



GALLSTONE PREDICTION

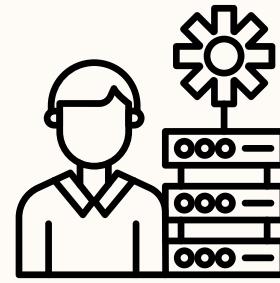
PRESENTED BY: GROUP 3



Project Team Overview

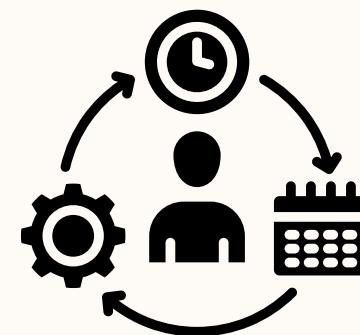


Team SignalX



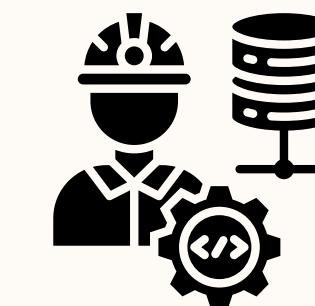
Yuyang Chen

GitHub: yuyangchen11



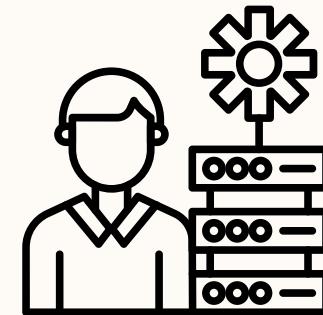
Ruihe Zhang

GitHub: Mudkipython



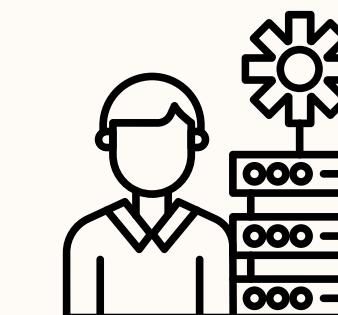
Zhaihan Gong

GitHub: yuehuatong



Wendy Xu

GitHub: ZihanXu12



Serena Sun

GitHub: YS - 02

Name of the GitHub repository: Gallstone_Prediction

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01

Business Context

Background

- Gallstones affect 10–15% of the global population
- Delayed diagnosis lead to complications and higher healthcare costs.

Current Limitations

- Ultrasound is accurate but not scalable for population screening
- Limited imaging capacity leads to delayed evaluation

Key Opportunity

- Use routine clinical data (e.g., BMI, lab tests, demographics) to build a **low-cost, scalable** risk stratification tool before imaging



02

Hypothesis

Research Question

Can routine non-acute clinical indicators support early gallstone risk stratification?

Core Hypothesis

- Routine demographic and metabolic indicators contain measurable predictive signal for pre-imaging risk stratification.

Hypothesis Testing Strategy

- H_0 : Routine non-acute metabolic and demographic indicators have no predictive power beyond random chance.
- Evaluate multiple models under stratified cross-validation.
- Compare logistic regression against tree-based ensembles.
- **Primary metric: PR-AUC**



We use PR-AUC because it better reflects positive-class detection performance in clinical screening settings.

03

Data

Dataset Snapshot

- Publicly available clinical dataset obtained from Kaggle, originally collected at the Internal Medicine Outpatient Clinic of Ankara VM Medical Park Hospital.
- 319 patients
- 161 positive (49.5%)
- 38 structured clinical features
- June 2022 – June 2023
- Ethics approved (E2-23-4632)
- Balanced disease distribution
- Non-imaging features only
- Cross-sectional; clinically diagnosed gallstone status



Demographics & Comorbidities

- Age
- Gender
- Comorbidity
- Coronary Artery Disease (CAD)
- Hypothyroidism
- Hyperlipidemia
- Diabetes Mellitus (DM)



