Part II: An Investigation into K-Nearest-Neighbours for Classification

4th January 2024

I will be investigating the K-Nearest-Neighbours Classification algorithm (KNN). I will be using it to perform classification of the IRIS dataset (9).

1 What is KNN classification?

KNN is a non-parametric, supervised learning classifier, which uses the distance between data-points to classify or predict the grouping of an individual point. Being non-parametric means that the algorithm makes no assumptions about the underlying data distribution. It should be noted that KNN can be used for both classification and regression; I will be focusing on classification. KNN is reliant on the idea that similar points will have similar labels or values, this is the main assumption of the model. The "K" in KNN, refers to the number of nearest neighbors that the algorithm considers when making a prediction. The basic idea is that if a majority of the "k"-nearest neighbors belong to a certain class (for classification), then the algorithm predicts that the new data point belongs to that class or has a similar value. It does this normally through calculating the Euclidean distance.

The Euclidean distance between two points $P = (p_1, p_2, ..., p_n)$ and $Q = (q_1, q_2, ..., q_n)$ in an n-dimensional space is calculated using the following formula:

Euclidean Distance =
$$\sqrt{\sum_{i=1}^{n} (p_i - q_i)^2}$$

In the case of KNN it uses this distance in the following way:

- 1. The distance is normally calculated between a test point and every point in the specified training sample, where each sample is represented as a vector. This is the main training phase of KNN. Unlike traditional model-based algorithms, KNN doesn't explicitly learn a model during training. Instead, it memorizes the positions of the entire training dataset and makes predictions based on the proximity of new data points to existing ones. (3)
- 2. The algorithm then identifies the "K"- nearest neighbours, i.e the points with the smallest Euclidean Distance from the point. I will later examine how we determine the "K".
- 3. It then gauges which class label (feature) is most common amongst these K-nearest-neighbours and assigns the point to this common class-label.

To be clear, Euclidean distance is the most commonly used distance metric, however in specific

cases other metrics such as Manhattan Distance (5)or Minkowski Distance (4) (amongst others) may be used.

KNN is well-suited for smaller to medium-sized datasets with non-linear patterns and classification challenges. It excels in instance-based learning, making predictions based on data similarity. Notable applications include anomaly detection, but it may be less effective in high-dimensional spaces, due to The Curse of Dimensionality (2).

2 Data Selection

I will be working with the Iris Data set. This is a world-famous dataset featured in R.A. Fisher's classic 1936 paper, "The Use of Multiple Measurements in Taxonomic Problems", and can also be found in many places. I will be importing it from the UCI machine learning repository (6), found on Kaggle:

https://www.kaggle.com/datasets/uciml/iris/data

2.1 Importing the data

```
[3]: import numpy as np
  import pandas as pd
  import seaborn as sns
  import matplotlib.pyplot as plt
  from sklearn.model_selection import train_test_split
  from sklearn.preprocessing import LabelEncoder
  from sklearn.metrics import accuracy_score, f1_score
  from sklearn.model_selection import GridSearchCV
  from sklearn.neighbors import KNeighborsClassifier

# Define k-NN classifier
knn = KNeighborsClassifier()

# Importing iris, I have mounted the csv file to my google colab session.
  iris = pd.read_csv('/content/drive/MyDrive/Iris.csv')
```

2.2 Data Exploration

```
[4]: #shape of dataset
shape = iris.shape

#Initial preview
head = iris.head(6)

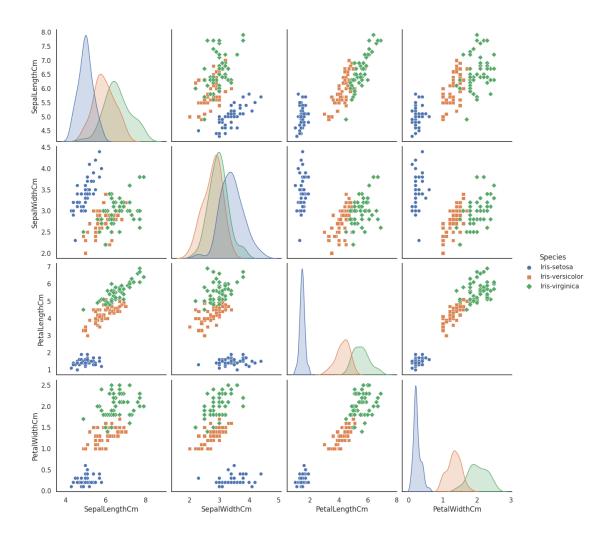
#how labels are split
label_split = iris.groupby('Species').size()

print(shape); print(head); print(label_split)
```

