

Breast Cancer Diagnosis Analysis

Predict breast cancer diagnosis (malignant vs benign) from digitized cell nuclei images using machine learning

Jotty SwarmMLComprehensive

February 05, 2026

Contents

| | | |
|-------|--|----|
| 0.1 | Executive Summary | 4 |
| 0.1.1 | Key Results | 4 |
| 0.2 | Data Quality Analysis | 4 |
| 0.2.1 | Dataset Overview | 4 |
| 0.2.2 | Distribution Analysis | 4 |
| 0.2.3 | Feature Distributions | 5 |
| 0.2.4 | Outlier Analysis | 5 |
| 0.2.5 | Outlier Distribution | 6 |
| 0.3 | Correlation & Multicollinearity Analysis | 7 |
| 0.3.1 | Correlation Matrix | 7 |
| 0.3.2 | Highly Correlated Feature Pairs ($ r \geq 0.7$) | 7 |
| 0.3.3 | Variance Inflation Factor (VIF) | 8 |
| 0.3.4 | VIF Visualization | 8 |
| 0.4 | Data Profile | 9 |
| 0.4.1 | Dataset Overview | 9 |
| 0.4.2 | Data Types | 9 |
| 0.4.3 | EDA Recommendations | 9 |
| 0.5 | Feature Importance Analysis | 9 |
| 0.5.1 | Top 20 Features | 10 |
| 0.5.2 | Feature Importance Visualization | 10 |
| 0.6 | Model Benchmarking | 10 |
| 0.6.1 | Model Comparison | 10 |
| 0.6.2 | Performance Visualization | 12 |
| 0.7 | Learning Curve Analysis | 12 |
| 0.7.1 | Bias-Variance Diagnosis | 12 |
| 0.7.2 | Learning Curve Visualization | 12 |
| 0.7.3 | Interpretation Guide | 13 |
| 0.8 | Cross-Validation Detailed Analysis | 13 |
| 0.8.1 | Fold-by-Fold Results | 13 |
| 0.8.2 | Stability Analysis | 13 |
| 0.8.3 | CV Performance Distribution | 13 |
| 0.9 | Classification Performance | 13 |

| | | |
|--------|---|----|
| 0.9.1 | Classification Report | 13 |
| 0.9.2 | Confusion Matrix | 14 |
| 0.10 | ROC Curve Analysis | 14 |
| 0.10.1 | Key Metrics | 14 |
| 0.10.2 | ROC Curve | 14 |
| 0.11 | Precision-Recall Analysis | 14 |
| 0.11.1 | Key Metrics | 14 |
| 0.11.2 | Precision-Recall Curve | 14 |
| 0.12 | Probability Calibration Analysis | 14 |
| 0.12.1 | Calibration Metrics | 14 |
| 0.12.2 | Calibration Curve | 15 |
| 0.12.3 | Interpretation | 15 |
| 0.13 | Lift & Gain Analysis | 15 |
| 0.13.1 | Key Metrics | 15 |
| 0.13.2 | Cumulative Gains & Lift Curves | 15 |
| 0.13.3 | Business Interpretation | 15 |
| 0.14 | Threshold Optimization | 15 |
| 0.14.1 | Optimal Thresholds | 15 |
| 0.14.2 | Threshold Impact Analysis | 15 |
| 0.14.3 | Threshold Visualization | 19 |
| 0.14.4 | Cost Parameters Used | 19 |
| 0.15 | Error Analysis | 19 |
| 0.15.1 | Misclassification Summary | 19 |
| 0.15.2 | Confusion Matrix Breakdown | 20 |
| 0.15.3 | Hardest to Classify Samples (Most Confident Errors) | 20 |
| 0.15.4 | Error Distribution Analysis | 20 |
| 0.16 | SHAP Deep Analysis | 20 |
| 0.16.1 | Global Feature Importance (Mean SHAP) | 21 |
| 0.16.2 | SHAP Summary Plot | 21 |
| 0.16.3 | SHAP Feature Importance Bar | 21 |
| 0.16.4 | SHAP Dependence Plots (Top 3 Features) | 21 |
| 0.16.5 | SHAP Waterfall (Sample Prediction) | 21 |
| 0.17 | Baseline Comparison | 21 |
| 0.17.1 | Performance Improvement | 21 |
| 0.18 | Recommendations & Next Steps | 25 |
| 0.19 | Reproducibility | 25 |
| 0.19.1 | Model Configuration | 25 |
| 0.19.2 | Random Seeds | 25 |
| 0.19.3 | Environment | 25 |
| 0.19.4 | Package Versions | 26 |
| 0.19.5 | Generation Timestamp | 26 |

Contents

0.1 Executive Summary

Predict breast cancer diagnosis (malignant vs benign) from digitized cell nuclei images using machine learning.

0.1.1 Key Results

Best Model: Logistic Regression

Performance Metrics:

| Metric | Value |
|-----------|--------|
| Accuracy | 0.9825 |
| Precision | 0.9861 |
| Recall | 0.9861 |
| F1 | 0.9861 |
| Auc Roc | 0.9954 |

Dataset: 30 features analyzed

0.2 Data Quality Analysis

A comprehensive analysis of data quality, identifying potential issues before modeling.

0.2.1 Dataset Overview

| Metric | Value |
|-----------------------|-----------|
| Total Samples | 114 |
| Total Features | 30 |
| Numeric Features | 30 |
| Categorical Features | 0 |
| Features with Missing | 0 |
| Total Missing Values | 0 (0.00%) |

0.2.2 Distribution Analysis

| Feature | Skewness | Kurtosis | Assessment |
|----------------|----------|----------|----------------------------|
| mean radius | 1.03 | 1.35 | Right-skewed |
| mean texture | 0.25 | -0.40 | Symmetric |
| mean perimeter | 1.07 | 1.56 | Right-skewed |
| mean area | 1.93 | 5.46 | Right-skewed, Heavy-tailed |

| Feature | Skewness | Kurtosis | Assessment |
|------------------------|----------|----------|----------------------------|
| mean smoothness | 0.53 | 1.50 | Right-skewed |
| mean compactness | 0.91 | 0.54 | Right-skewed |
| mean concavity | 1.44 | 2.46 | Right-skewed |
| mean concave points | 1.15 | 1.18 | Right-skewed |
| mean symmetry | 0.52 | 0.34 | Right-skewed |
| mean fractal dimension | 1.26 | 3.76 | Right-skewed, Heavy-tailed |
| radius error | 3.30 | 17.46 | Right-skewed, Heavy-tailed |
| texture error | 0.75 | 1.02 | Right-skewed |
| perimeter error | 3.60 | 20.78 | Right-skewed, Heavy-tailed |
| area error | 5.87 | 45.73 | Right-skewed, Heavy-tailed |
| smoothness error | 1.03 | 1.22 | Right-skewed |

