

BREAST_CANCER _CLASSIFIER

Classification Project with Multiple ML Models & Streamlit
Deployment

PROJECT OVERVIEW:

Predict whether a tumor is
Malignant (M) ● or Benign (B) ●
using the Breast Cancer Wisconsin
(Diagnostic) dataset.

AGENDA:

1. Dataset
2. Data Preprocessing
3. Exploratory Data Analysis (EDA)
4. Feature Selection & Explanation
5. Model Training & Evaluation
6. Streamlit App Deployment
7. Conclusion

1. DATASET :

- 30 numeric features such as: radius, texture, smoothness, compactness, symmetry, fractal dimension
- Target: diagnosis (M = Malignant, B = Benign)
- Rows: 569

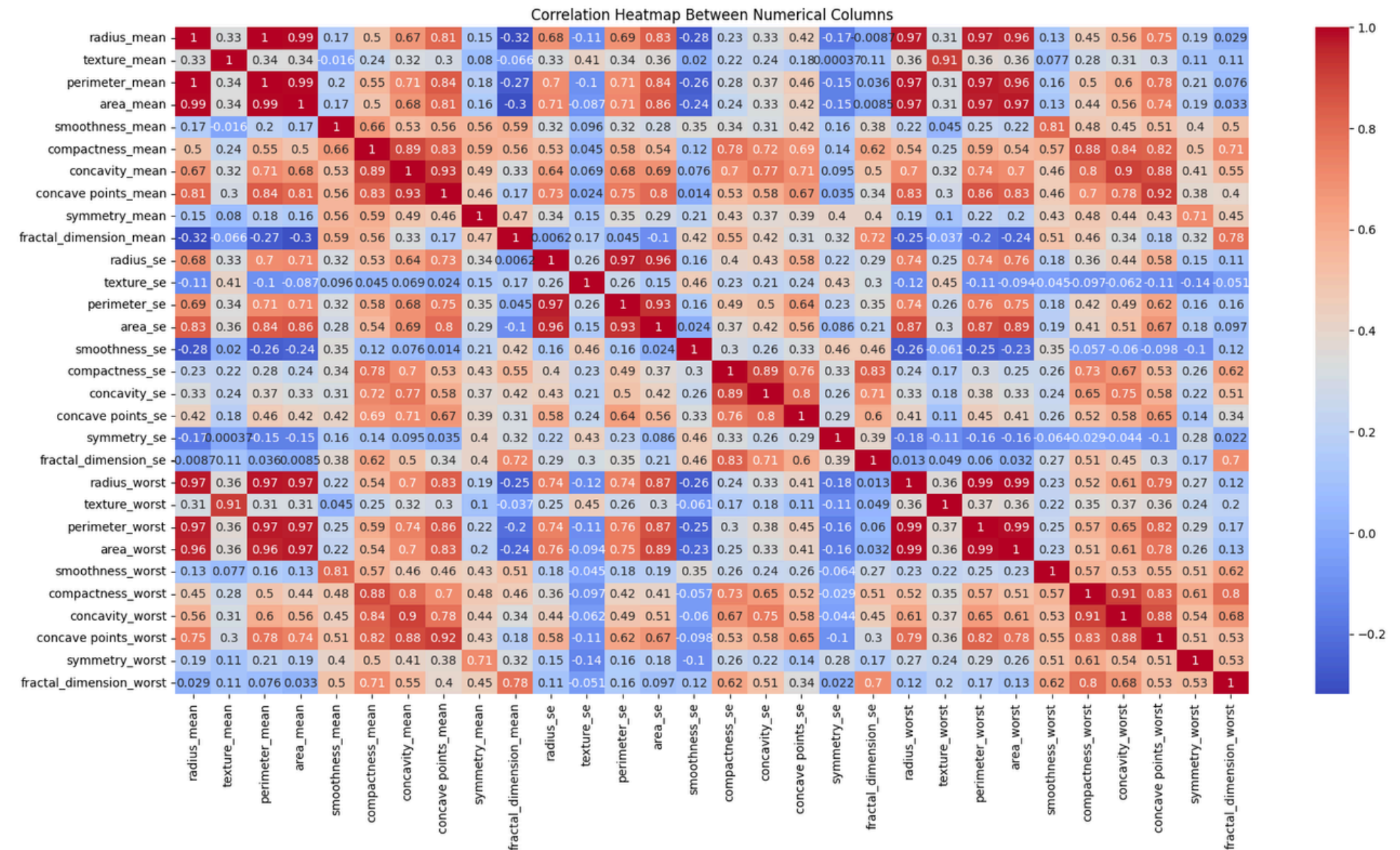
2. DATA

PREPROCESSING:

- Removed unnecessary columns (id, Unnamed:32)
- Checked missing values & duplicates
- Handled outliers with Boxplots (replaced extreme values with boundary values)
- Normalized features using Min-Max Scaling
- Encoded target variable (Label Encoding)

