Lecture 13

KNN, Distance metrics

Let's start with an overview of KNN, or K-Nearest Neighbors.

## What is KNN?

- K-Nearest Neighbors (KNN) is a simple, non-parametric, and lazy machine learning algorithm primarily used for classification tasks but can also be applied to regression.
- **Non-parametric** means that it does not assume a fixed form (or distribution) for the data, making it versatile for various types of datasets.
- Lazy algorithm implies that it makes no assumptions during the training phase and instead waits until a query point needs to be predicted.

# How Does KNN Work?

- Given a query point, KNN looks for the 'K' closest data points in the training dataset and uses their labels (in classification) or values (in regression) to make a prediction.
- The core idea: "Birds of a feather flock together." That is, similar instances are likely to have similar labels.

While KNN is typically deployed as a classification algorithm, it can be modified to work as a regression algorithm. We'll look at this case in a future lecture, but for the moment, we'll stick with the classification context.

We looked in a prior lecture at distance metrics for this and other related applications. The way that KNN determines closeness or similarity is done by distance metrics. Let's review some common choices used with KNN.

# What Are Distance Metrics?

• The concept of "closeness" is defined using distance metrics. Different distance metrics can lead to different results.

# **Common Distance Metrics:**

- Euclidean Distance: Most common and intuitive.  $d(x, y) = \sqrt{\sum_{i=1}^{n} (x_i y_i)^2}$
- Manhattan Distance: Sum of absolute differences.  $d(x, y) = \sum_{i=1}^{n} |x_i y_i|$
- *Minkowski Distance*: A generalization of both Euclidean and Manhattan.  $d(x, y) = \sqrt[p]{\sum_{i=1}^{n} |x_i y_i|^p}$
- Chebyshev Distance: The maximum difference along any coordinate dimension.  $d(x, y) = \max(|x_i y_i|)$
- Cosine Similarity: Measures the cosine of the angle between two vectors (used when the magnitude of the vectors matters less than their direction).  $d(x, y) = 1 \frac{x \cdot y}{\|x\| \|y\|}$

See the earlier lecture for additional options.

# Implementation Considerations:

The choice of distance metric should align with the nature of your data. For example, Euclidean distance works well for continuous data, while Hamming distance is more suitable for categorical data.

The next element to consider is the value of k: how many nearest neighbors will be considered? **Choosing the Value of 'K':** 

- The parameter 'K' determines the number of neighbors to consider.
  - Low 'K' value (e.g., K = 1): Can lead to overfitting, as the model might be too sensitive to noise in the training data.
  - *High 'K' value (e.g., K = total number of samples):* Can lead to underfitting, as the model becomes too generalized.
- Common practice: Use cross-validation to select an optimal value for 'K'.

Because of tie-breaking methods, it's common, particularly in binary classification, to avoid even values of k to avoid instability in the model. However, when classification is not binary, choosing the best value of k is more flexible, since ties can happen with any value of k except 1.

# Impact of Scaling:

- Distance metrics are sensitive to the scale of the data. Features with larger ranges will dominate the distance calculations.
- Standardization/Normalization:
  - *Standardization*: Transform features to have a mean of 0 and a standard deviation of 1.
  - *Normalization*: Transform features to fall within a specific range, typically [0, 1].

**Practical Consideration**: Before applying KNN, always scale your features to avoid bias in distance calculation.

Scaling variables (however they are scaled, it should be consistent) can have a huge impact on the quality of your model. It's fine to try it without scaling first, but always check scaling to look for improvement.

## Classification (using packages)

```
# Example classification with K = 3
knn_classification <- function(train_data, test_point, k = 3) {
    distances <- sqrt(rowSums((train_data[, -1] - test_point) ^ 2))
    neighbors <- train_data[order(distances), ][1:k, ]
    prediction <- names(sort(table(neighbors[, 1]), decreasing = TRUE))[1]
    return(prediction)
}</pre>
```

(This is a generic example. We'll look at the algorithm and an example in more detail.)

```
Regression (using packages)
```

```
# Example regression with K = 3
knn_regression <- function(train_data, test_point, k = 3) {
    distances <- sqrt(rowSums((train_data[, -1] - test_point) ^ 2))
    neighbors <- train_data[order(distances), ][1:k, ]
    prediction <- mean(neighbors[, 1])
    return(prediction)
}</pre>
```

The main difference between regression and classification is in the final step of the algorithm. In classification, the nearest neighbors "vote" on which class the new element belongs to, with the one with the most votes winning (subject to potential tiebreaking). In regression, the nearest neighbors are averaged together to get the regression prediction.

## Advantages of KNN:

- *Simplicity*: Easy to understand and implement.
- *Versatility:* Can be used for classification and regression.
- No Training Phase: Works directly on raw data, which can be advantageous in some scenarios.