0.2.3 Feature Distributions

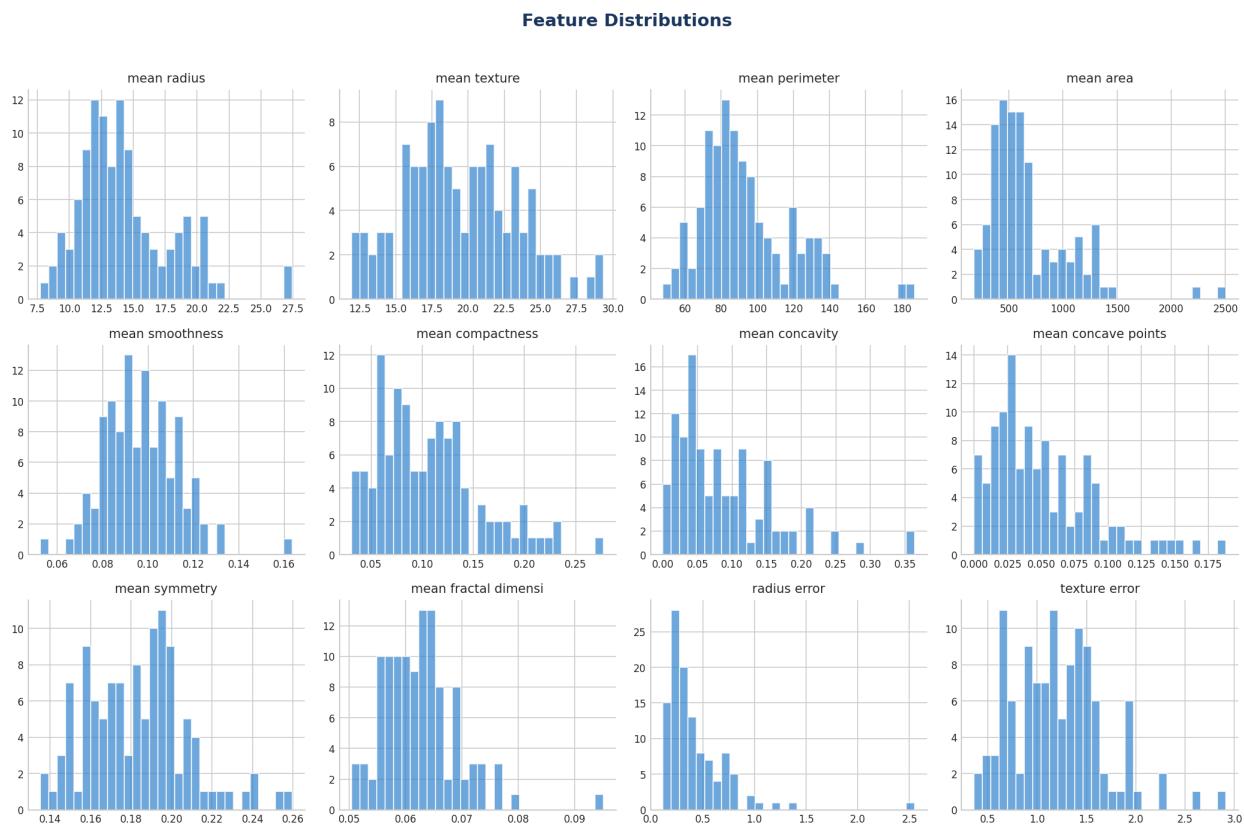


Figure 1: Feature Distributions

0.2.4 Outlier Analysis

Method: Interquartile Range (IQR) with 1.5x multiplier

Total Outliers Detected: 96 across 27 features

| Feature | Outliers | % of Data | Min | Max |
|-------------------------|----------|-----------|--------|---------|
| area error | 10 | 8.8% | 8.61 | 542.20 |
| fractal dimension error | 7 | 6.1% | 0.00 | 0.01 |
| concavity error | 6 | 5.3% | 0.00 | 0.09 |
| mean concavity | 5 | 4.4% | 0.00 | 0.36 |
| worst symmetry | 5 | 4.4% | 0.20 | 0.48 |
| mean area | 4 | 3.5% | 181.00 | 2501.00 |
| radius error | 4 | 3.5% | 0.12 | 2.55 |
| perimeter error | 4 | 3.5% | 0.77 | 18.65 |
| smoothness error | 4 | 3.5% | 0.00 | 0.02 |
| symmetry error | 4 | 3.5% | 0.01 | 0.04 |
| worst area | 4 | 3.5% | 268.60 | 4254.00 |
| worst smoothness | 4 | 3.5% | 0.09 | 0.22 |
| worst fractal dimension | 4 | 3.5% | 0.06 | 0.14 |
| mean compactness | 3 | 2.6% | 0.03 | 0.28 |
| mean concave points | 3 | 2.6% | 0.00 | 0.19 |

0.2.5 Outlier Distribution

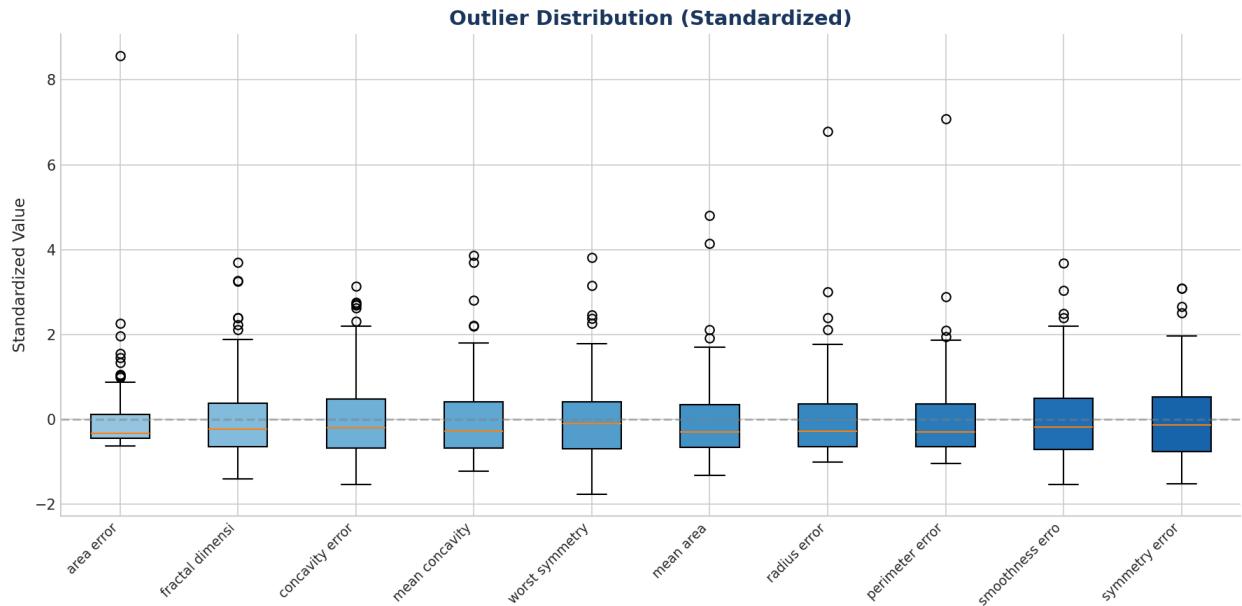


Figure 2: Outlier Boxplot

0.3 Correlation & Multicollinearity Analysis

Understanding feature relationships is critical for model interpretation and feature selection.

0.3.1 Correlation Matrix

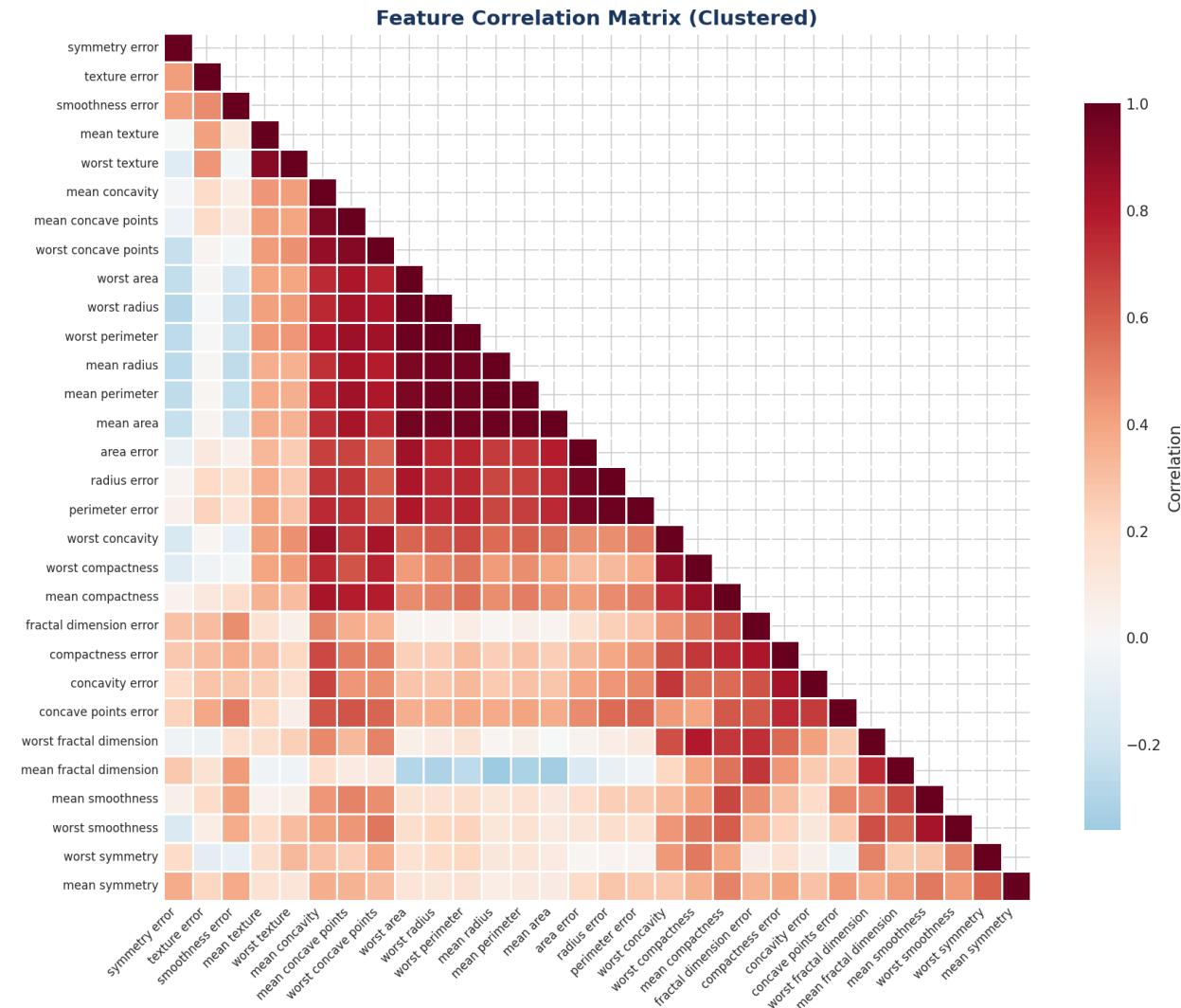


Figure 3: Correlation Matrix

0.3.2 Highly Correlated Feature Pairs ($|r| \geq 0.7$)

| Feature 1 | Feature 2 | Correlation |
|----------------|-----------------|-------------|
| mean radius | mean perimeter | 0.998 |
| worst radius | worst perimeter | 0.994 |
| mean perimeter | mean area | 0.983 |

| Feature 1 | Feature 2 | Correlation |
|-----------------|-----------------|-------------|
| mean radius | mean area | 0.983 |
| worst radius | worst area | 0.980 |
| radius error | perimeter error | 0.979 |
| worst perimeter | worst area | 0.976 |
| mean perimeter | worst radius | 0.969 |
| mean perimeter | worst perimeter | 0.969 |
| mean radius | worst radius | 0.968 |
| mean area | worst area | 0.966 |
| mean area | worst radius | 0.963 |
| mean radius | worst perimeter | 0.962 |
| mean area | worst perimeter | 0.959 |
| radius error | area error | 0.953 |

0.3.3 Variance Inflation Factor (VIF)

VIF measures multicollinearity. VIF > 5 indicates moderate, VIF > 10 indicates severe multicollinearity.

