

# Week4: Metrics and Naive Bayes

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## 1. Why Evaluation Deserves Its Own Discussion

So far in this module, you have trained some models and understood their behavior.

In doing so, we often report **accuracy** — not because it is the best metric, but because it is simple and easy to interpret at first glance.

However, this raises an important question:

*How do we decide whether a model is actually “good”?*

Answering this requires more than training a model.

In addition to other factors, it requires **choosing the right way to evaluate it**.

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### 1.1 Models Do Not Speak for Themselves

A trained model does not tell us:

- whether it is reliable,
- whether it generalises,
- or whether it is useful in practice.

All of these judgements come from **evaluation metrics**.

Crucially:

**Different metrics answer different questions.**

Using the wrong metric can:

- make a weak model look strong,
  - hide important failure modes,
  - or encourage the wrong modelling decisions.
- 

## 2. Classification Evaluation — Confusion Matrix & Core Metrics

In a **classification task**, a model predicts a *class label* (e.g. **yes** / **no**, **positive** / **negative**).

To evaluate such predictions meaningfully, we must first understand **how predictions can be right or wrong**.

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## 2.1 The Confusion Matrix: The Foundation

At the heart of classification evaluation lies the **confusion matrix**.

It compares: - what the model *predicted*, - against what actually *happened*.

For a binary classification problem, there are four possible outcomes:

	Actual: Yes	Actual: No
Predicted: Yes	True Positive (TP)	False Positive (FP)
Predicted: No	False Negative (FN)	True Negative (TN)

Every classification metric is derived from these four quantities.

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## 2.2 Interpreting the Outcomes

Understanding these outcomes is more important than memorising formulas.

- **True Positive (TP)**  
The model predicted *yes*, and the outcome was *yes*.
- **False Positive (FP)**  
The model predicted *yes*, but the outcome was *no*.  
(A *false alarm*.)
- **False Negative (FN)**  
The model predicted *no*, but the outcome was *yes*.  
(A *missed case*.)
- **True Negative (TN)**  
The model predicted *no*, and the outcome was *no*.

Different applications care about these errors **very differently**.

## 2.3 Accuracy: The Simplest Metric

**Accuracy** measures how often the model is correct overall. Conceptually:

*Out of all predictions, how many were correct?*

Accuracy is intuitive — but also dangerous.

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### Why Accuracy Can Be Misleading

Accuracy treats all errors equally. This becomes problematic when:

- the dataset is **imbalanced**,
- one class dominates the data,
- some mistakes are more costly than others.

A model that predicts the majority class every time may achieve high accuracy while being practically useless.

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## 2.4 Precision, Recall, and F1 Score: A Loan Approval Example

Consider a loan default prediction model used by a bank.

- **Positive** (“**yes**”) = the applicant will **default**
- **Negative** (“**no**”) = the applicant will **repay**

This framing lets us interpret each metric in business terms.

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## Precision: Are Default Warnings Trustworthy?

**Precision** answers:

*When the model predicts that an applicant will default, how often is it correct?*

Suppose the model flags **100 applicants as likely to default**:

- 80 actually default
- 20 would have repaid

**Precision = 80%**

High precision means:

- few **false positives**,
- fewer safe customers wrongly rejected,
- risk flags can be trusted.

Low precision means the bank loses good customers by rejecting loans unnecessarily.

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## Recall: How Many Risky Loans Did We Catch?

**Recall** answers:

*Of all applicants who will actually default, how many did the model identify?*

Suppose **200 applicants would eventually default**:

- the model flags 80 of them,
- 120 risky applicants are missed.

**Recall = 40%**

High recall means:

- few **false negatives**,
- fewer bad loans slip through,
- lower financial losses.

Low recall means the bank approves many loans that later default.

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## The Precision–Recall Trade-off

Precision and recall often conflict. Which is preferable depends on context:

- Risk-averse institutions prioritize **recall**
- Growth-focused institutions prioritize **precision**

This choice reflects business values, not algorithmic correctness.

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## F1 Score: A Balanced Summary

The **F1 score** combines precision and recall into a single metric. It is useful when:

- defaults are rare (class imbalance),
- both rejecting good customers and approving bad ones are costly,
- accuracy alone is misleading.

However, using F1 implicitly assumes that **false positives and false negatives matter equally**—a judgement that may or may not align with real-world priorities.

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## 3. Naive Bayes Demo — Classification Evaluation in Practice

We now apply the ideas from the previous section to a **concrete example** using Naive Bayes. Our goal here is **not** to build a high-performing model. It is to see how **evaluation metrics behave** in practice.

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### 3.1 Dataset and Feature Choice

For this demonstration, we use the same **UCI Bank Marketing dataset**.

To keep the demo interpretable, we restrict ourselves to a **small set of categorical features**:

- job
- education
- housing
- loan

- contact

This is a **modelling choice**, not a requirement.