## **Disadvantages:**

- *Computationally Expensive:* Particularly with large datasets, as distance calculations must be repeated for each prediction.
- *Memory Intensive:* Stores the entire training dataset.
- Sensitive to Noise: Outliers can significantly affect the prediction.

## Improvements:

- *KD-Trees or Ball Trees* for efficient neighbor searches.
- *Weighted KNN:* Assign weights to neighbors based on their distance, giving closer neighbors more influence on the prediction.

Let's start with classification and look at two examples, one where we employ the Euclidean distance metric and a second that uses a different one.

Euclidean distance:

```
# Prepare data
set.seed(123)
mtcars$am <- as.factor(mtcars$am) # Convert 'am' to a factor (for classification)</pre>
```

```
# Use 'hp', 'wt', and 'qsec' as predictors and 'am' as the target
data <- mtcars[, c("hp", "wt", "qsec", "am")]</pre>
```

# Scale the predictors
data[, 1:3] <- scale(data[, 1:3])</pre>

```
# Euclidean distance function
euclidean_distance <- function(x1, x2) {
   sqrt(sum((x1 - x2) ^ 2))
}</pre>
```

```
# KNN function using Euclidean distance
```

```
knn_euclidean <- function(train_data, test_point, k = 3) {
    distances <- apply(train_data[, -ncol(train_data)], 1, function(row) euclidean_distance(row,
    test_point))
    sorted_indices <- order(distances)
    nearest neighbors <- train data[sorted indices[1:k], ]</pre>
```

#### # Return the majority vote for classification

prediction <- names(sort(table(nearest\_neighbors[, ncol(nearest\_neighbors)]), decreasing =
TRUE))[1]</pre>

return(as.numeric(prediction))

}

## # Split the data into train and test sets

train\_data <- data[-1, ] # Use all rows except the first as training data test\_point <- data[1, -4] # Use the first row (without the target column) as the test point

# Predict using KNN with Euclidean distance
predicted\_value <- knn\_euclidean(train\_data, test\_point, k = 3)
predicted\_value</pre>

For the second case, let's try cosine similarity.

#### # Cosine similarity function

```
cosine_similarity <- function(x1, x2) {
    sum(x1 * x2) / (sqrt(sum(x1 ^ 2)) * sqrt(sum(x2 ^ 2)))
}</pre>
```

#### # Split the data into train and test sets

train\_data <- data[-1, ] # Use all rows except the first as training data test\_point <- data[1, -4] # Use the first row (without the target column) as the test point

#### # Predict using KNN with Euclidean distance

```
predicted_value <- knn_euclidean(train_data, test_point, k = 3)
predicted_value</pre>
```

## # KNN function using Cosine Similarity

```
knn_cosine <- function(train_data, test_point, k = 3) {
    similarities <- apply(train_data[, -ncol(train_data)], 1, function(row) cosine_similarity(row,
    test_point))
    sorted_indices <- order(-similarities) # Sort in decreasing order because higher cosine similarity
    is better
    nearest neighbors <- train data[sorted indices[1:k], ]</pre>
```

### # Return the majority vote for classification

```
prediction <- names(sort(table(nearest_neighbors[, ncol(nearest_neighbors)]), decreasing =
TRUE))[1]</pre>
```

```
return(as.numeric(prediction))
```

```
}
```

#### # Predict using KNN with Cosine similarity

```
predicted_value_cosine <- knn_cosine(train_data, test_point, k = 3)
predicted_value_cosine</pre>
```

In this example, the first row of mtcars was removed as a test point, and the model was built on the other available datapoints. Both methods of distance metrics correctly predicted the value of 1 for the am variable (compare with the first row of mtcars for verification).