| Feature | VIF | Assessment |
|-------------------------|----------|------------|
| mean radius | 79262.63 | Critical |
| mean perimeter | 72990.12 | Critical |
| worst radius | 22342.62 | Critical |
| worst perimeter | 12079.97 | Critical |
| worst area | 2216.27 | Critical |
| mean area | 1946.14 | Critical |
| worst fractal dimension | 1185.56 | Critical |
| mean fractal dimension | 1108.49 | Critical |
| worst smoothness | 849.75 | Critical |
| mean smoothness | 811.67 | Critical |
| worst texture | 667.77 | Critical |
| radius error | 547.22 | Critical |
| perimeter error | 478.85 | Critical |
| mean texture | 478.71 | Critical |
| worst symmetry | 319.25 | Critical |

0.3.4 VIF Visualization

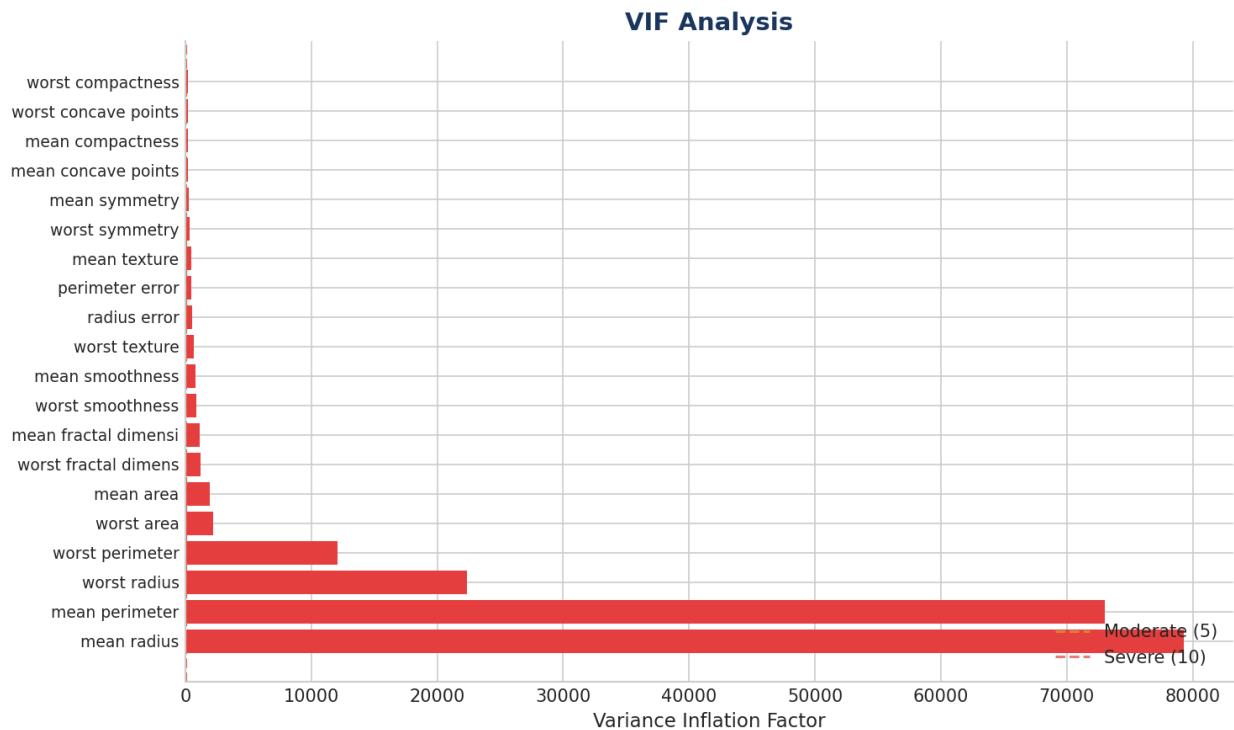


Figure 4: VIF Analysis

0.4 Data Profile

0.4.1 Dataset Overview

- **Total Samples:** 569
- **Total Features:** 30

0.4.2 Data Types

| Data Type | Count |
|-----------|-------|
| float64 | 30 |

0.4.3 EDA Recommendations

- All features are numeric - no encoding needed
- Features describe cell nuclei measurements
- Consider feature scaling for distance-based models

0.5 Feature Importance Analysis

Feature importance measures how much each feature contributes to the model's predictions. Higher values indicate more influential features.

0.5.1 Top 20 Features

| Rank | Feature | Importance |
|------|----------------------|------------|
| 1 | worst texture | 1.2551 |
| 2 | radius error | 1.0830 |
| 3 | worst concave points | 0.9537 |
| 4 | worst area | 0.9478 |
| 5 | worst radius | 0.9476 |
| 6 | worst symmetry | 0.9392 |
| 7 | area error | 0.9291 |
| 8 | worst concavity | 0.8232 |
| 9 | worst perimeter | 0.7632 |
| 10 | worst smoothness | 0.7466 |
| 11 | mean concave points | 0.7042 |
| 12 | mean compactness | 0.6483 |
| 13 | compactness error | 0.6472 |
| 14 | mean concavity | 0.6021 |
| 15 | mean texture | 0.5527 |
| 16 | perimeter error | 0.5443 |
| 17 | mean area | 0.5411 |
| 18 | mean radius | 0.5115 |
| 19 | mean perimeter | 0.4763 |
| 20 | concave points error | 0.4438 |

0.5.2 Feature Importance Visualization

0.6 Model Benchmarking

Multiple machine learning algorithms were evaluated using 5-fold cross-validation. The table below shows the performance of each model.

0.6.1 Model Comparison

| Model | CV Score | Std Dev | Test Score | Time (s) |
|---------------------|----------|--------------|------------|----------|
| Logistic Regression | 0.9802 | ± 0.0128 | 0.9825 | 0.53 |
| Random Forest | 0.9538 | ± 0.0235 | 0.9561 | 0.17 |
| Gradient Boosting | 0.9560 | ± 0.0139 | 0.9561 | 0.49 |