Laboratory Markers

- Glucose
- Total Cholesterol (TC)
- LDL
- HDL
- Triglyceride
- AST
- ALT
- ALP
- Creatinine
- Glomerular Filtration Rate (GFR)
- C-Reactive Protein (CRP)
- Hemoglobin (HGB)
- Vitamin D



Anthropometrics

- Height
- Weight
- Body Mass Index (BMI)
- Obesity (%)



Bioimpedance – Body Composition

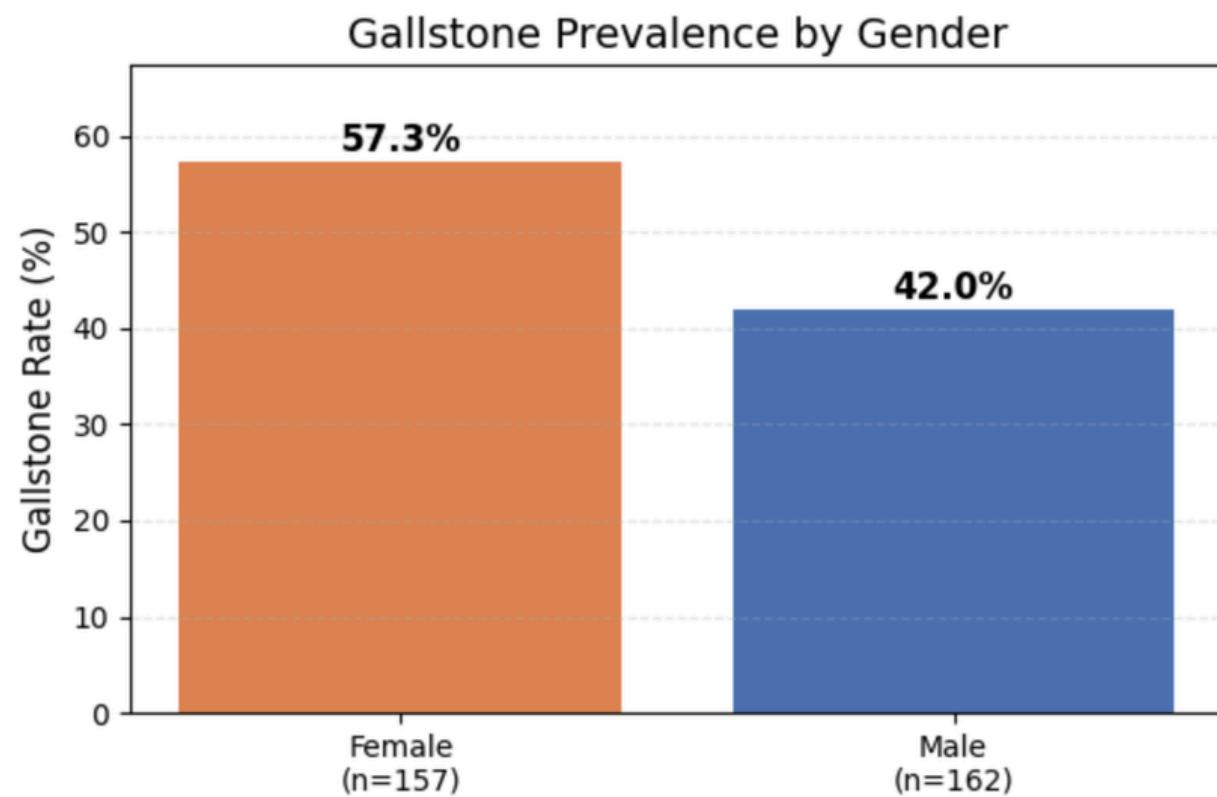
- Total Body Water (TBW)
- Extracellular Water (ECW)
- Intracellular Water (ICW)
- Extracellular Fluid / TBW (ECF/TBW)
- Total Body Fat Ratio (TBFR) (%)
- Total Fat Content (TFC)
- Lean Mass (LM) (%)
- Muscle Mass (MM)
- Bone Mass (BM)
- Body Protein (%)
- Visceral Fat Rating (VFR)
- Visceral Fat Area (VFA)
- Visceral Muscle Area (VMA)
- Hepatic Fat Accumulation (HFA)