### 3.2 Loading Data

We reuse the same train/validation split introduced earlier.

#### R: Loading Data and Creating Splits

```
library(tidyverse)
library(dplyr)

bank <- read.csv("data/raw/bank-additional.csv", sep = ";", stringsAsFactors = FALSE)

model_data <- bank %>%
  select(-duration) %>%
  mutate(y = as.factor(y))

set.seed(42)

n <- nrow(model_data)
idx <- sample(n)

train_idx <- idx[1:floor(0.6 * n)]
val_idx   <- idx[(floor(0.6 * n) + 1):floor(0.8 * n)]
test_idx  <- idx[(floor(0.8 * n) + 1):n]

train_data <- model_data[train_idx, ]
val_data   <- model_data[val_idx, ]
test_data  <- model_data[test_idx, ]
```

#### Python: Loading Data and Creating Splits

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.compose import ColumnTransformer
from sklearn.preprocessing import OneHotEncoder
from sklearn.pipeline import Pipeline

bank = pd.read_csv("data/raw/bank-additional.csv", sep=";")
```

```

model_data = bank.drop(columns=["duration"])
model_data["y"] = model_data["y"].astype("category")

# Split features / target
X = model_data.drop(columns=["y"])
y = model_data["y"]

# Identify column types
cat_cols = X.select_dtypes(include=["object", "category"]).columns
num_cols = X.select_dtypes(exclude=["object", "category"]).columns

# Preprocessing: encode categoricals, pass through numerics
preprocess = ColumnTransformer(
    transformers=[
        ("num", "passthrough", num_cols),
        ("cat", OneHotEncoder(handle_unknown="ignore"), cat_cols),
    ]
)

X_train, X_temp, y_train, y_temp = train_test_split(
    X, y, test_size=0.4, random_state=42, stratify=y
)

X_val, X_test, y_val, y_test = train_test_split(
    X_temp, y_temp, test_size=0.5, random_state=42, stratify=y_temp
)

```

---

### 3.3 Preparing the Data

#### R: Select Features and Split Data

```

library(e1071)

features <- c("job", "education", "housing", "loan", "contact")

nb_train <- train_data[, c(features, "y")]
nb_val    <- val_data[, c(features, "y")]

```



**i** What this code is doing

Selects a small subset of categorical predictors.  
Uses the existing training and validation splits.  
Keeps the target variable unchanged.  
Why this matters:  
Fewer features make probability estimates easier to interpret.  
This avoids accidental leakage or overfitting.

## Python: Select Features and Split Data

```
from sklearn.naive_bayes import CategoricalNB
from sklearn.preprocessing import OrdinalEncoder

features = ["job", "education", "housing", "loan", "contact"]

X_train_nb = X_train[features]
X_val_nb    = X_val[features]
```

**i** What this code is doing

Restricts the feature set to categorical variables.  
Prepares separate training and validation inputs.

## 3.4 Training a Naive Bayes Classifier

### R: Training the Model

```
nb_model <- naiveBayes(
  y ~ .,
  data = nb_train
)
```

**i** What this code is doing

Fits a Naive Bayes classifier.  
Estimates class priors and conditional probabilities.  
Assumes conditional independence between features.

## Python: Training the Model

```
nb_model = Pipeline(  
    steps=[  
        ("enc", OrdinalEncoder()),  
        ("nb", CategoricalNB())  
    ]  
)  
  
nb_model.fit(X_train_nb, y_train)
```

```
Pipeline(steps=[('enc', OrdinalEncoder()), ('nb', CategoricalNB())])
```

**i** What this code is doing

Encodes categorical variables numerically because scikit-learn requires numeric inputs.  
Trains a categorical Naive Bayes classifier.  
Keeps preprocessing and modelling in one pipeline.

## 3.5 From Probabilities to Predictions

Naive Bayes produces probabilities, not hard decisions. To obtain class predictions, we apply a threshold.

### R: Predicted Probabilities and Classes

```
nb_probs <- predict(nb_model, nb_val, type = "raw")  
nb_preds <- predict(nb_model, nb_val, type = "class")
```

## Python: Predicted Probabilities and Classes

```
nb_probs = nb_model.predict_proba(X_val_nb)  
nb_preds = nb_model.predict(X_val_nb)
```

**i** Why this step matters

Probabilities express uncertainty.  
Class labels depend on a threshold.

Changing the threshold changes precision and recall.  
This directly links model output to evaluation metrics.

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## 4. Computing Classification Metrics (R & Python)

We now evaluate the Naive Bayes model using the metrics introduced earlier. The key idea in this section is simple:

**The same predictions can look “good” or “bad” depending on the metric we use.**

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### 4.1 Confusion Matrix

We begin with the confusion matrix, which shows **how predictions break down**.

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#### R: Confusion Matrix