(150, 6)							
	Id	SepalLengt	hCm	${\tt SepalWidthCm}$	${\tt PetalLengthCm}$	${\tt PetalWidthCm}$	Species
0	1		5.1	3.5	1.4	0.2	Iris-setosa
1	2		4.9	3.0	1.4	0.2	Iris-setosa
2	3		4.7	3.2	1.3	0.2	Iris-setosa
3	4		4.6	3.1	1.5	0.2	Iris-setosa
4	5		5.0	3.6	1.4	0.2	Iris-setosa
5	6		5.4	3.9	1.7	0.4	Iris-setosa
Species							
Iris-setosa			50				
Iris-versicolor			50				
Iris-virginica			50				
dt	ype:	int64					

Initially, we can see that the dataset has 150 rows by 6 columns. This means there are 150 examples or entries for individual iris flowers. There is the one column indexing the flower, 4 columns of continuous data containing features (Sepal length, Sepal Width, Petal lenth and Petal Width) and the one column classifying the plants into a one of three specific species (setosa, versicolor and virginica). When eventually classifying, the Species column, containing categorical data, will act as labels.

The following pairplot showin in Figure 1 can be useful for visualising initial clusters.



For example, if I observe distinct clusters of points for different species in the scatterplots, this could indicate that the corresponding pair of features is informative for classifying the iris species. Here we can see that setosa features tend to be a distinct cluster when examining the relative features, whereas versicolor and virginica have a lot of overlap.

2.3 Train/Test Split

As is standard I will now split the data into training, validation and test sets; ready to undergo classification. Before doing this I split the data into features (X) and labels (y). I also perform label encoding on y. Label encoding is a way of taking a categorical variable that is represented non-numerically and converting it into a numerical format. Hence, Iris-setosa will correspond to 0, Iris-versicolor to 1 and Iris-virginica to 2.

```
[6]: # selecting features and labels arrays:
    X = iris.iloc[:, 1:5].values
    y = iris.iloc[:, 5].values

# Perform label encoding for the target variable y
label_encoder = LabelEncoder()
```

I have split the data into training, validation and test sets.

1. Training Set (X train, y train):

The model learns patterns and relationships in the data from this set. In the code, train_test_split is used to split the original dataset into training and test sets.

2. Validation Set (X val, y val):

The validation set is a subset of the training set, used to test different hyperparameters etc, to find optimal results. It prevents the test data from being used to train, hence avoiding contamination in the simulation.

3. Test Set (X test, y test):

The test set is composed of previously unseen data to the model. We use this after training and validation has been complete, to gauge how well the model works on data that is completely new.

3 Variance Investigation

Instead of building my own KNN function, I have opted to use *KNeighborsClassifier* from the sklearn.neighbours package.

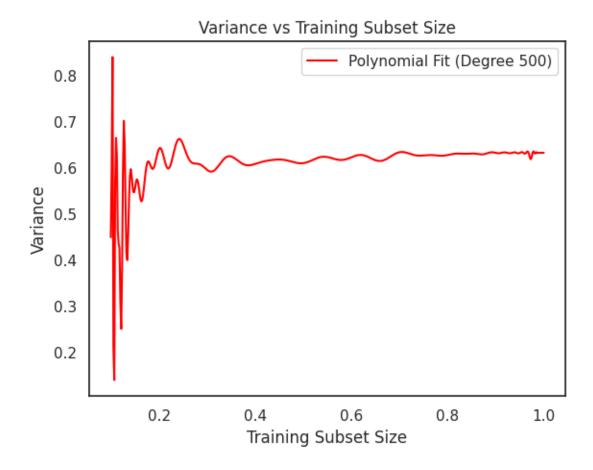
Later on I will investigate optimising hyperparamters and cross-validation techniques, but before I do, I will investigate the result of differing sizes of training data on the variance.