We can also include metrics for comparison if we tweak this a bit. ROC and AUC are most easily done on binary problems, so let's look at those first, before we move on a case with three or more classes.

```
# Load the mtcars dataset
data(mtcars)
# Convert 'am' to a factor for classification
mtcars$am <- as.factor(mtcars$am)
# Select the numeric predictor variables and normalize them
mtcars[, c("mpg", "hp")] <- scale(mtcars[, c("mpg", "hp")])</pre>
# Split the data into training and testing sets (70% train, 30% test)
set.seed(123)
train indices <- sample(1:nrow(mtcars), 0.7 * nrow(mtcars))</pre>
train_data <- mtcars[train_indices, c("mpg", "hp", "am")]</pre>
test data <- mtcars[-train_indices, c("mpg", "hp", "am")]</pre>
euclidean_distance <- function(x1, x2) {</pre>
 sqrt(sum((as.numeric(x1) - as.numeric(x2)) ^ 2))
}
cosine_similarity <- function(x1, x2) {</pre>
 sum(as.numeric(x1) * as.numeric(x2)) / (sqrt(sum(as.numeric(x1)^2)) *
sqrt(sum(as.numeric(x2)^2)))
}
knn_classifier <- function(train_data, test_point, k = 3, distance_func) {</pre>
 distances <- apply(train_data[, -ncol(train_data)], 1, function(row) distance_func(row,
test_point))
 # Determine whether to sort distances ascending or descending
 if (identical(distance func, cosine similarity)) {
  nearest_neighbors <- order(distances, decreasing = TRUE)</pre>
 } else {
  nearest neighbors <- order(distances)</pre>
 }
 nearest_neighbors <- train_data[nearest_neighbors[1:k], "am"]</pre>
 # Majority voting
 prediction <- as.character(sort(table(nearest neighbors), decreasing = TRUE)[1])
 return(prediction)
}
```

```
predict_knn <- function(train_data, test_data, k = 3, distance_func) {
    predictions <- sapply(1:nrow(test_data), function(i) {
        knn_classifier(train_data, test_data[i, -ncol(test_data)], k, distance_func)
    })
    return(as.factor(predictions))
}</pre>
```

#### # Predictions using Euclidean distance

```
predictions_euclidean <- predict_knn(train_data, test_data, k = 3, distance_func =
euclidean_distance)</pre>
```

```
# Predictions using Cosine similarity
predictions_cosine <- predict_knn(train_data, test_data, k = 3, distance_func = cosine_similarity)</pre>
```

```
library(caret)
```

```
evaluate_model <- function(predictions, true_labels) {
    confusion <- confusionMatrix(predictions, true_labels)
    accuracy <- confusion$overall["Accuracy"]
    f1 <- confusion$byClass["F1"]</pre>
```

```
return(list(Confusion_Matrix = confusion$table, Accuracy = accuracy, F1_Score = f1))
}
```

library(caret)

```
evaluate_model <- function(predictions, true_labels) {
    # Ensure that predictions and true labels have the same levels
    levels(predictions) <- levels(true_labels)</pre>
```

```
# Calculate confusion matrix, accuracy, and F1 score
confusion <- confusionMatrix(predictions, true_labels)
accuracy <- confusion$overall["Accuracy"]
f1 <- confusion$byClass["F1"]</pre>
```

```
return(list(Confusion_Matrix = confusion$table, Accuracy = accuracy, F1_Score = f1))
}
```

```
# Euclidean results
```

```
euclidean_results <- evaluate_model(predictions_euclidean, test_data$am)
print("Euclidean Results")
print(euclidean_results)</pre>
```

## # Cosine results

```
cosine_results <- evaluate_model(predictions_cosine, test_data$am)
print("Cosine Results")
print(cosine_results)</pre>
```

### library(pROC)

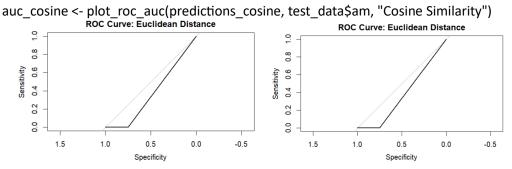
```
plot_roc_auc <- function(predictions, true_labels, title) {
  roc_obj <- roc(as.numeric(true_labels) - 1, as.numeric(predictions) - 1)
  auc_value <- auc(roc_obj)
  plot(roc_obj, main = paste("ROC Curve:", title))
  return(auc_value)</pre>
```

}

#### # ROC and AUC for Euclidean

auc\_euclidean <- plot\_roc\_auc(predictions\_euclidean, test\_data\$am, "Euclidean Distance")

#### # ROC and AUC for Cosine



Let's look at an example that generates some classification metrics for us, so that we can compare how well various distance metrics work. We'll also use the iris dataset which has three classes to work with.

#### # Load the dataset

data(iris) set.seed(123)

```
# Convert species to a factor (for classification)
iris$Species <- as.factor(iris$Species)</pre>
```

```
# Scale the predictors
iris[, 1:4] <- scale(iris[, 1:4])</pre>
```

```
# Split the data into training and testing sets (70% train, 30% test)
train_indices <- sample(1:nrow(iris), 0.7 * nrow(iris))
train_data <- iris[train_indices, ]</pre>
```

```
test_data <- iris[-train_indices, ]</pre>
```

```
# Manhattan distance function
manhattan_distance <- function(x1, x2) {
    sum(abs(x1 - x2))
}</pre>
```

```
# Minkowski distance function with p = 3
minkowski_distance <- function(x1, x2, p = 3) {
    sum(abs(x1 - x2) ^ p)^(1/p)
}</pre>
```