```
table(  
  Predicted = nb_preds,  
  Actual    = nb_val$y  
)
```

	Actual	
Predicted	no	yes
no	736	88
yes	0	0

#### **i** What this shows

- How many predictions fall into each TP, FP, FN, TN category.
- The raw material from which all classification metrics are computed.

Why this matters:

- Metrics summarise behaviour.
- The confusion matrix *reveals* behaviour.

---

## Python: Confusion Matrix

```
from sklearn.metrics import confusion_matrix  
  
confusion_matrix(y_val, nb_preds)
```

```
array([[733,  1],  
       [ 90,  0]])
```

### **i** Why start with the confusion matrix

Looking only at a single metric hides important failure modes. The confusion matrix forces us to ask:

- *What kinds of mistakes is the model making?*

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## 4.2 Accuracy

**Accuracy** measures how often the model is correct overall.

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### R: Accuracy

```
mean(nb_preds == nb_val$y)
```

```
[1] 0.8932039
```

## Python: Accuracy

```
from sklearn.metrics import accuracy_score  
  
accuracy_score(y_val, nb_preds)
```

0.8895631067961165

**i** How to interpret accuracy here

- Accuracy treats all mistakes equally.
- With class imbalance, this can be misleading.

Use accuracy as a **baseline**, not a final judgement.

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## 4.3 Precision and Recall

Accuracy alone does not tell us **what kind of mistakes** the model is making.

Precision and recall help us look deeper.

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## R: Precision and Recall

```
#install.packages("recipes")  
library(caret)  
  
conf <- confusionMatrix(nb_preds, nb_val$y, positive = "yes")  
  
conf$byClass["Precision"]
```

Precision  
NA

```
conf$byClass["Recall"]
```

Recall  
0

---

## Python: Precision and Recall

```
from sklearn.metrics import precision_score, recall_score  
  
precision_score(y_val, nb_preds, pos_label="yes")
```

0.0

```
recall_score(y_val, nb_preds, pos_label="yes")
```

0.0

### **i** Interpreting precision vs recall

- Precision focuses on **false positives**.
- Recall focuses on **false negatives**.

Which one matters more depends on the **context**, not the model.

---

## 4.4 F1 Score

The **F1 score** combines precision and recall into a single number.

It is useful when:

- classes are imbalanced,
  - both types of errors matter.
-

## R: F1 Score

```
conf$byClass["F1"]
```

F1

NA

---

## Python: F1 Score

```
from sklearn.metrics import f1_score  
  
f1_score(y_val, nb_preds, pos_label="yes")
```

0.0

---

## 4.5 A Note on Thresholds (Conceptual Only)

Remember that Naive Bayes produces **probabilities**. Changing the classification threshold would:

- change the confusion matrix,
- alter precision and recall,
- leave the underlying probabilities unchanged.

We do **not** tune thresholds here — but it is important to know that metrics depend on this choice.

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## 5. Regression Metrics — What Changes When the Target Is Numeric

So far, we have focused on **classification**, where predictions are discrete labels (e.g. **yes** / **no**). In **regression**, the situation is different:

There is no notion of “correct” or “incorrect” — only *how far off* a prediction is.

This requires a different way of evaluating models.

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### 5.1 Why Accuracy Does Not Apply

In regression tasks: - predictions are numeric, - errors are continuous, - mistakes vary in magnitude.

Asking whether a prediction is “right” no longer makes sense.

Instead, we ask:

*How large are the prediction errors?*

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### 5.2 Absolute Error vs Squared Error

The core idea in regression evaluation is **measuring error size**.

We briefly introduce three commonly used metrics.

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#### Mean Absolute Error (MAE)

**MAE** measures the average absolute difference between predictions and actual values.

Conceptually: - treats all errors equally, - easy to interpret in original units, - robust to occasional large errors.

MAE answers: > *On average, how far off are we?*

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## Mean Squared Error (MSE)

**MSE** measures the average of squared errors.

Conceptually: - penalises large errors more heavily, - sensitive to outliers, - useful when large mistakes are particularly costly.

MSE answers: > *How much do large errors matter to us?*

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## Root Mean Squared Error (RMSE)

**RMSE** is the square root of MSE.

Why it is used: - brings error back to the original scale, - still penalises large errors more than MAE.

RMSE balances: - interpretability, - sensitivity to large deviations.

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## 5.3 Choosing Between MAE and RMSE

There is no universally correct choice.

A useful rule of thumb:

- Use **MAE** when:
  - interpretability matters,
  - all errors are equally costly.
- Use **RMSE** when:
  - large errors are particularly undesirable,
  - outliers should be penalised more strongly.

Metric choice reflects **modelling priorities**, not mathematical correctness.

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## 5.4 Connecting Back to Your Project Dataset

In the **Online Retail dataset**, regression-style targets might include: - invoice value, - total customer spend, - basket size.

If you choose such a target: - MAE tells you average monetary error, - RMSE highlights whether large mistakes dominate.

As with classification, the metric must match the **question you are asking**.

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