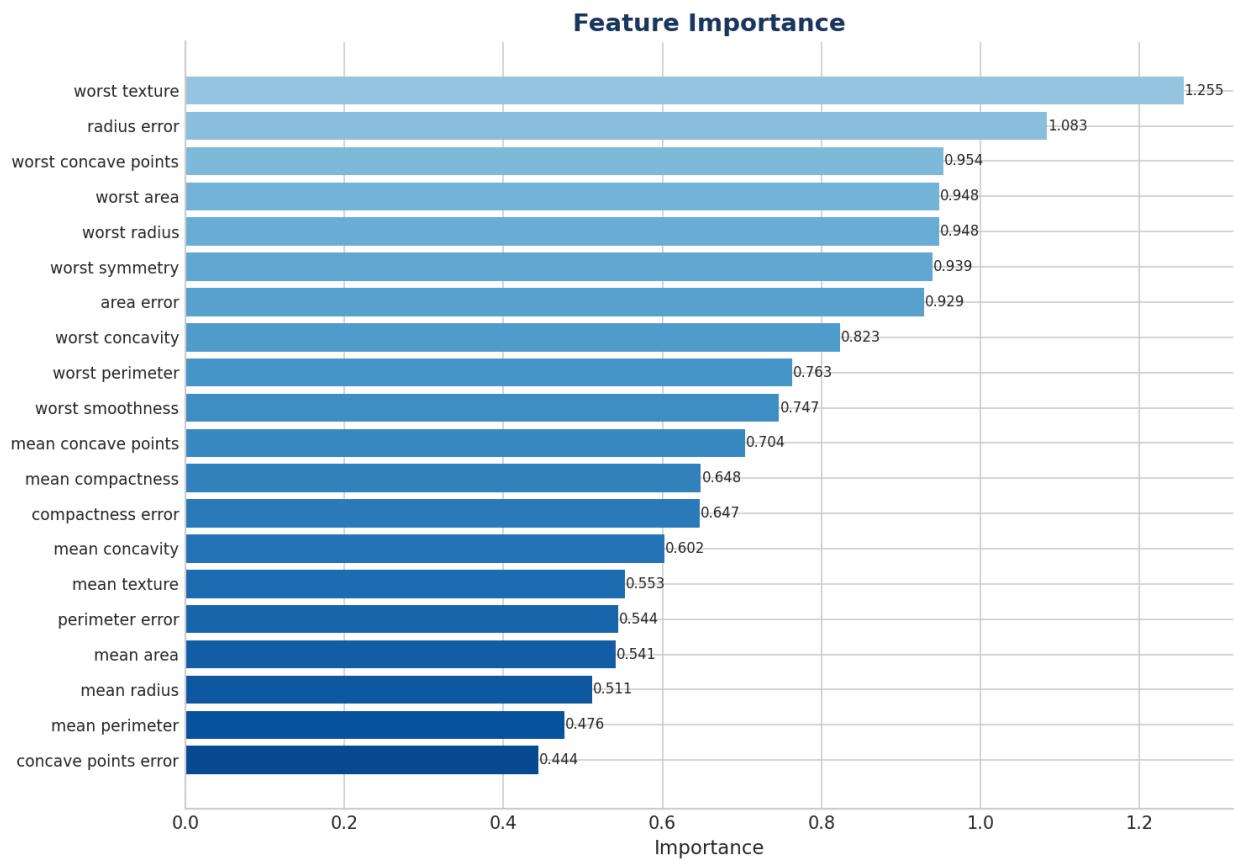


Figure 5: Feature Importance

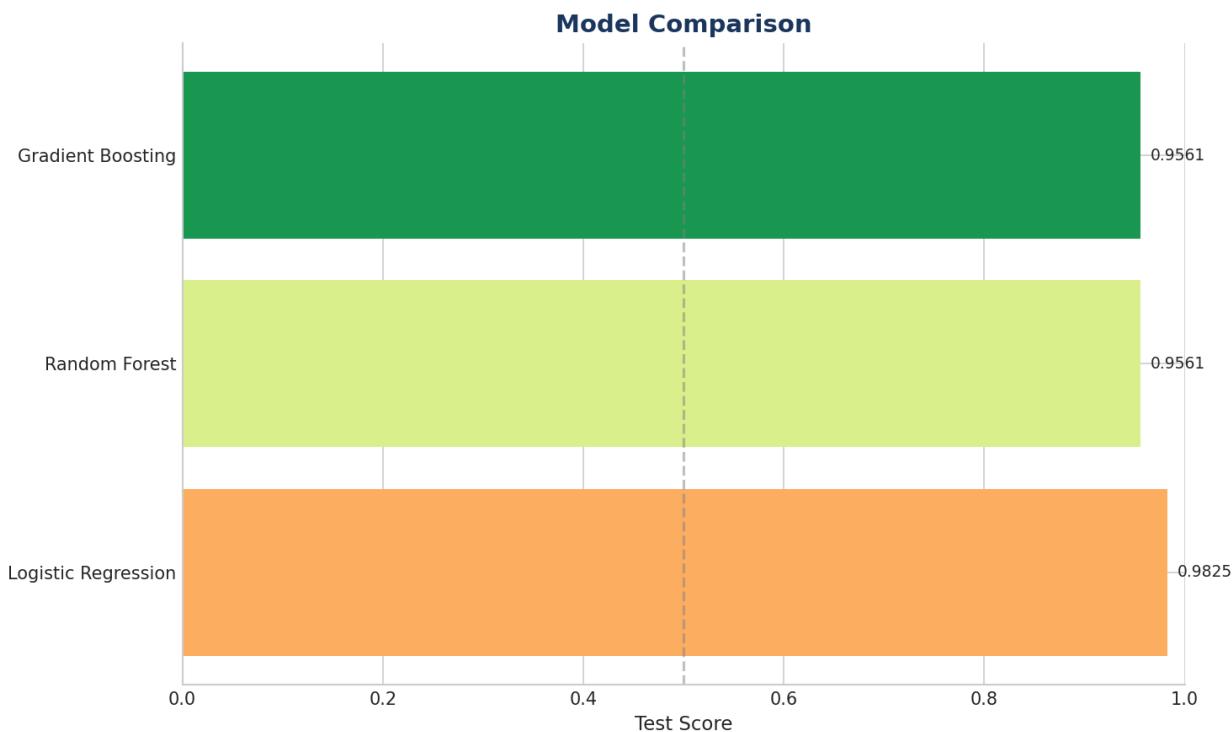


Figure 6: Model Benchmarking

0.6.2 Performance Visualization

0.7 Learning Curve Analysis

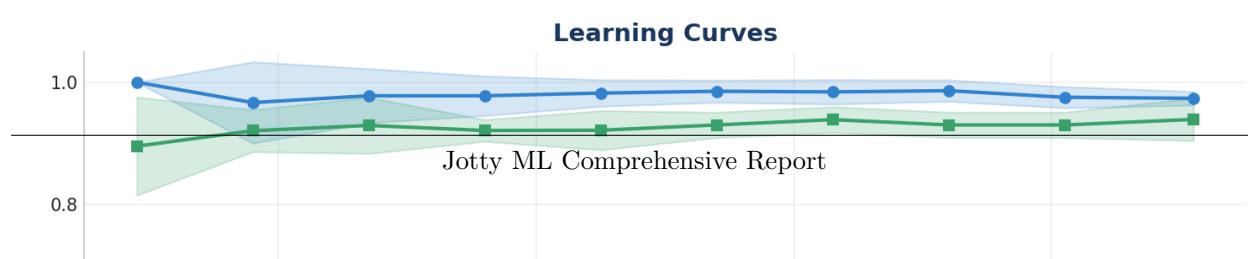
Learning curves reveal how model performance changes with training data size, helping diagnose underfitting vs overfitting.

0.7.1 Bias-Variance Diagnosis

Good Fit: Model has balanced bias-variance tradeoff.

| Metric | Value |
|------------------------|--------|
| Final Training Score | 0.9736 |
| Final Validation Score | 0.9391 |
| Gap (Train - Val) | 0.0345 |
| Training Samples Used | 91 |

0.7.2 Learning Curve Visualization



Jotty ML Comprehensive Report

0.7.3 Interpretation Guide

- **Converging curves** with small gap → Good fit
 - **Flat training curve** at low score → High bias, need more complex model
 - **Large gap** between curves → High variance, need regularization or more data
 - **Curves still improving** → May benefit from more training data
-

0.8 Cross-Validation Detailed Analysis

5-fold cross-validation provides robust performance estimates and helps detect instability.

0.8.1 Fold-by-Fold Results

| Fold | Train Accuracy | Test Accuracy | Train F1 | Test F1 |
|------|----------------|---------------|----------|---------|
| 1 | 0.9780 | 0.9130 | 0.9762 | 0.9115 |
| 2 | 0.9670 | 0.9565 | 0.9641 | 0.9533 |
| 3 | 0.9890 | 0.9130 | 0.9883 | 0.9042 |
| 4 | 0.9780 | 0.9130 | 0.9762 | 0.9042 |
| 5 | 0.9565 | 1.0000 | 0.9527 | 1.0000 |

0.8.2 Stability Analysis

| Metric | Value |
|----------------|------------------|
| Mean Accuracy | 0.9391 |
| Std Deviation | 0.0348 |
| CV Coefficient | 3.70% |
| 95% CI | [0.8710, 1.0073] |

Stability Assessment: Good

0.8.3 CV Performance Distribution

0.9 Classification Performance

0.9.1 Classification Report

| Class | Precision | Recall | F1-Score | Support |
|-----------|-----------|--------|----------|---------|
| Malignant | 0.976 | 0.976 | 0.976 | 42 |
| Benign | 0.986 | 0.986 | 0.986 | 72 |

| Class | Precision | Recall | F1-Score | Support |
|-----------------|-----------|--------|--------------|---------|
| Accuracy | | | 0.982 | |

0.9.2 Confusion Matrix

0.10 ROC Curve Analysis

The Receiver Operating Characteristic (ROC) curve shows the trade-off between true positive rate and false positive rate at various classification thresholds.

0.10.1 Key Metrics

- **AUC-ROC:** 0.9954
- **Optimal Threshold:** 0.3659

0.10.2 ROC Curve

0.11 Precision-Recall Analysis

The Precision-Recall curve is especially useful for imbalanced datasets, showing the trade-off between precision and recall.

0.11.1 Key Metrics

- **Average Precision:** 0.9971

0.11.2 Precision-Recall Curve

0.12 Probability Calibration Analysis

Well-calibrated probabilities are essential for reliable decision-making. A perfectly calibrated model's predicted probabilities should match actual outcome frequencies.

0.12.1 Calibration Metrics

| Metric | Value | Interpretation |
|----------------------------|--------|-------------------------------|
| Brier Score | 0.0215 | Lower is better (0 = perfect) |
| Expected Calibration Error | 0.0222 | Lower is better |

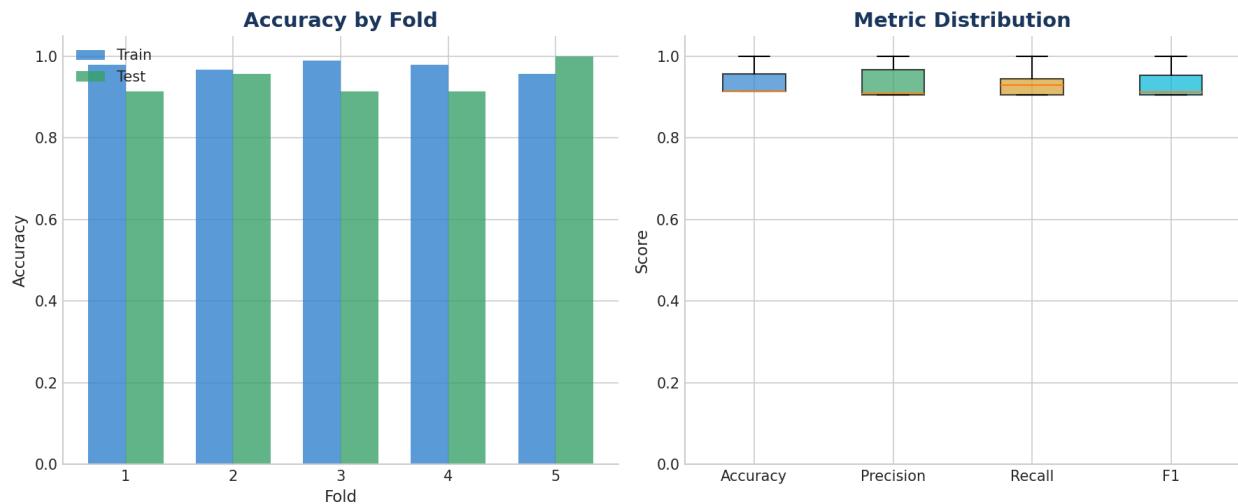


Figure 8: CV Analysis

0.12.2 Calibration Curve

0.12.3 Interpretation

- Points on diagonal = perfectly calibrated
- Points above diagonal = underconfident (probabilities too low)
- Points below diagonal = overconfident (probabilities too high)

0.13 Lift & Gain Analysis

These charts help evaluate model effectiveness for targeted campaigns and prioritization.