3. EXPLORATORY DATA ANALYSIS (EDA):

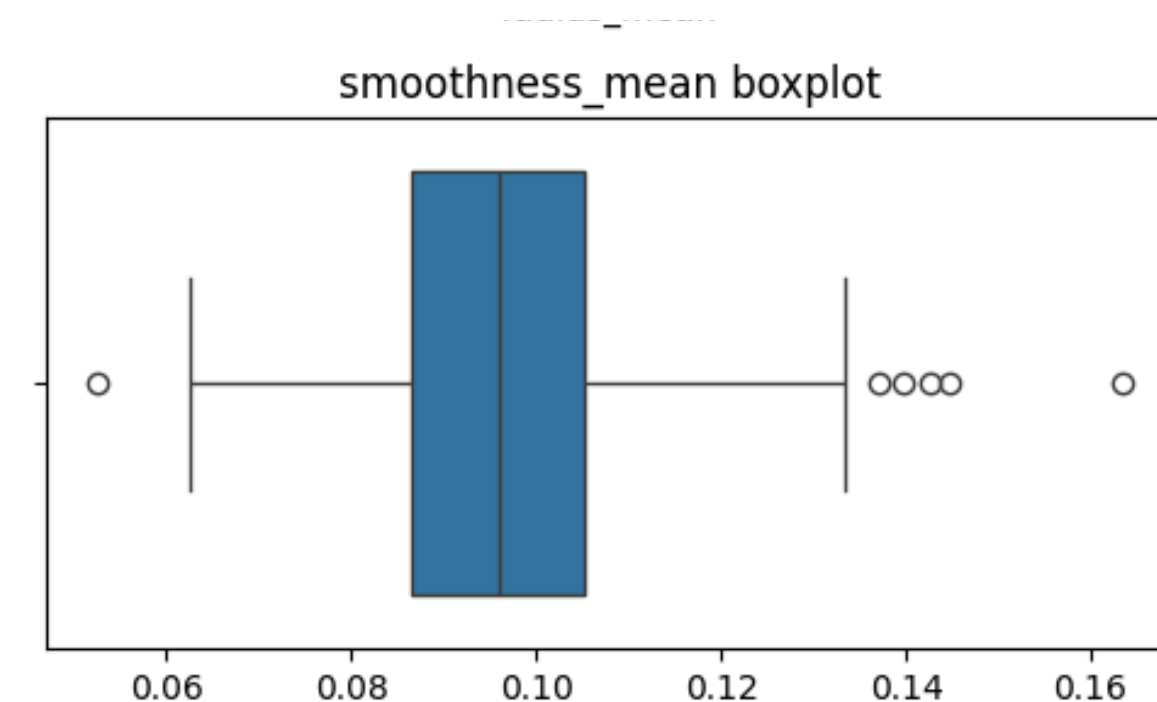