03

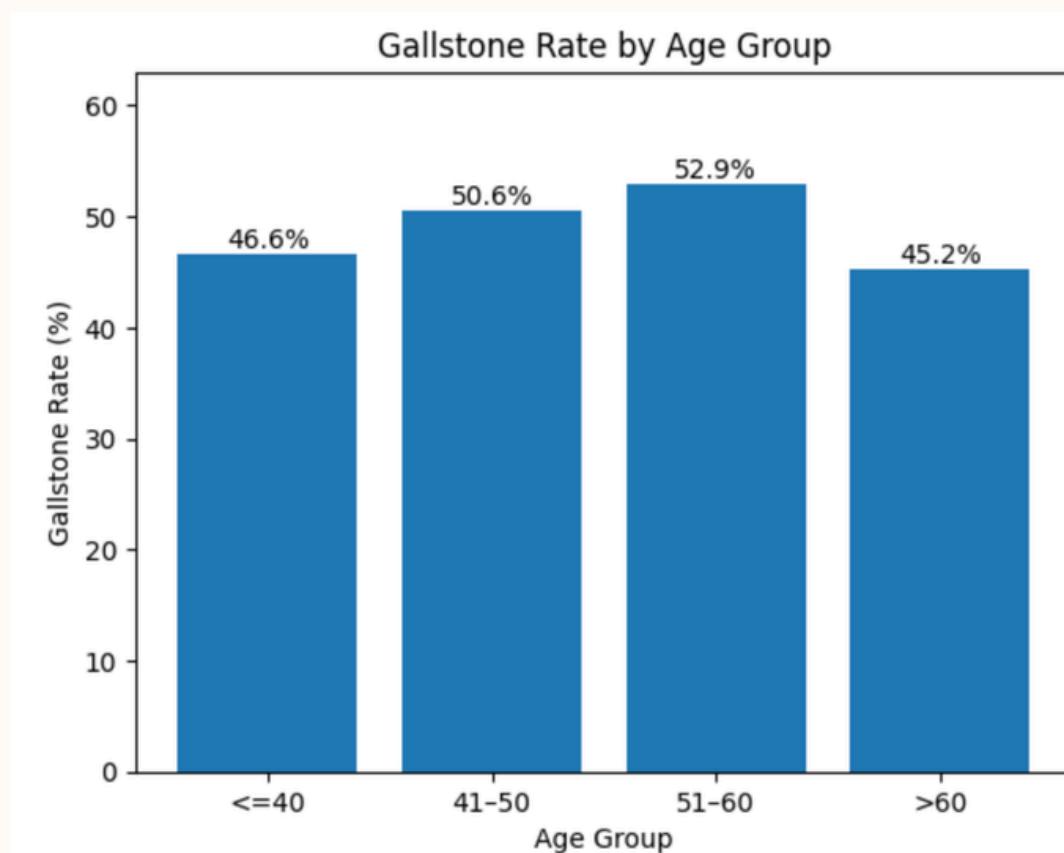
Data - Demographic Risk Patterns



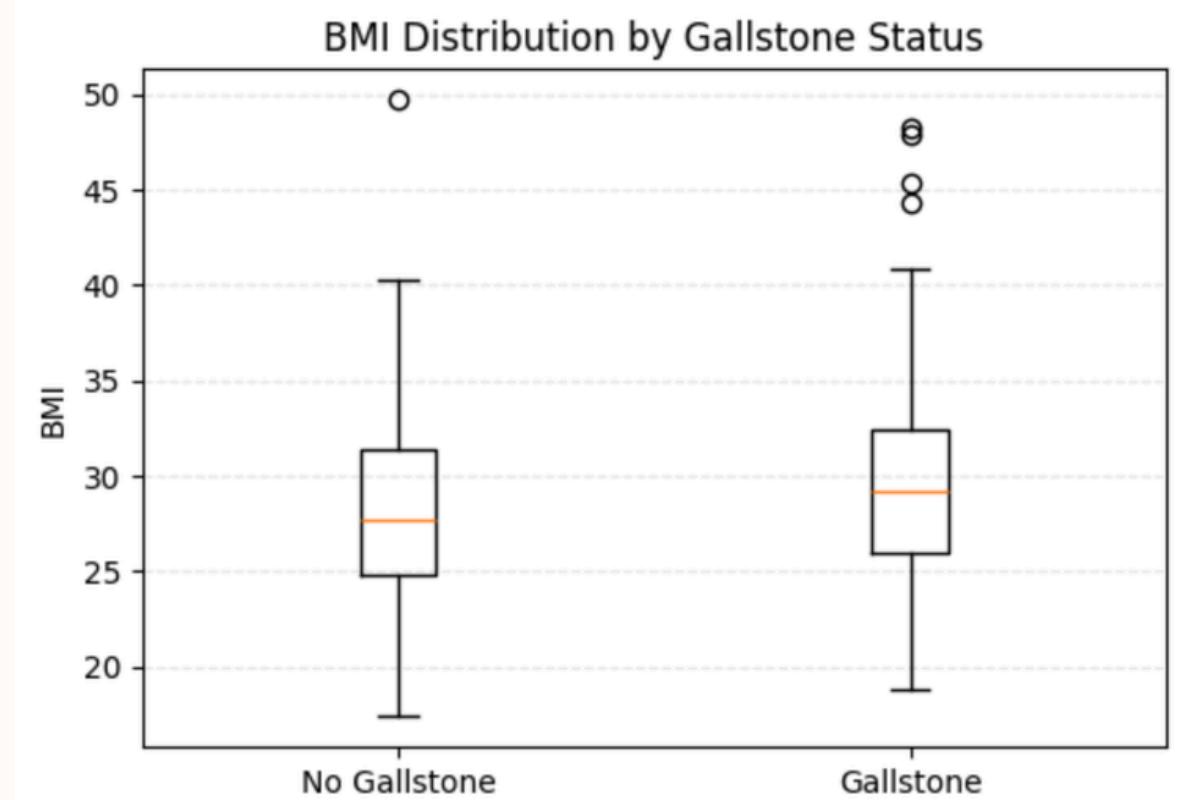
