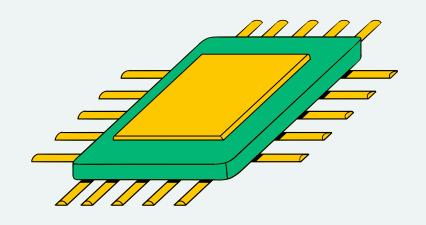
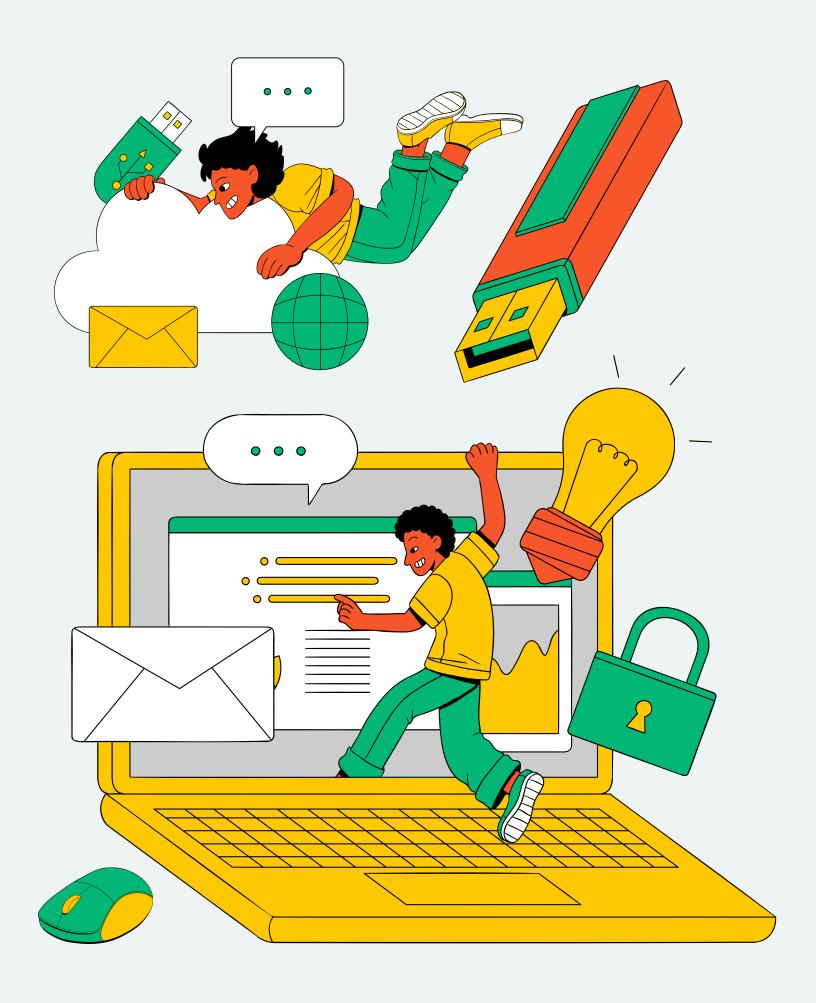


COMP 466 BUSINESS INTELLIGENCE ACTIVITY 1 PRESENTATION

PRESENTED BY:

TARIK BUĞRA AY





PRESENTATION OUTLINE

- Introduction
- Project Objective
- Dataset Overview
- Data Preparation
- Feature Engineering
- Clustering
- Clustering vs Actual Class
- Classification Results
- Model Comparison
- Conclusion



INTRODUCTION

This project explores how machine learning can be applied to analyze breast cancer data using both clustering and classification techniques.

The goal is to:

- Discover natural groupings in the data
- Build predictive models to classify outcomes
- Evaluate and compare performance using common ML metrics





OBJECTIVE OF THE PROJECT

The aim of this project is to apply machine learning techniques to a breast cancer dataset to:

- Identify natural groupings using clustering algorithms
- Use those clusters as pseudo-labels for classification
- Compare results with real class labels
- Evaluate models using:
 - Accuracy
 - ROC AUC Score
 - RMSE





DATASET OVERVIEW

- Dataset Name: breast_cancer.arff
- Source: University Medical Centre, Institute of Oncology, Ljubljana
- Instances: 286 patients
- Features: 9 input attributes + 1 class label
- Class Labels:
 - no-recurrence-events
 - recurrence-events