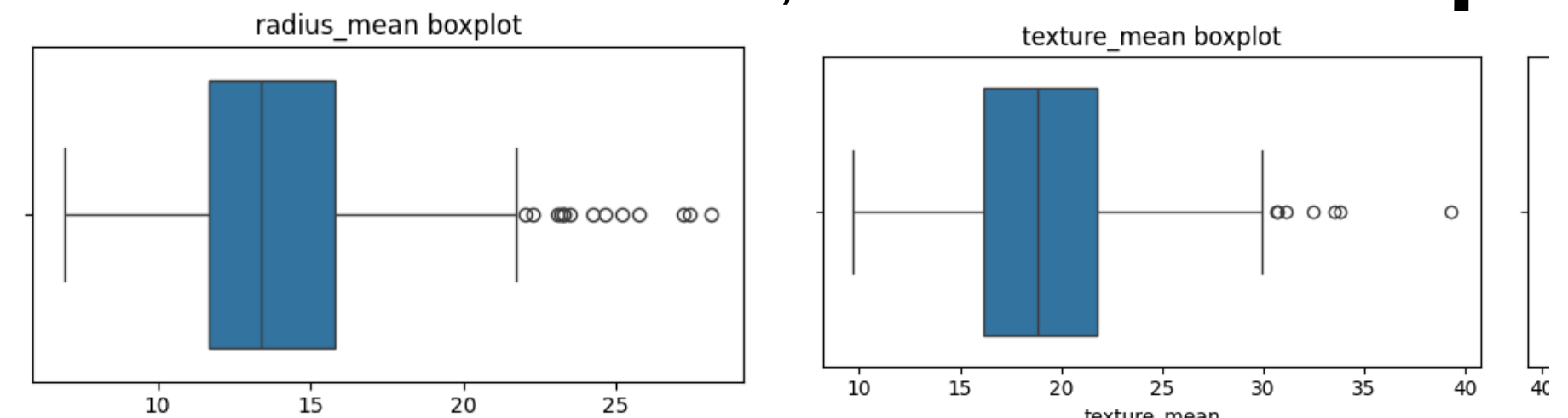
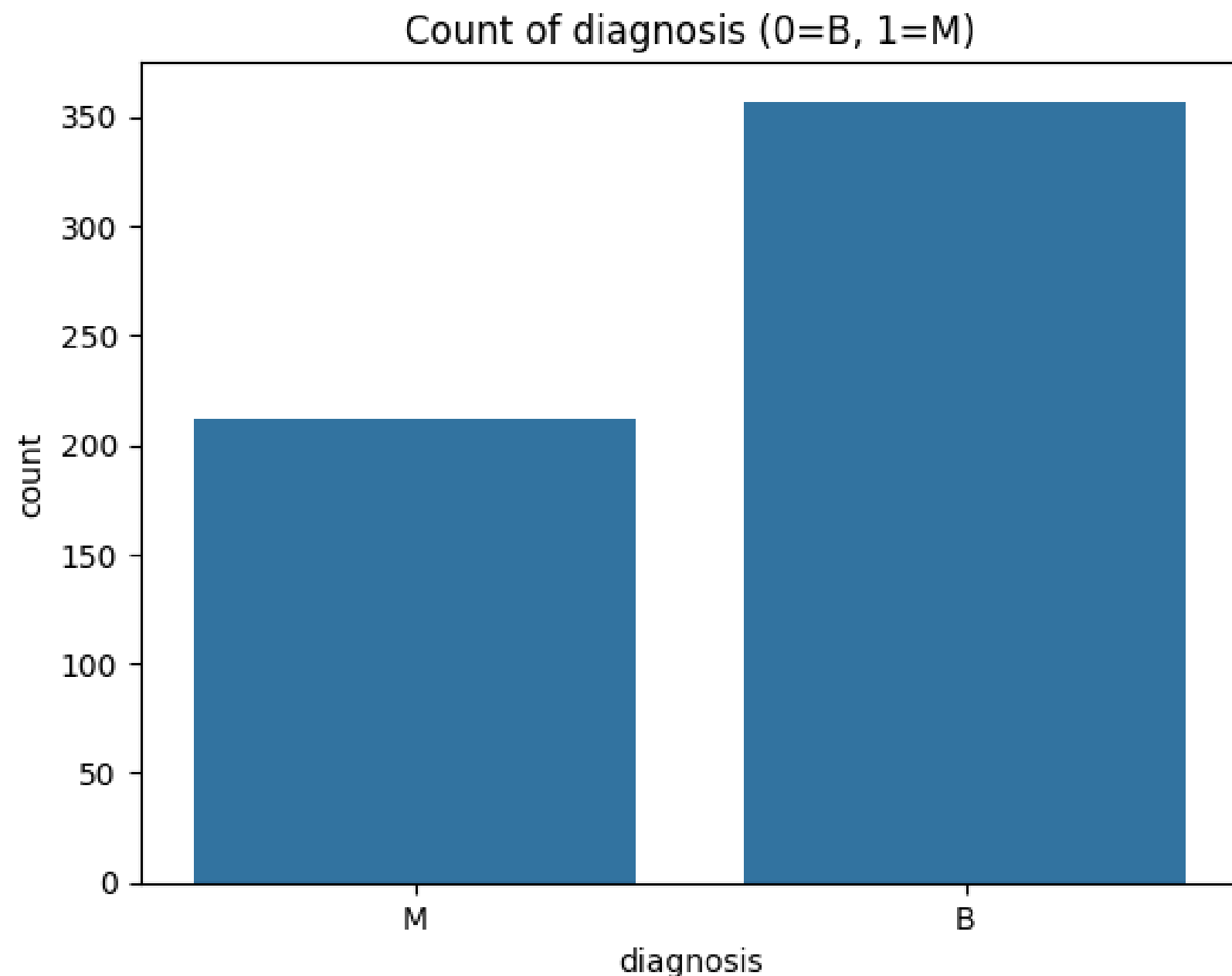
- Correlation Heatmap