I do this by first creating a numpy array of evenly distributed subset sizes from 0.1 to 1, i.e 10%-100% of the training data size. I then create a for loop which, continuously makes random subsets of these sizes, trains the KNN classifier on this subset and then tests this on the validation set. I compute the variances for each predicted sample, take an average and append this to my variances list.

The plot below shows the distribution of variance as the subset size increases.

```
[9]: from sklearn.neighbors import KNeighborsClassifier
  import numpy as np
  import matplotlib.pyplot as plt
  from numpy.polynomial import Polynomial
```

```
#setting seed for reproducability
np.random.seed(0)
# Generate 500 evenly spaced subset sizes from 0.1 to 1, KNN does not work with _{\sqcup}
subsets = np.linspace(0.1, 1, 500)
subsets = np.array(subsets)
variances = []
for i in subsets:
      # Create a random subset of the training data
      subsets_random = np.random.choice(len(X_train), size=int(i *_
→len(X_train)), replace=False)
      X_subset = X_train[subsets_random]
      y_subset = y_train[subsets_random]
      # Train the KNN model
      knn_classifier = KNeighborsClassifier(n_neighbors=3)
      knn_classifier.fit(X_subset, y_subset)
      # Make predictions
      y_val_pred = knn_classifier.predict(X_val)
      # Compute the variance
      pred_variances = np.var(y_val_pred, axis=0)
      # Average the variances over the validation set
      average_variance = np.mean(pred_variances)
      variances.append(average_variance)
# Fit a polynomial to the data for a smoother curve
p = Polynomial.fit(subsets, variances, 500)
# Plotting the variance against subset sizes
plt.plot(subsets, p(subsets), color='red')
plt.title('Variance vs Training Subset Size')
plt.xlabel('Training Subset Size')
plt.ylabel('Variance')
plt.legend()
plt.show()
```



As shown by the above graph, the magnitude in the varying degree of variance, begins to settle at a variance value of around 0.6, as the subset size reaches 40% of its original size.

Initially, with very small subset sizes, the model might suffer from underfitting(8). The limited amount of data leads to high variability in the model's predictions on the validation set.

As the subset size increases, the model becomes more exposed to the training data, allowing it to capture more meaningful patterns. With a larger subset, the model's predictions on the validation set become more stable and consistent, resulting in a decrease in variance.

Beyond a certain subset size, increasing the size further may not provide substantial additional information for the model to improve its predictions and could lead to overfitting. The variance settles at a certain value, indicating that the model has reached a point of stability and additional data may not lead to significant changes in performance.

Hence, from this experiment we can conclude that the model has adequate data at around 40% of its original size.

4 Investigating KNN in more detail (Incorporating Question's 4 and 5)

Whilst the above validation experiment provides a solid insight into how we reduce variance amongst training results, it is basic and does not account for many things, for example the number of neighbours in that simulation was set at an arbitary value of 3 and not investigated.

I will now investigate how best to accurately train the model focusing on the "k" of the KNN as my hyperparameter to be optimised. I will do this through cross-validation and analyse the results on the validation and test sets through metrics of accuracy and F1-score.

4.1 Hyperparamter Tuning

In the context of k-nearest neighbors (KNN), the parameter "k" serves a pivotal role, as it determines the number of nearest neighbors considered when making predictions for new data points. This choice is integral due to the inherent tradeoff between bias and variance. A smaller value of "k" (e.g., 1 or 3) increases the complexity of the model and renders the algorithm more susceptible to noise and outliers, potentially resulting in overfitting. Conversely, a larger "k" (e.g., 10 or 20) imparts robustness by considering a broader neighborhood, but this may lead to underfitting as the decision boundaries become smoother and miss local patterns in the data. It is, therefore, of paramount necessity that k is optimised.

To do this I will be using a grid-search cross-validation technique. It is, therefore, necessary for me to define both Cross-Validation and Grid-search.

4.2 Cross Validation

Cross Validation involves splitting the data into many training and validation sets, like we did before, and then assessing the best result for numbers of k. I will be using a specific form, known as k-fold cross validation (1).