Gender vs Gallstone Rate



Age Group vs Gallstone Rate



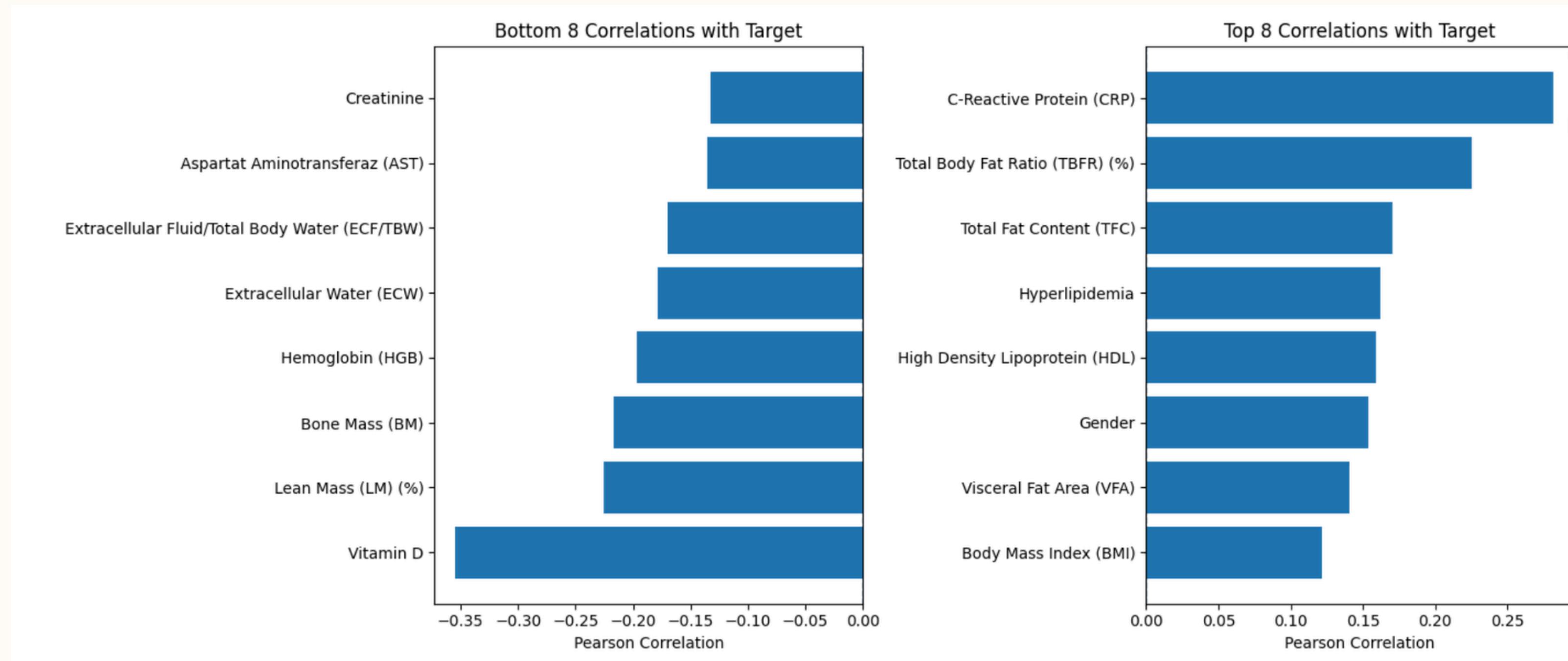