3. EXPLORATORY DATA ANALYSIS (EDA):

- Count plot for target distribution: B = 357 ●, M = 212 ●

- Boxplots of key features (radius_mean, texture_mean, smoothness_mean)






4. FEATURE SELECTION & EXPLANATION:

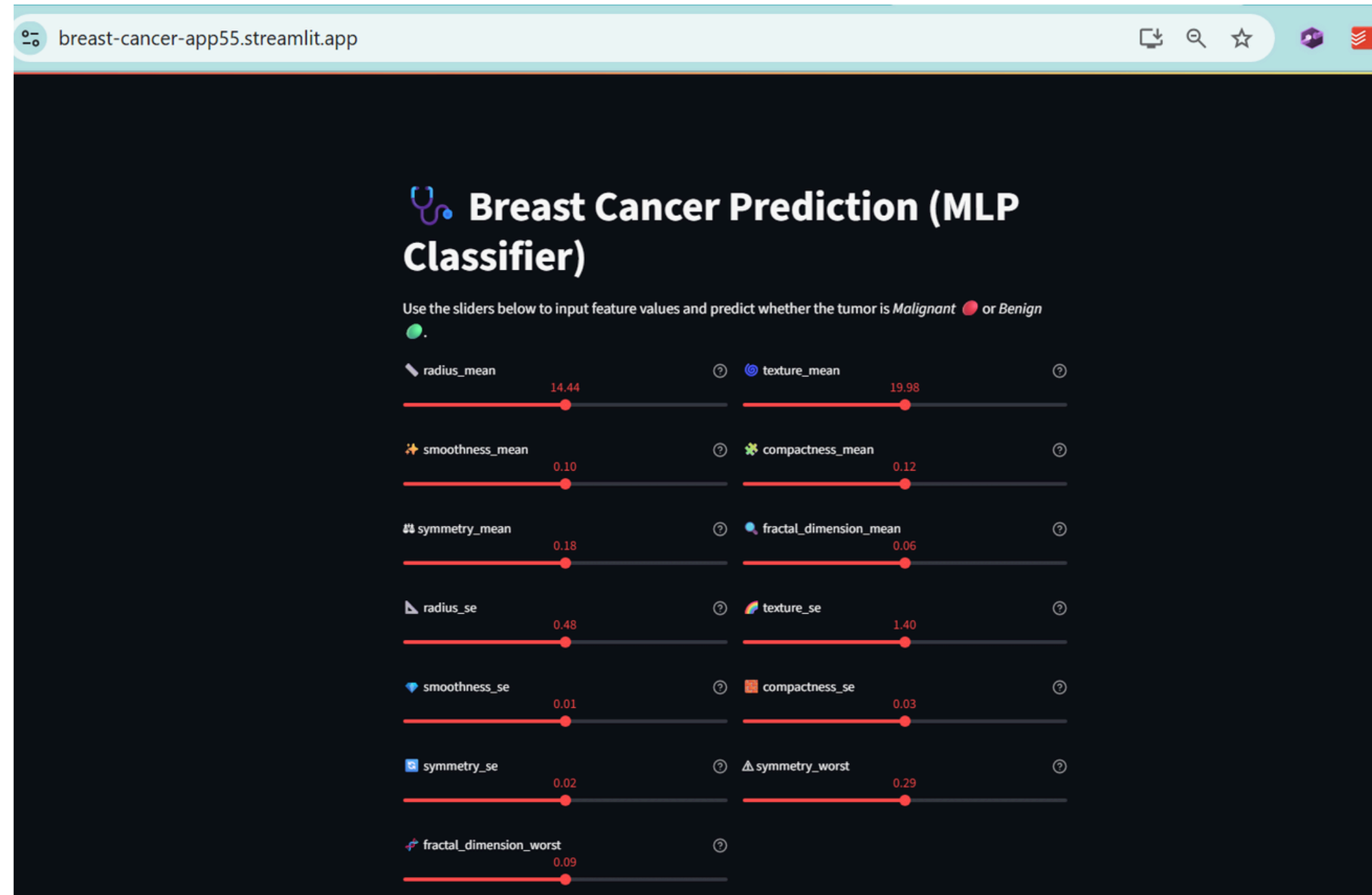
- Removed highly correlated features (≥ 0.8)
- Selected 13 final features for MLP model:
 - 1.radius_mean :Average radius of the cells
 - 2.texture_mean: Variation in texture
 - 3.smoothness_mean: How smooth the cell edges are
 - 4.compactness_mean: Ratio of area to perimeter²
 - 5.symmetry_mean: Symmetry of the cells
 - 6.fractal_dimension_mean: Complexity of cell boundary
 - 7.radius_se: Standard error of radius
 - 8.texture_se: Standard error of texture
 - 9.smoothness_se: Standard error of smoothness
 - 10.compactness_se: Standard error of compactness
 - 11.symmetry_se: Standard error of symmetry
 - 12.symmetry_worst: Worst (largest) symmetry
 - 13.fractal_dimension_worst: Worst fractal dimension

5. MODEL TRAINING & EVALUATION:

- Train,Test Split: 80% training , 20% testing
 - Trained multiple classifiers: Random Forest, Decision Tree, XGBoost, Gradient Boosting, Perceptron, MLP
 - Metrics: Accuracy, Precision, Recall, F1 Score
 - Model Test Accuracy F1 Score:
 - 1.Random Forest 93.86% 0.916
 - 2.Decision Tree 92.11% 0.897
 - 3.XGBoost 95.61% 0.941
 - 4.Gradient Boosting 94.74% 0.930
 - 5.Perceptron 94.74% 0.929
 - 6.MLP 96.49% 0.953
 - MLP is best due to:
High Test Accuracy
Balanced Precision & Recall

6. STREAMLIT APP DEPLOYMENT:

- Interactive sliders for input
- Shows predicted tumor type   and probabilities 
- Features have descriptive tooltips
- Streamlit link : <https://breast-cancer-app55.streamlit.app/>



7. CONCLUSION:

- Built robust breast cancer classifier with MLP
- Demonstrated: data preprocessing, feature selection, model comparison
- Deployed with Streamlit for easy interaction
- MLP chosen for best performance: high accuracy & F1 Score



THANK YOU