0.13.1 Key Metrics

| Metric | Value | Interpretation |
|--------------|--------|---|
| KS Statistic | 0.3596 | Maximum separation between model and random |
| KS at Decile | 64% | Optimal cutoff point |
| Top 10% Lift | 1.58x | Model advantage in top 10% |
| Top 20% Lift | 1.58x | Model advantage in top 20% |

0.13.2 Cumulative Gains & Lift Curves

0.13.3 Business Interpretation

- **Gains Curve:** Shows % of positives captured by targeting top X% of predictions
- **Lift Curve:** Shows how much better the model is vs random selection
- **KS Statistic:** Higher values indicate better model discrimination

0.14 Threshold Optimization

Jotty ML Comprehensive Report
Choosing the right classification threshold depends on business objectives.

0.14.1 Optimal Thresholds

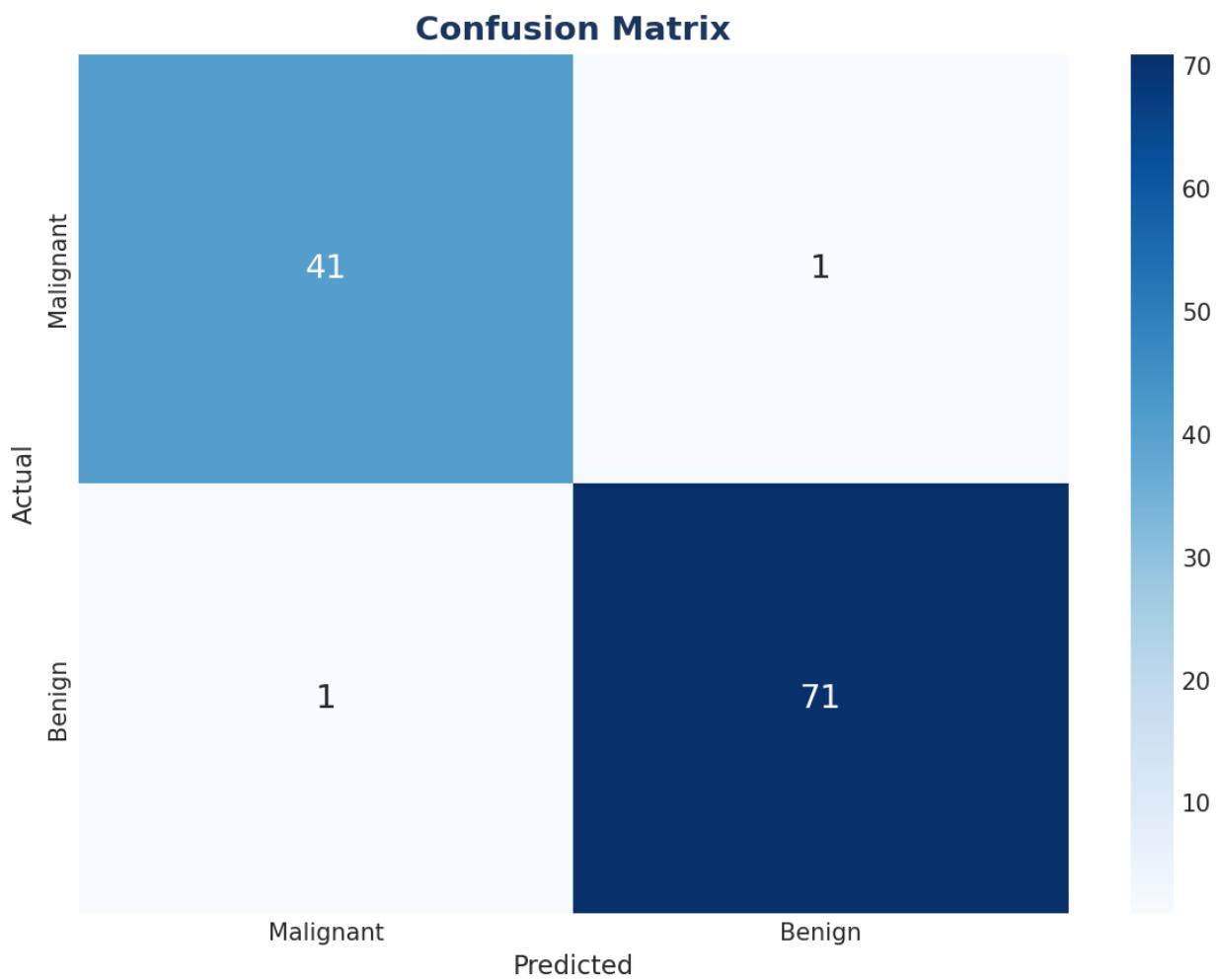


Figure 9: Confusion Matrix

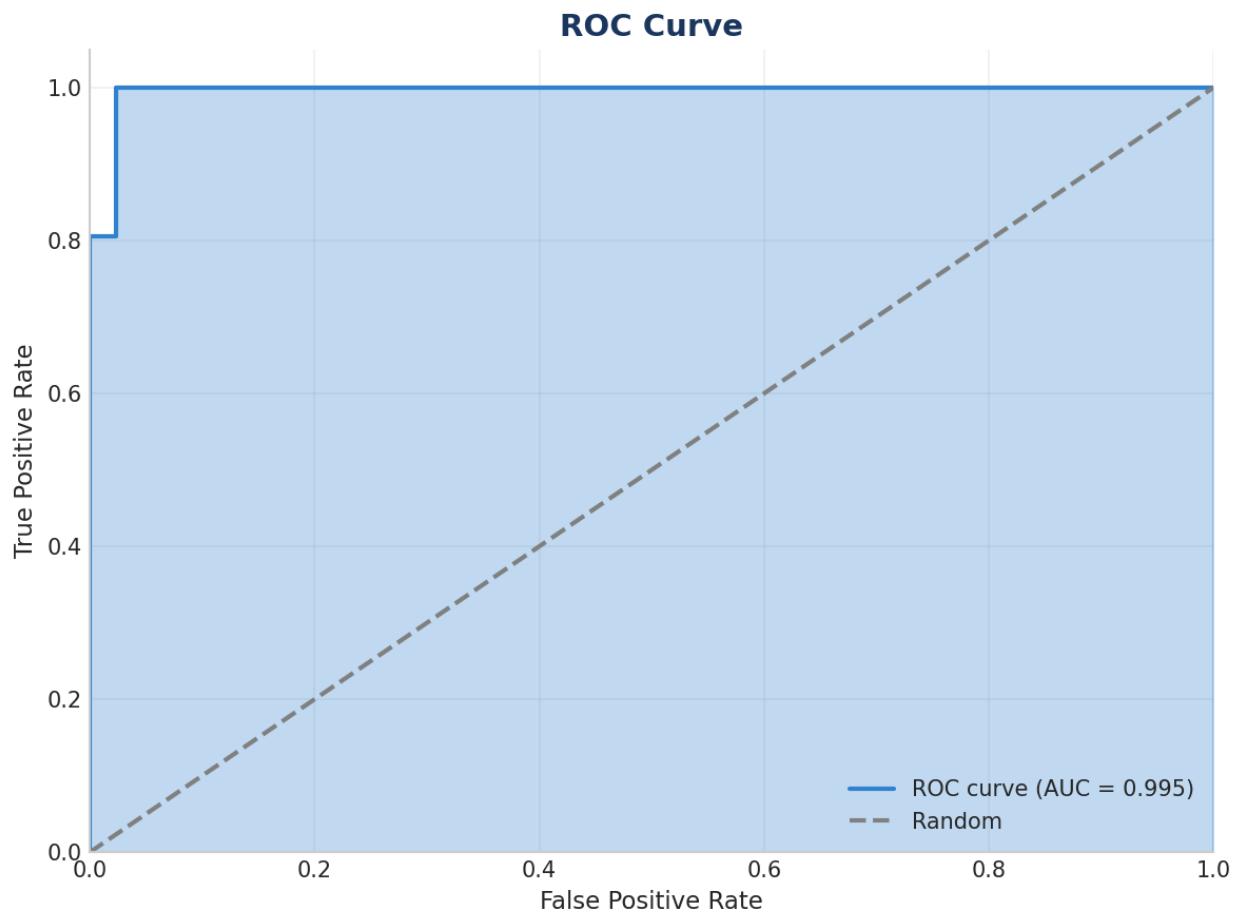