The method involves the following steps:

- 1. Choose a number of folds, denoted as k. Common choices for k include 5 or 10, although any number less than the dataset's length can be selected.
- 2. Split the dataset into k approximately equal parts, referred to as folds.
- 3. Select k-1 folds as the training set, with the remaining fold designated as the test set.
- 4. Train the model on the training set. For each iteration of cross-validation, a new model is trained independently of the models from previous iterations.
- 5. Validate the model on the test set.
- 6. Save the result of the validation.
- 7. Repeat steps 3 through 6 k times, using the remaining fold as the test set in each iteration. By the end, the model has been validated on every fold.

8. Calculate the final score by averaging the results obtained in step 6.

4.3 Grid Search

Grid Search is an exhaustive algorithm, that places all possible hyperparameters on a grid and then attempts every combination of them over a specified number of cross-validations, until the best score is returned (11). In the case of our grid search algorithm, we will be examining the Accuracy score.

4.4 Accuracy, Precision, Recall and F1-Score

Accuracy is defined as the ratio between the number of correct predictions to the total number of predictions. It is best applied when classes are of equal importance and should be used skeptically if the data is known to be imbalanced.

The precision measures the model's accuracy in classifying a sample as positive (10). It is the ratio between the True Positives and True Positives + False Positives.

Recall is only concerned with how many positive samples are correctly classified as positive. This means it is independent of negative samples being misclassified as positive.

Precision and Recall are combined in the F1 metric. This metric is perfect for when Accuracy falls down; when classes are imbalanced. It is calculated as the harmonic mean of Precision and Recall scores, which implies that if one of these two metrics are low then the F1-score will be correspondingly low as well.

Now from the earlier data analysis, we know that each of the 3 classes has 50 samples, so they are perfectly balanced. Even still, I will use the F1 score as a sanity check on my accuracy scores.

```
[10]: from sklearn.model_selection import GridSearchCV
    from sklearn.neighbors import KNeighborsClassifier

# Define k-NN classifier
knn = KNeighborsClassifier()

# Define hyperparameter grid
grid = {'n_neighbors': [1, 3, 5, 7, 9, 11, 13, 15]}

# Perform grid search with cross-validation
grid_search = GridSearchCV(knn, grid, cv=5, scoring='accuracy')
grid_search.fit(X_train, y_train)

# Get the best hyperparameter
best_k = grid_search.best_params_['n_neighbors']
print(best_k)
```

5

My grid search with cross-validation returns the optimal k value as 5. It should be noted that I only tested for odd values of k, as this is a common practice to avoid ties when determining which

class a data point should belong to.

I now train the KNN classifier with this optimal k and test it on my validation data.

```
[11]: # Train k-NN with the best hyperparameter
best_knn = KNeighborsClassifier(n_neighbors=best_k)
best_knn.fit(X_train, y_train)

# Predictions on the validation set
y_val_pred = best_knn.predict(X_val)

# Calculate accuracy and F1 score
accuracy = accuracy_score(y_val, y_val_pred)
f1 = f1_score(y_val, y_val_pred, average='weighted')

print('Accuracy:', accuracy, 'F1-score:',f1)
```

As shown above, a high accuracy score and a matchin f1 score are achieved. The next step will be to use this trained model on my test data. However, before doing this I need to be completely content with the model, or else I'll waste the usage of the test data. For this reason, I hold off testing until the final part of this report.

5 Further Investigation of Distance Metrics (Question 6)

As aforementioned, this experiment used KNN with Euclidean distance as the distance metric. This is the most commonly used distance metric, yet it is not the only one. Of course there are many different ways of calculating distance that could be used, giving different insights when used correctly. I would now like to examine, the other most popular ways of calculating distance.