# # Chebyshev distance function

```
chebyshev_distance <- function(x1, x2) {
  max(abs(x1 - x2))
}</pre>
```

```
# General KNN function
```

```
knn_general <- function(train_data, test_point, k = 3, distance_func) {
    distances <- apply(train_data[, -ncol(train_data)], 1, function(row) distance_func(row,
    test_point))
    sorted_indices <- order(distances)
    nearest_neighbors <- train_data[sorted_indices[1:k], ]</pre>
```

## # Return the majority vote for classification

```
prediction <- names(sort(table(nearest_neighbors[, ncol(nearest_neighbors)]), decreasing =
TRUE))[1]</pre>
```

return(prediction)

}

## # Function to calculate evaluation metrics

```
evaluate_knn <- function(train_data, test_data, k, distance_func) {
    predictions <- sapply(1:nrow(test_data), function(i) {
        knn_general(train_data, test_data[i, -ncol(test_data)], k, distance_func)
    })</pre>
```

```
true_labels <- test_data$Species</pre>
```

## # Calculate confusion matrix

```
confusion_matrix <- table(Predicted = predictions, Actual = true_labels)
```

```
# Calculate accuracy
```

```
accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)</pre>
```

# # Calculate F1 score (macro-averaged)

```
f1_score <- mean(sapply(levels(true_labels), function(class) {
    precision <- confusion_matrix[class, class] / sum(confusion_matrix[class, ])
    recall <- confusion_matrix[class, class] / sum(confusion_matrix[, class])
    f1 <- ifelse(precision + recall > 0, 2 * precision * recall / (precision + recall), 0)
    return(f1)
```

```
}))
```

list(predictions = predictions, accuracy = accuracy, f1\_score = f1\_score, confusion\_matrix = confusion\_matrix)

## }

#### # Evaluate with Manhattan Distance

results\_manhattan <- evaluate\_knn(train\_data, test\_data, k = 5, distance\_func = manhattan\_distance) results\_manhattan\$accuracy results\_manhattan\$f1\_score results\_manhattan\$confusion\_matrix

#### # Evaluate with Minkowski Distance (p = 3)

results\_minkowski <- evaluate\_knn(train\_data, test\_data, k = 5, distance\_func = function(x1, x2) minkowski\_distance(x1, x2, p = 3)) results\_minkowski\$accuracy results\_minkowski\$f1\_score results\_minkowski\$confusion\_matrix

#### # Evaluate with Chebyshev Distance

results\_chebyshev <- evaluate\_knn(train\_data, test\_data, k = 5, distance\_func =
chebyshev\_distance)
results\_chebyshev\$accuracy
results\_chebyshev\$f1\_score
results\_chebyshev\$confusion\_matrix</pre>

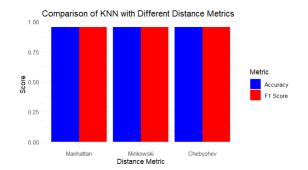
### # Prepare data for plotting

```
plot_data <- data.frame(
    Metric = rep(c("Accuracy", "F1 Score"), each = 3),
    Value = c(results_manhattan$accuracy, results_minkowski$accuracy,
    results_chebyshev$accuracy,
        results_manhattan$f1_score, results_minkowski$f1_score, results_chebyshev$f1_score),
    Distance = factor(rep(c("Manhattan", "Minkowski", "Chebyshev"), 2), levels = c("Manhattan",
    "Minkowski", "Chebyshev"))
    )
```

# # Plot the results

```
library(ggplot2)
```

```
ggplot(plot_data, aes(x = Distance, y = Value, fill = Metric)) +
geom_bar(stat = "identity", position = "dodge") +
theme_minimal() +
labs(title = "Comparison of KNN with Different Distance Metrics", y = "Score", x = "Distance
Metric") +
scale_fill_manual(values = c("Accuracy" = "blue", "F1 Score" = "red"))
```



As it turns out here, the metrics turn out largely the same for all three distance metrics, but there is a slight difference in one of the confusion matrices. The important thing to keep in mind is that you can't know which distance metric will produce the best results (or the same results) unless you test them.

Resources:

- 1. <u>https://www.kdnuggets.com/2020/11/most-popular-distance-metrics-knn.html</u>
- 2. <u>https://www.analyticsvidhya.com/blog/2021/08/how-knn-uses-distance-measures/</u>
- 3. <u>https://medium.com/@luigi.fiori.lf0303/distance-metrics-and-k-nearest-neighbor-knn-1b840969c0f4</u>
- 4. https://towardsdatascience.com/knn-k-nearest-neighbors-1-a4707b24bd1d
- 5. <u>https://www.linkedin.com/advice/3/what-most-effective-distance-metrics-optimizing-xndwc</u>