Figure 10: ROC Curve

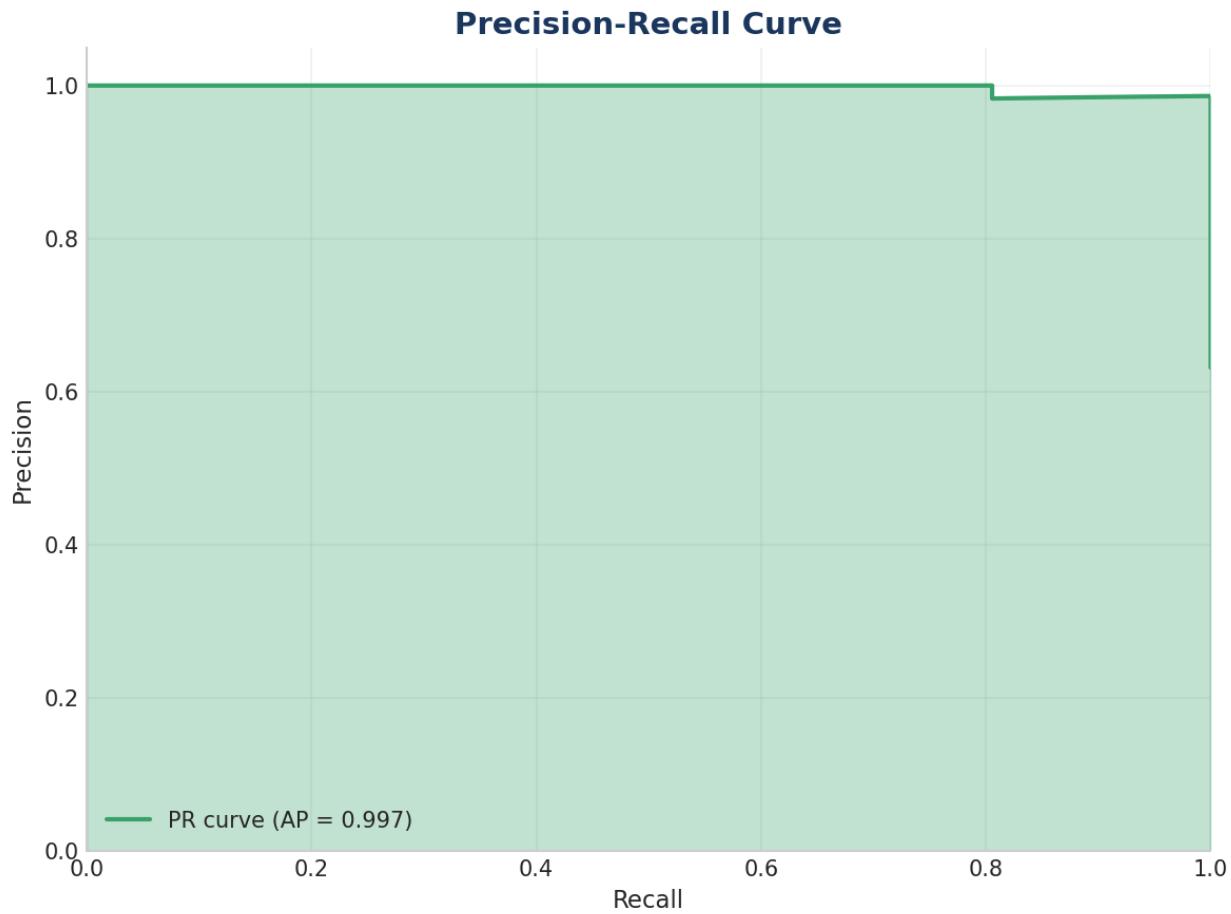


Figure 11: Precision-Recall Curve

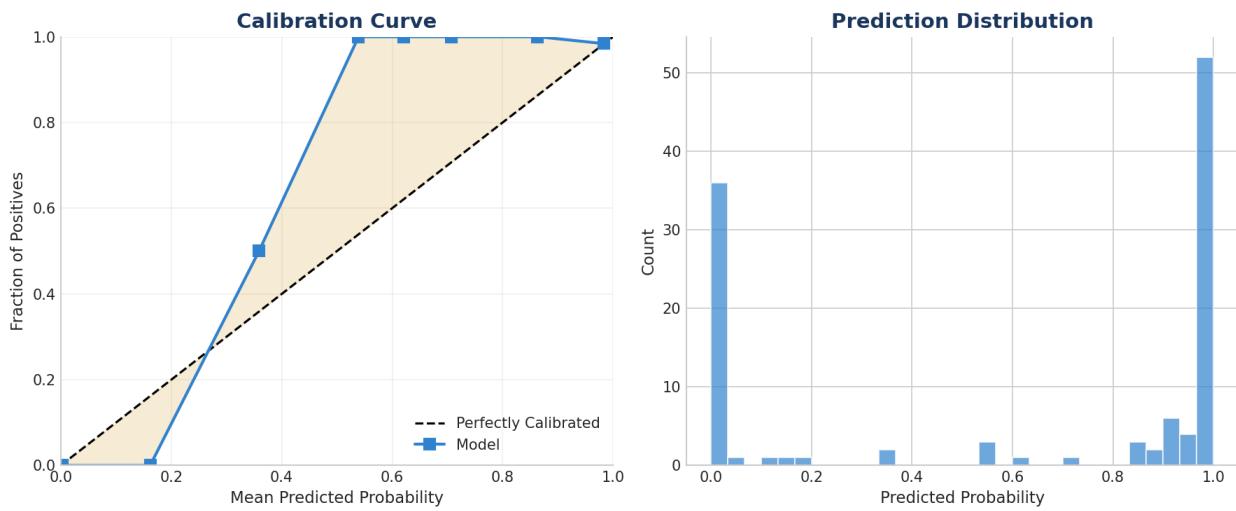


Figure 12: Calibration Curve

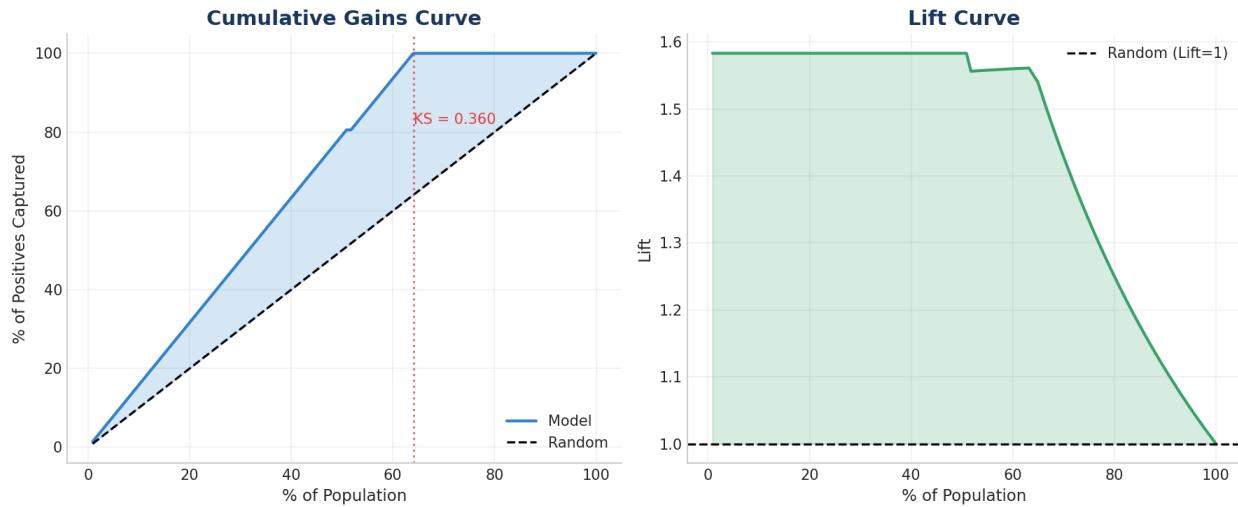
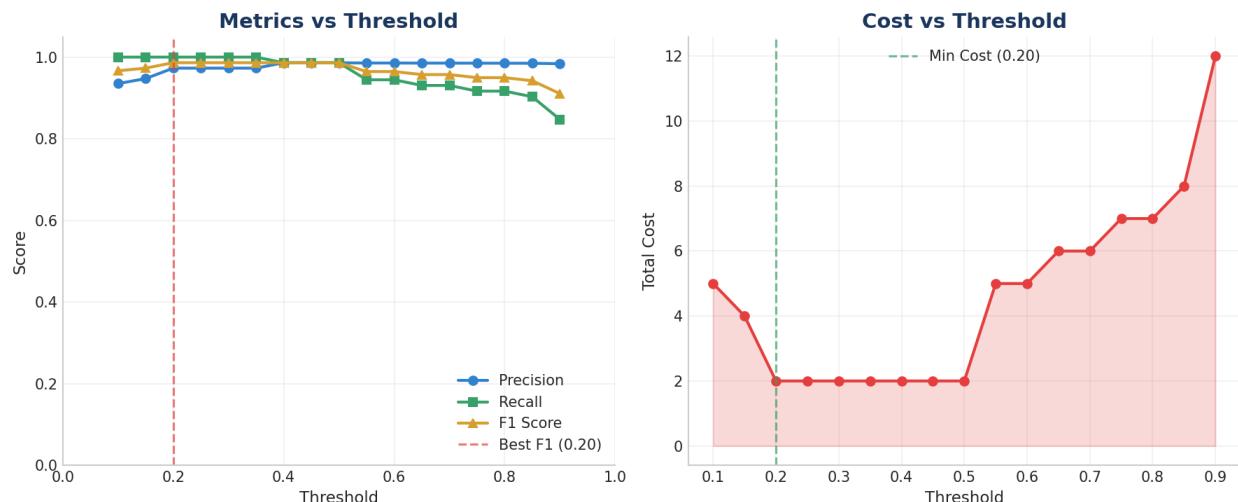


Figure 13: Lift and Gain Charts

| Threshold | TP | FP | FN | TN | Precision | Recall | F1 |
|-----------|----|----|----|----|-----------|--------|-------|
| 0.10 | 72 | 5 | 0 | 37 | 0.935 | 1.000 | 0.966 |
| 0.20 | 72 | 2 | 0 | 40 | 0.973 | 1.000 | 0.986 |
| 0.30 | 72 | 2 | 0 | 40 | 0.973 | 1.000 | 0.986 |
| 0.40 | 71 | 1 | 1 | 41 | 0.986 | 0.986 | 0.986 |
| 0.50 | 71 | 1 | 1 | 41 | 0.986 | 0.986 | 0.986 |
| 0.60 | 68 | 1 | 4 | 41 | 0.986 | 0.944 | 0.965 |
| 0.70 | 67 | 1 | 5 | 41 | 0.985 | 0.931 | 0.957 |
| 0.80 | 66 | 1 | 6 | 41 | 0.985 | 0.917 | 0.950 |
| 0.90 | 61 | 1 | 11 | 41 | 0.984 | 0.847 | 0.910 |