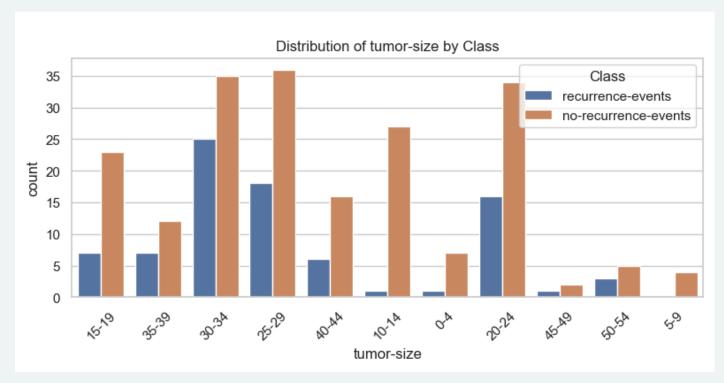


Figure 1. Distribution of tumor size by class.

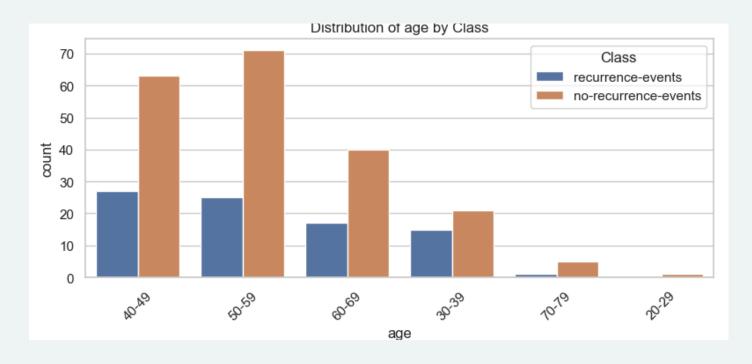


Figure 2. Distribution of age by class.



DATA PREPARATION

Steps Taken to Prepare the Data

- Parsed `.arff` file manually to extract data and attributes
- Handled missing values:
 - `node-caps`: filled with mode
 - o breast-quad: filled with mode
- Removed extra quotes and cleaned column names
- Saved cleaned version as CSV for further processing
- Resulting dataset:
 - Shape: 286 rows × 10 columns

```
opause,tumor-size,inv-nodes,node-caps,deg-malig,breast,breast-quad,irradiat,
40-49, premeno, 15-19, 0-2, yes, 3, right, left_up, no, recurrence-events
50-59,ge40,15-19,0-2,no,1,right,central,no,no-recurrence-events
50-59,ge40,35-39,0-2,no,2,left,left_low,no,recurrence-events
40-49, premeno, 35-39, 0-2, yes, 3, right, left_low, yes, no-recurrence-events
40-49, premeno, 30-34, 3-5, yes, 2, left, right_up, no, recurrence-events
50-59, premeno, 25-29, 3-5, no, 2, right, left_up, yes, no-recurrence-events
50-59,ge40,40-44,0-2,no,3,left,left_up,no,no-recurrence-events
40-49, premeno, 10-14, 0-2, no, 2, left, left_up, no, no-recurrence-events
40-49, premeno, 0-4, 0-2, no, 2, right, right_low, no, no-recurrence-events
40-49,ge40,40-44,15-17,yes,2,right,left_up,yes,no-recurrence-events
50-59, premeno, 25-29, 0-2, no, 2, left, left_low, no, no-recurrence-events
60-69,ge40,15-19,0-2,no,2,right,left_up,no,no-recurrence-events
50-59, ge40, 30-34, 0-2, no, 1, right, central, no, no-recurrence-events
50-59,ge40,25-29,0-2,no,2,right,left_up,no,no-recurrence-events
40-49, premeno, 25-29, 0-2, no, 2, left, left_low, yes, recurrence-events
30-39, premeno, 20-24, 0-2, no, 3, left, central, no, no-recurrence-events
50-59, premeno, 10-14, 3-5, no, 1, right, left_up, no, no-recurrence-events
60-69,ge40,15-19,0-2,no,2,right,left_up,no,no-recurrence-events
50-59, premeno, 40-44, 0-2, no, 2, left, left_up, no, no-recurrence-events
50-59,ge40,20-24,0-2,no,3,left,left_up,no,no-recurrence-events
50-59,1t40,20-24,0-2,no,1,left,left_low,no,recurrence-events
60-69,ge40,40-44,3-5,no,2,right,left_up,yes,no-recurrence-events
50-59,ge40,15-19,0-2,no,2,right,left_low,no,no-recurrence-events
40-49, premeno, 10-14, 0-2, no, 1, right, left_up, no, no-recurrence-events
30-39, premeno, 15-19, 6-8, yes, 3, left, left_low, yes, recurrence-events
```

Figure 3. A snippet from data.



