    # Initialize dictionary and get the unique labels
    model = {}
```

```
unique labels = dataframe[labels].unique()
    # Calculate the total number of samples
    total samples = len(dataframe)
    # Initialize dictionary for log probabilities of each class
    log prob class = {}
    # Calculate the mean, standard deviation, and log probability for
each class
    for label in unique labels:
        class data = dataframe[dataframe[labels] == label]
        features = class data.drop(labels, axis=1)
        mean = features.mean(numeric only=True)
        std = features.std(numeric only=True)
        # Store mean and std in the model
        model[label] = {'mean': mean, 'std': std}
        # Calculate and store the log probability of the class
        log prob class[label] = np.log(len(class data) /
total samples)
    return model, log prob class
def score doc label numerical(document, label, model):
    Computes the log probability of a numerical document given a
label.
    mean = model[label]['mean']
    std = model[label]['std']
    log prob = 0
    # Calculate the log probability for each feature
    for feature value, feature mean, feature std in zip(document,
mean, std):
        # Apply the Gaussian Naive Bayes formula
        log_prob += np.log(1 / (np.sqrt(2 * np.pi) * feature_std)) -
((feature value - feature mean) ** 2) / (2 * feature std ** 2)
    return log prob
def classify nb numerical(document, model):
    Classify a numerical document using a Naive Bayes model.
    # Initialize a dictionary to store log probabilities for each
label
    log probs = {}
```

```
# Compute the log probability for each class
    for label in model.keys():
        log probs[label] = score doc label numerical(document, label,
model)
    # Return the label with the highest log probability
    return max(log probs, key=log probs.get)
# Train the model
model, num_log_prob_class = train_nb_numerical(df, 'species')
# Initialize an empty list to store predictions
predictions = []
# Iterate over the rows in the dataframe
for index, row in df.iterrows():
    iris_document = row.drop('species').tolist()
    # Compute the log probability for each class
    log probs = {label: score doc label numerical(iris document,
label, model) for label in model}
    predicted class = max(log probs, key=log probs.get)
    predictions.append(predicted class)
# Add the predictions to the dataframe
df['predicted species'] = predictions
df['correct prediction'] = df['species'] == df['predicted species']
# Plot the correct and incorrect predictions
plt.figure(figsize=(10, 6))
sns.scatterplot(data=df, x='petal length', y='petal width',
hue='correct prediction', style='species', palette={True: "green",
False: "red"})
plt.title('Iris Species Predictions: Correct vs Incorrect')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
handles, = plt.gca().get legend handles labels()
plt.legend(handles=handles[1:], labels=['Incorrect', 'Correct'],
title='Correct Prediction', loc='upper left')
plt.show()
```

#### Iris Species Predictions: Correct vs Incorrect



```
def cross validation 10 fold numerical(dataframe, label column, N=10):
    Performs 10-fold cross-validation on a DataFrame with numerical
data.
    accuracy_scores, all_precision_scores, all_recall_scores,
all f1 scores = [], [], [],
    n = len(dataframe)
    # Shuffle the DataFrame
    dataframe = dataframe.sample(frac=1,
random state=1).reset index(drop=True)
    for fold nbr in range(N):
        split_point_1 = int(fold_nbr / N * n)
        split_point_2 = int((fold nbr + 1) / N * n)
        # Splitting DataFrame into training and validation sets
        train df = pd.concat([dataframe.iloc[:split point 1],
dataframe.iloc[split point 2:]])
        val_df = dataframe.iloc[split_point_1:split_point_2]
        # Training the model
        model, num log prob class = train nb numerical(train df,
label column)
```

```
# Initialize an empty list to store predictions
        predicted labels = []
        # Extracting validation labels
        val labels = val df[label column].tolist()
        # Classifying each document in the validation set
        for index, row in val df.iterrows():
            # Select the feature columns
            document = row[['sepal_length', 'sepal_width',
'petal length', 'petal width']].tolist()
            # Classify the document
            predicted_label = classify_nb_numerical(document, model)
            predicted labels.append(predicted label)
        # Calculating accuracy for the fold
        accuracy scores.append(accuracy(val labels, predicted labels))
        # Calculate precision, recall, and F1 scores for the fold
        precision, recall, f1 = precision recall f1(val labels,
predicted labels)
        all precision scores.append(precision)
        all recall scores.append(recall)
        all f1 scores.append(f1)
    # Calculate the average scores
    average_accuracy = sum(accuracy_scores) / N
    num labels = len(set(dataframe[label column]))
    average precision = sum(sum(precision.values()) for precision in
all precision scores) / (N * num labels)
    average_recall = sum(sum(recall.values()) for recall in
all recall scores) / (N * num labels)
    average f1 = sum(sum(f1.values()) for f1 in all f1 scores) / (N *
num labels)
    return average accuracy, average precision, average recall,
average f1
average accuracy, average precision, average recall, average f1 =
cross validation 10 fold numerical(df, 'species')
print(f"Average Accuracy: {average accuracy}")
print(f"Average Precision: {average_precision}")
print(f"Average Recall: {average recall}")
print(f"Average F1 Score: {average f1}")
Average Accuracy: 0.953333333333333
Average Precision: 0.947222222222222
```

Average Recall: 0.9523809523809524 Average F1 Score: 0.945929625929626

It seems like the Gaussian Naive Bayes classifier is performing well on the iris dataset, achieving a high average accuracy of 95.33% and F1 score of 94.59% over the 10-fold cross-validation. Indicating that the model is effectively learning and making accurate predictions on the given dataset.

- The high accuracy indicates that the classifier is effective in correctly predicting the species of the iris flowers. It suggests that the model generalizes well on the validation data, which is a positive outcome.
- The F1 score as it is mentioned early is a metric that considers both precision and recall. The high F1 score suggests a good balance between precision and recall, indicating that the model performs well in correctly identifying positive instances while minimizing false positives and false negatives.
- Gaussian Naive Bayes assumes that the features follow a Gaussian distribution, the good performance might indicate that this assumption is reasonable for the iris dataset because the model is capturing the patterns in the data effectively. However, further analysis of the feature distributions could provide more insights into this assumption.
- The use of 10-fold cross-validation provides a robust evaluation of the model. It helps ensure that the model's performance is consistent across different subsets of the data, reducing the risk of overfitting.

## References:

• Laplace Smoothing: https://courses.engr.illinois.edu/cs447/fa2018/Slides/Lecture04.pdf