BMI Boxplot by Target



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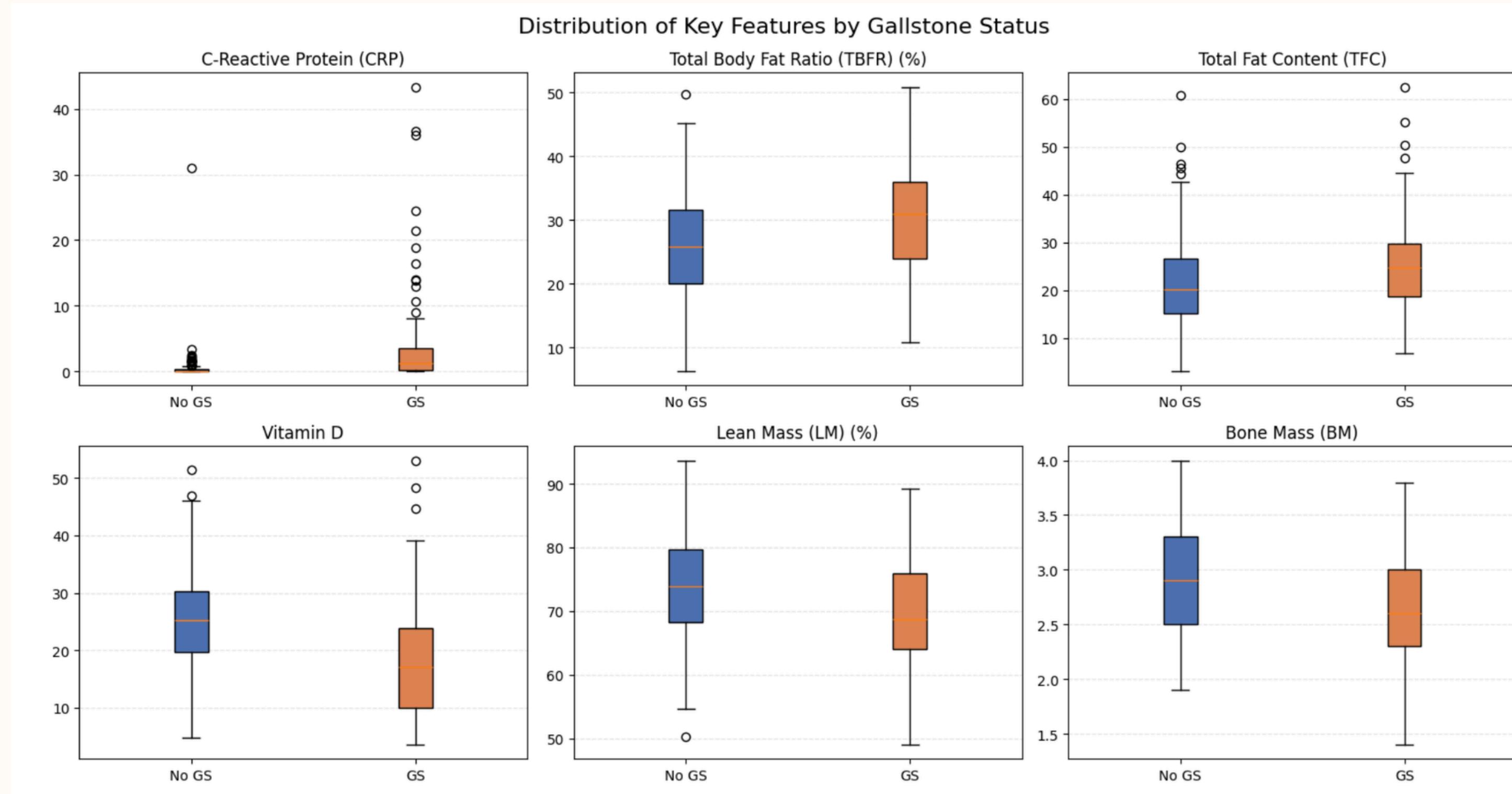
Data