5.1 Minkowski Distance

The Minkowski Distance is a normed vector space, which is a generalization of both the Euclidean Distance and Manhattan Distance(4). To be normed means distances are represented as a vector that has non-negative length and satisfies the following:

• positivity: |a, b| > 0

• triangle inequality: $|a+b| \le |a| + |b|$

• identity: $|a, b| = 0 \iff a == b$

• symmetry: |a, b| = |b, a|

The formula for the Minkowski Distance is as follows:

For two points, (x_1, y_1) and (x_2, y_2) in a two-dimensional space:

$$D(x, y, p) = (|x_1 - x_2|^p + |y_1 - y_2|^p)^{1/p}$$

which is generalised as:

$$D(\mathbf{X}, \mathbf{Y}, p) = \left(\sum_{i=1}^{n} |X_i - Y_i|^p\right)^{1/p}$$

As you can see, setting p = 2, results in the Euclidean distance and p = 1 results in the Manhattan Distance, which I will discuss next.

5.2 Manhattan Distance

The Manhattan Distance is the distance between two points defined to be the sum of the absolute differences of their Cartesian coordinates. It is often referred to as *Taxicab Geometry* (7), as it can be understood by placing all points on a two-dimensional grid and finding the shortest, non-diagonal path to traverse between two points, much like a taxicab would in the Island of Manhattan.

It is defined mathematically below;

$$D(\mathbf{X}, \mathbf{Y}) = \sum_{i=1}^{n} |X_i - Y_i|$$

The Manhattan Distance, is typically utilised in datasets with high-dimensionallity. This is because it does not take any squares and hence does not amplify the difference between features. ref.

5.3 Chebyshev Distance

Chebyshev distance, also referred to as chessboard distance, quantifies the distance between data points by assessing the greatest distance among their respective coordinates. It can also be derived by substituting ∞ into the Minkowski equation (12).

$$D(\mathbf{X}, \mathbf{Y}) = \max_{i=1}^{n} |X_i - Y_i|$$

5.4 Cosine Distance

The cosine distance is typically used to determine the dissimilarity between two vectors.

$$D(\mathbf{X}, \mathbf{Y}) = 1 - \frac{\mathbf{X} \cdot \mathbf{Y}}{\|\mathbf{X}\| \cdot \|\mathbf{Y}\|}$$

where

$$\cos\theta = \frac{\mathbf{X} \cdot \mathbf{Y}}{\|\mathbf{X}\| \cdot \|\mathbf{Y}\|}$$

It is defined mathematically above as the inverse of cosine similarity. Hence, a lower distance score indicates a higher similarity between two vectors. A common usage of cosine distance in in sentiment analysis, analysing the similarity between two corpuses of text.

5.5 Jaccard Distance

Jaccard Distance or Similarity, is another similarity measure, normally used to measure the similarity between two objects or sets.

$$D(\mathbf{A}, \mathbf{B}) = 1 - \frac{|\mathbf{A} \cap \mathbf{B}|}{|\mathbf{A} \cup \mathbf{B}|}$$

5.5.1 Results

Now that I have covered some of the more popular distance metrics, I will run some tests to demonstrate their applicability to certain situations.

From the above descriptions of the distance metrics, when using them as the specified distance metric for KNN, we would expect some variation in performance. We would expect, given the well-separated clusters of iris species in the feature space, that Euclidean distance would be the best metric for this analysis. We should also find that Manhattan, Chebyshev and Cosine Distances works well but do not offer significant advantages over Euclidean. As Jaccard is typically used in datasets, where an exclsusion of features matters, we might find that it is not suitable for this setting.

I adjust the above code to view how both the optimal number of k and the accuracies change for the KNN model, when experiementing with different distance metrics.

```
[14]: # Define a list of distance metrics to test
      distance_metrics = ['euclidean', 'manhattan', 'chebyshev', 'cosine', 'jaccard']
      # Iterate over distance metrics
      for j in distance_metrics:
          print(f"Testing with {j} distance:")
          # Define k-NN classifier
          knn = KNeighborsClassifier(metric=j)
          # Define hyperparameter grid
          grid_q6 = {'n_neighbors': [1, 3, 5, 7, 9, 11, 13, 15]}
          # Perform grid search with cross-validation
          grid_search = GridSearchCV(knn, grid_q6, cv=5, scoring='accuracy')
          grid_search.fit(X_train, y_train)
          # Get the best k
          best_k = grid_search.best_params_['n_neighbors']
          print(f"Best k for {j} distance: {best_k}")
          # Train k-NN with the best hyperparameter
```

```
best_knn = KNeighborsClassifier(n_neighbors=best_k, metric=j)
best_knn.fit(X_train, y_train)

# Predictions on the validation set
y_val_pred = best_knn.predict(X_val)

# Calculate accuracy
accuracy = accuracy_score(y_val, y_val_pred)
print(f"Accuracy for {j} distance: {accuracy}")
```