0.14.3 Threshold Visualization

Jotty ML Comprehensive Report
Figure 14: Threshold Analysis

0.14.4 Cost Parameters Used

| Metric | Value |
|--------------|--------|
| Total Errors | 2 |
| Error Rate | 1.75% |
| Accuracy | 98.25% |

0.15.2 Confusion Matrix Breakdown

- Class 0 misclassified as Class 1: 1 (2.4%)
- Class 1 misclassified as Class 0: 1 (1.4%)

0.15.3 Hardest to Classify Samples (Most Confident Errors)

| Sample | True | Predicted | Probability | Confidence |
|--------|------|-----------|-------------|------------|
| 53 | 0 | 1 | 0.909 | 0.817 |
| 16 | 1 | 0 | 0.366 | 0.268 |

0.15.4 Error Distribution Analysis

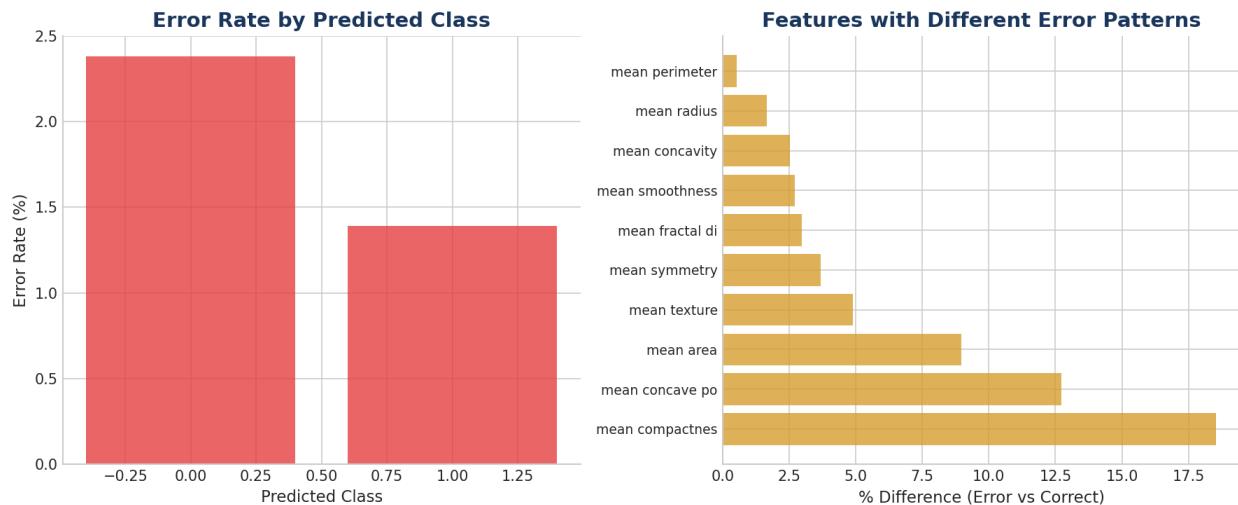


Figure 15: Error Analysis

0.16 SHAP Deep Analysis

SHAP (SHapley Additive exPlanations) provides consistent, locally accurate feature attributions for any machine learning model.

0.16.1 Global Feature Importance (Mean |SHAP|)

| Rank | Feature | Mean | SHAP |
|------|----------------------|--------|-------|
| 1 | worst texture | 0.9617 | 7.8% |
| 2 | radius error | 0.7973 | 14.2% |
| 3 | worst radius | 0.7614 | 20.3% |
| 4 | worst area | 0.7453 | 26.3% |
| 5 | worst concave points | 0.7220 | 32.1% |
| 6 | area error | 0.6391 | 37.3% |
| 7 | worst perimeter | 0.6109 | 42.2% |
| 8 | worst smoothness | 0.6033 | 47.1% |
| 9 | worst concavity | 0.5463 | 51.5% |
| 10 | worst symmetry | 0.5358 | 55.8% |
| 11 | mean concave points | 0.5358 | 60.1% |
| 12 | mean compactness | 0.4774 | 64.0% |
| 13 | compactness error | 0.4310 | 67.4% |
| 14 | mean area | 0.4174 | 70.8% |
| 15 | mean concavity | 0.3991 | 74.0% |

0.16.2 SHAP Summary Plot

Shows feature impact on predictions. Color indicates feature value (red=high, blue=low).

0.16.3 SHAP Feature Importance Bar

0.16.4 SHAP Dependence Plots (Top 3 Features)

Shows how feature values affect SHAP values, revealing non-linear relationships.

0.16.5 SHAP Waterfall (Sample Prediction)

Shows how features contribute to a single prediction.

0.17 Baseline Comparison

0.17.1 Performance Improvement

| Model | Score | Improvement |
|-------------------|---------------|-------------------------|
| Baseline | 0.6270 | - |
| Best Model | 0.9825 | +0.3555 (+56.7%) |

The final model achieves a **56.7%** improvement over the baseline.