FEATURE ENGINEERING

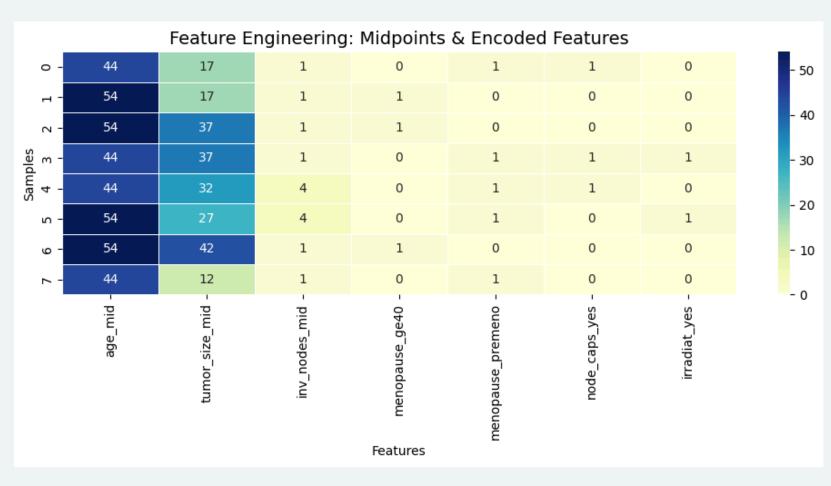


Figure 4. Feature Engineered Columns

Feature Engineering Steps

To prepare the dataset for clustering and classification, we:

- bConverted categorical ranges to numeric midpoints:
 - o `age`, `tumor-size`, `inv-nodes`
- Applied one-hot encoding to:
 - menopause`, node-caps`, irradiat`
 - breast`,`breast-quad`
- Retained `deg-malig` as numeric

Final dataset:

→ 286 rows × 27 columns



CLUSTERING

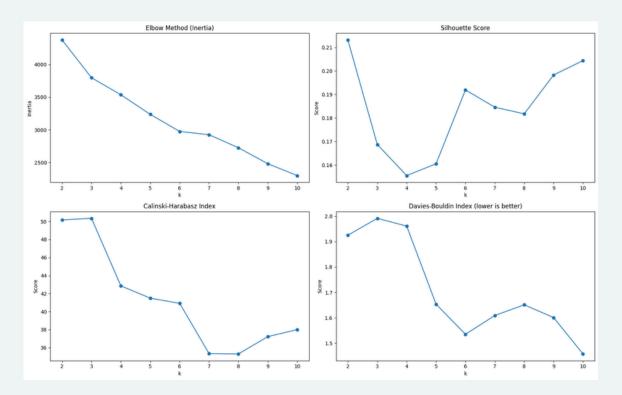


Figure 5. Evaluation metric chart (4 subplots) — Elbow, Silhouette, CH, DBI

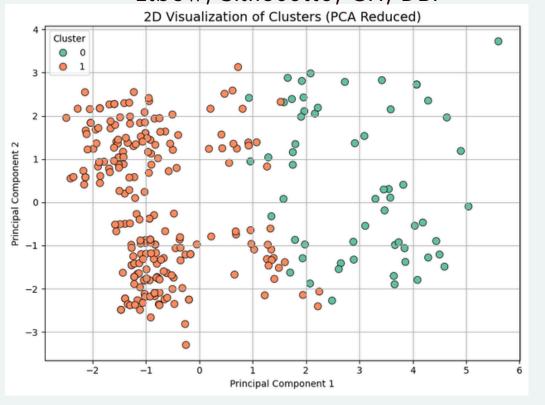


Figure 6. PCA cluster visualization

Clustering with KMeans

- Standardized numeric features before clustering
- Applied KMeans to discover natural groupings in the data
- Evaluated multiple values of `k` using:
 - Elbow Method (Inertia)
 - Silhouette Score
 - Calinski-Harabasz Index
 - Davies-Bouldin Index

Chosen Number of Clusters: k = 2

- Best balance across metrics
- Matches the known recurrence/no-recurrence classes
- Simple for downstream analysis

CLUSTERING VS ACTUAL CLASS

Evaluating Clustering Against Actual Class

- After clustering, we compared the generated cluster labels with the real medical outcomes (`Class` column).
- Mapped clusters to match the closest class distribution
- Adjusted cluster labels where needed
- Evaluation metrics:
- Accuracy: 0.71
- ROC AUC Score: 0.69
- RMSE: 0.46
- This showed partial alignment between clusters and real-world outcomes.