As expected the accuracies between Euclidean, Manhattan, Chebyshev and Cosine are practically the same, with Jaccard distance giving a very poor accuracy. What is interesting to note is the varying value of the best-k. The k-value can vary for many reasons. The ideal k value in KNN can differ for various distance metrics due to specific dataset characteristics. For instance, Euclidean distance, sensitive to spatial separation, might perform differently to Manhattan distance, which considers distances along coordinate axes.

I will store the models generated for Euclidean, Manhattan, Chebyshev and Jaccard and test them on the test data. This will be the ultimate test on which model is most suitable for the given dataset.

${\bf 5.5.2 \quad Euclidean, \, k_number} = {\bf 5}$

```
[15]: # Train k-NN with the best hyperparameter
best_knn = KNeighborsClassifier(n_neighbors=5, metric ='euclidean')
best_knn.fit(X_train, y_train)

# Predictions on the validation set
y_t_pred = best_knn.predict(X_test)

# Calculate accuracy and F1 score
```

```
accuracy = accuracy_score(y_test, y_t_pred)
f1 = f1_score(y_test, y_t_pred, average='weighted')
print('Accuracy:', accuracy, 'F1-score:',f1)
```

Accuracy: 0.9666666666666667 F1-score: 0.9664109121909632

5.5.3 Manhattan, k number = 9

```
[16]: # Train k-NN with the best hyperparameter
best_knn = KNeighborsClassifier(n_neighbors=9, metric = 'manhattan')
best_knn.fit(X_train, y_train)

# Predictions on the validation set
y_t_pred = best_knn.predict(X_test)

# Calculate accuracy and F1 score
accuracy = accuracy_score(y_test, y_t_pred)
f1 = f1_score(y_test, y_t_pred, average='weighted')

print('Accuracy:', accuracy, 'F1-score:',f1)
```

Accuracy: 1.0 F1-score: 1.0

5.5.4 Chebyshev, k number = 3

```
[17]: # Train k-NN with the best hyperparameter
best_knn = KNeighborsClassifier(n_neighbors=3, metric ='chebyshev')
best_knn.fit(X_train, y_train)

# Predictions on the validation set
y_t_pred = best_knn.predict(X_test)

# Calculate accuracy and F1 score
accuracy = accuracy_score(y_test, y_t_pred)
f1 = f1_score(y_test, y_t_pred, average='weighted')

print('Accuracy:', accuracy, 'F1-score:',f1)
```

Accuracy: 0.9666666666666667 F1-score: 0.9664109121909632

5.5.5 Jaccard, k number = 1

```
[18]: # Train k-NN with the best hyperparameter
best_knn = KNeighborsClassifier(n_neighbors=1, metric ='jaccard')
best_knn.fit(X_train, y_train)
```

```
# Predictions on the validation set
y_t_pred = best_knn.predict(X_test)

# Calculate accuracy and F1 score
accuracy = accuracy_score(y_test, y_t_pred)
f1 = f1_score(y_test, y_t_pred, average='weighted')

print('Accuracy:', accuracy, 'F1-score:',f1)
```

The final results above exceed the expectations delivered from the validation set, with even higher accuracy results. Manhattan Distance performs the best with a perfect accuracy score. This can most probably be attributed to the fact that the Iris dataset is one of low-dimesions and Mnahattan diatnce is most commonly used for high-dimensional probelms. Hence, it is probably computational overkill for this situation, but nonetheless performs perfectly.

In this final part of the study I have investigated different distance metrics that can be used in the KNN classifier. The distance metric is just one of the many possible hyperparameters to focus on and optimise and in further studies it would be worth investigating others, such as weightings, in more detail. For the purposes of this, I have not investigated others, allowing for more detail on my chosen metrices.

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