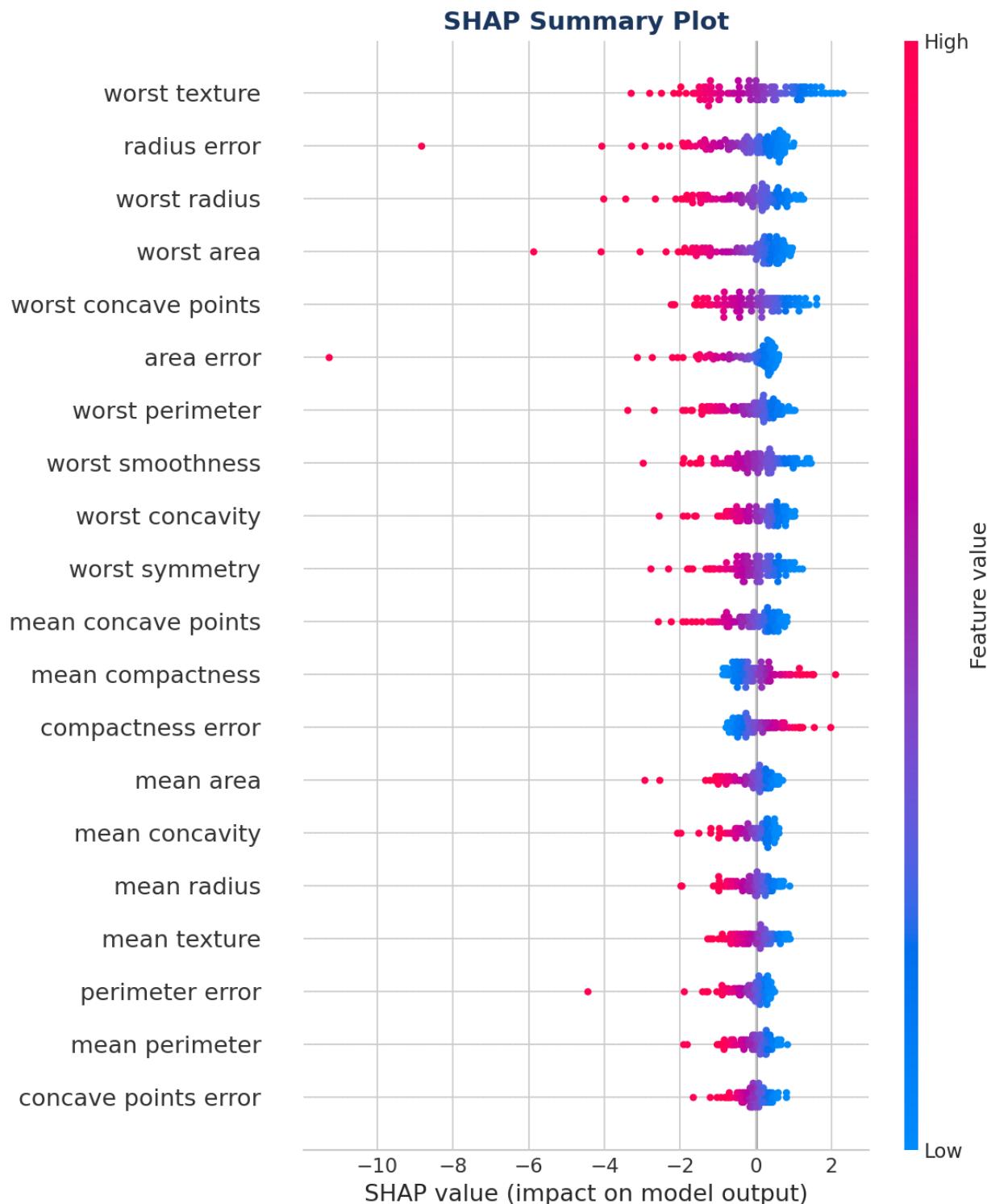


Figure 16: SHAP Summary

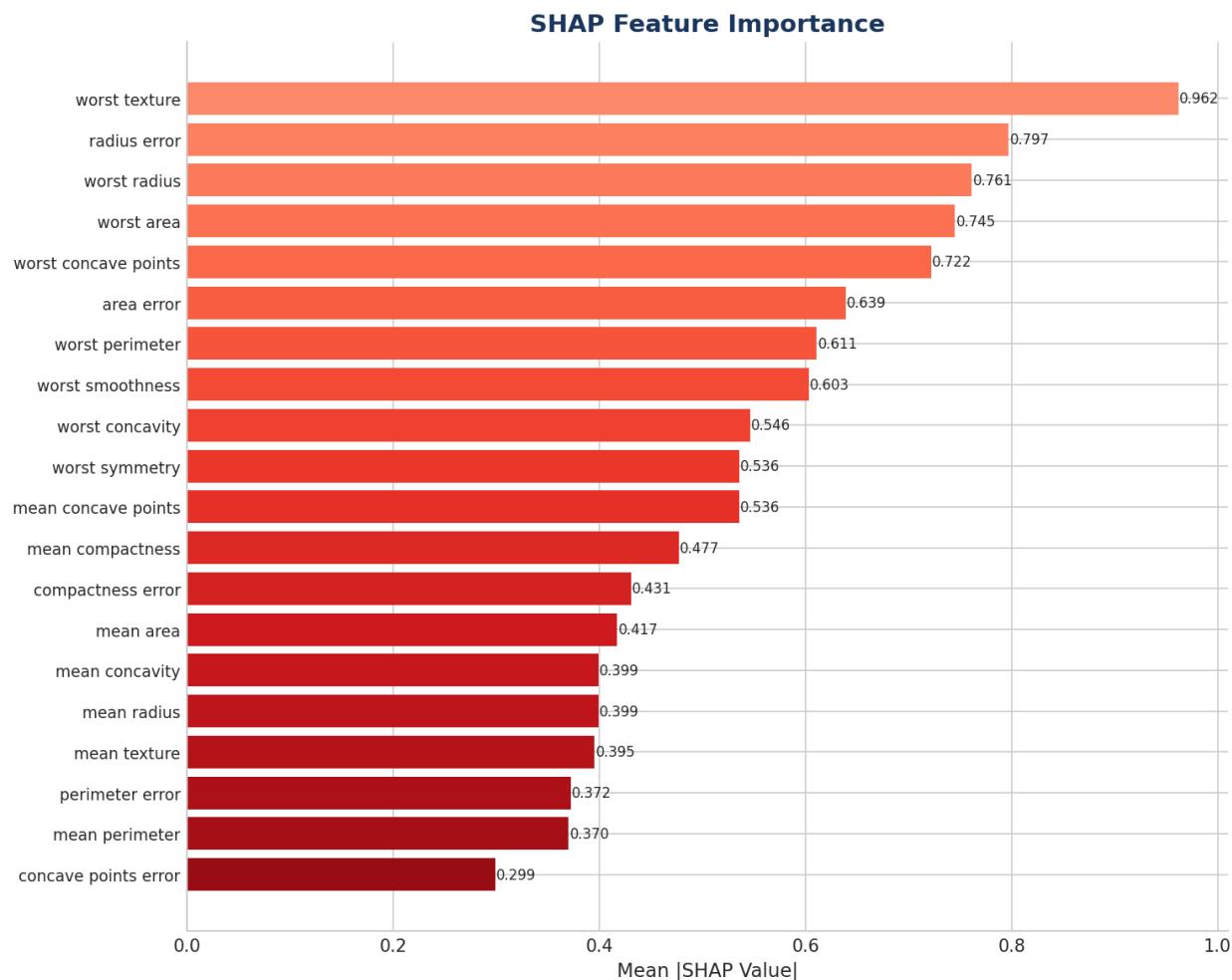


Figure 17: SHAP Bar Plot

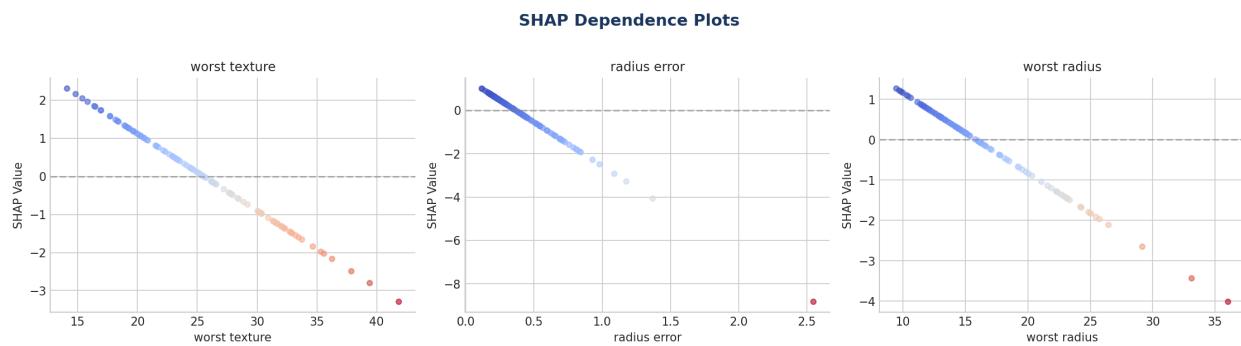


Figure 18: SHAP Dependence

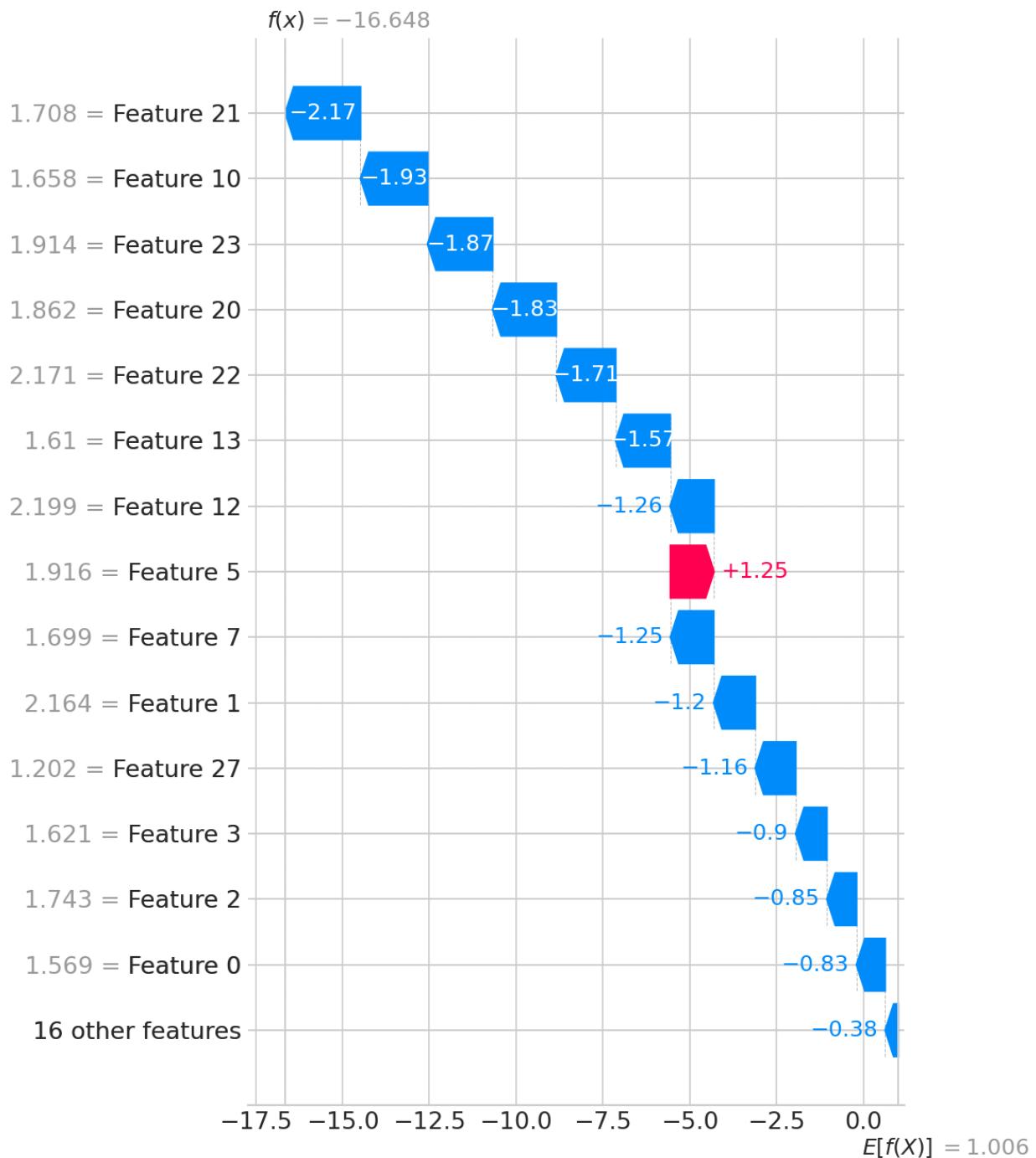


Figure 19: SHAP Waterfall

0.18 Recommendations & Next Steps

1. Excellent performance (98.2%) achieved - monitor for overfitting
 2. Logistic Regression provides good interpretability - ideal for regulated industries
 3. Top predictive features: worst texture, radius error, worst concave points
 4. Excellent discrimination (AUC=0.995) - suitable for production
 5. Monitor model performance over time for concept drift
 6. Validate on held-out data before production deployment
 7. Document model decisions for regulatory compliance
-

Report generated by Jotty SwarmMLComprehensive on 2026-02-05 03:18:31

0.19 Reproducibility

Full information for reproducing this analysis.

0.19.1 Model Configuration

Model Type: LogisticRegression

Hyperparameters:

| Parameter | Value |
|-----------|-------|
| | |

0.19.2 Random Seeds

| Component | Seed |
|-------------------|------|
| Main Random State | 42 |
| NumPy | 42 |
| Train/Test Split | 42 |

0.19.3 Environment

| Component | Version |
|----------------|---|
| Python Version | 3.11.2 |
| Platform | Linux-5.4.17-2136.312.3.4.el8uek.aarch64-aarch64-with-glibc2.36 |
| Processor | |

0.19.4 Package Versions

| Package | Version |
|------------|---------|
| numpy | 1.26.4 |
| pandas | 2.3.3 |
| matplotlib | 3.10.8 |
| seaborn | 0.13.2 |
| shap | 0.50.0 |

0.19.5 Generation Timestamp

Report Generated: 2026-02-05 03:18:31