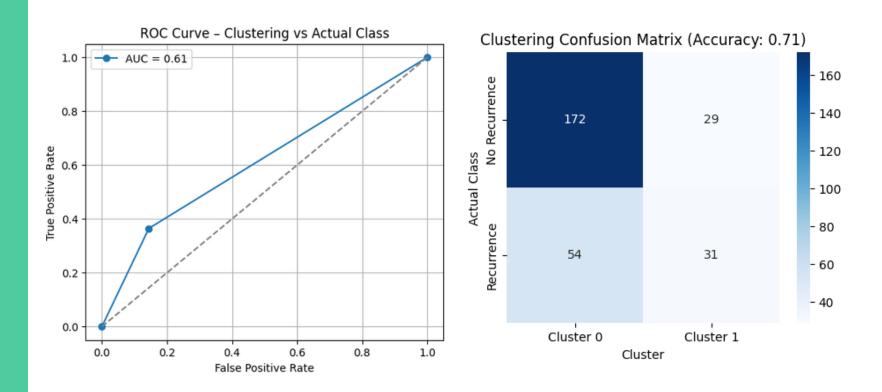


Figure 7. ROC Curve and Confusion Matrix



CLASSIFICATION RESULTS

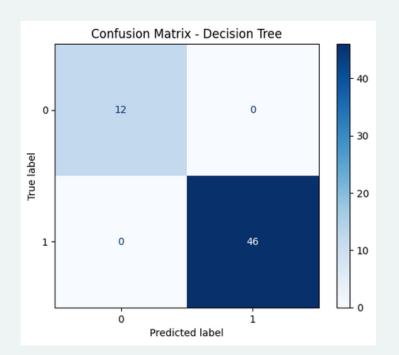
Decision Tree Classifier Results

We trained two classification models using the same features:

- 1. Model 1 Predicting Cluster Labels
 - Accuracy: 1.00
 - ROC AUC: 1.00
 - RMSE: 0.00
- 2. Model 2 Predicting Actual Class Labels
 - Accuracy: 0.71
 - ROC AUC: 0.64
 - RMSE: 0.53

Model 1 replicated clustering perfectly.

Model 2 showed more realistic performance.



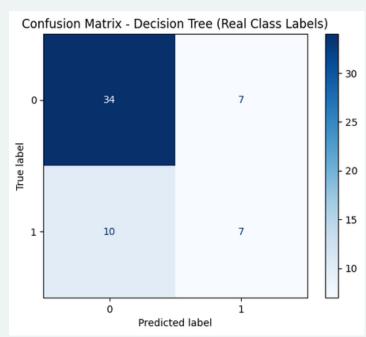
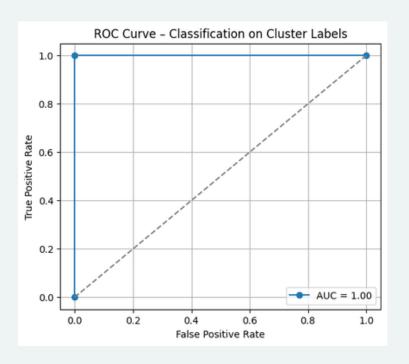


Figure 8. Confusion Matrixes for Classifications.



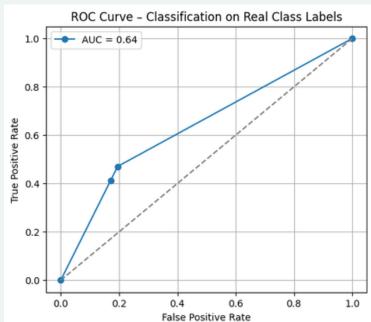


Figure 9. ROC Curves for Classifications.



CONCLUSION



- KMeans clustering revealed patterns in the breast cancer dataset that partially aligned with real recurrence labels.
- Classification models performed very differently:
 - Cluster labels were easy to learn and replicate.
 - Real class labels were harder to predict but gave more realistic results.
- Feature engineering played a key role in shaping both clustering and classification performance.
- Project Repository GitHub:
 - https://github.com/TarikBugraAy/breast-cancerclustering-classification?tab=readme-ov-file