Top/Bottom Correlated Features



Data

Top 3 positive & top 3 negative correlated features



03

Data

Final Feature Selection Strategy

Leakage Control – Removed Features

To ensure early-stage risk prediction and avoid data leakage, we excluded acute diagnostic markers:

- C-Reactive Protein (CRP)
- AST
- ALT
- ALP
- AST/ALT ratio

These markers reflect acute inflammatory or hepatobiliary status and may capture post-diagnostic signals. Removing them ensures we are truly modeling early-stage risk rather than symptomatic status.

Cross-Validated Performance (PR-AUC)

Method	# Features	PR-AUC
Mutual Information	20	0.829
RFECV	33	0.842
LASSO (Selected)	28	0.843

Final Decision

- LASSO achieved highest PR-AUC
- Provided automatic sparsity
- Reduced dimensionality ($33 \rightarrow 28$)
- Maintained interpretability

Data → Baseline Models → Cross-Validation → Feature Refinement → Final Model Selection

Modeling Strategy

- Supervised learning
- Binary classification (Gallstone: Yes/No)
- Stratified K-fold cross-validation
- Out-of-fold prediction for unbiased evaluation

Benchmark Models

- Logistic Regression
- Random Forest
- Histogram-based Gradient Boosting
- Light Gradient Boosting Machine
- XGBoost

Model Evaluation

- 319 samples, 5-fold stratified cross-validation
- 49.5% positive
- Hyperparameter tuning via cross-validation

Evaluation Metrics

- Primary: PR-AUC
 - precision–recall trade-off
- ROC-AUC
- Recall
- Calibration: Brier

Model Validation

Model Benchmark Results

Model Comparison

Model	ROC-AUC	PR-AUC	Recall	Brier
Logistic Regression	0.837	0.843	0.829	0.163
LightGBM	0.789	0.788	0.766	0.209
XGBoost	0.793	0.783	0.778	0.192
Random Forest	0.777	0.752	0.835	0.195
HistGB	0.789	0.789	0.937	0.199

Model Selection Strategy

Final Selection - Logistic Regression

Performance

- Best ROC-AUC (0.837) and PR-AUC (0.843)
- Lowest Brier Score → well-calibrated probabilities

Robustness

- Stable after removing confounders
- Less overfitting than tree-based models

Interpretability

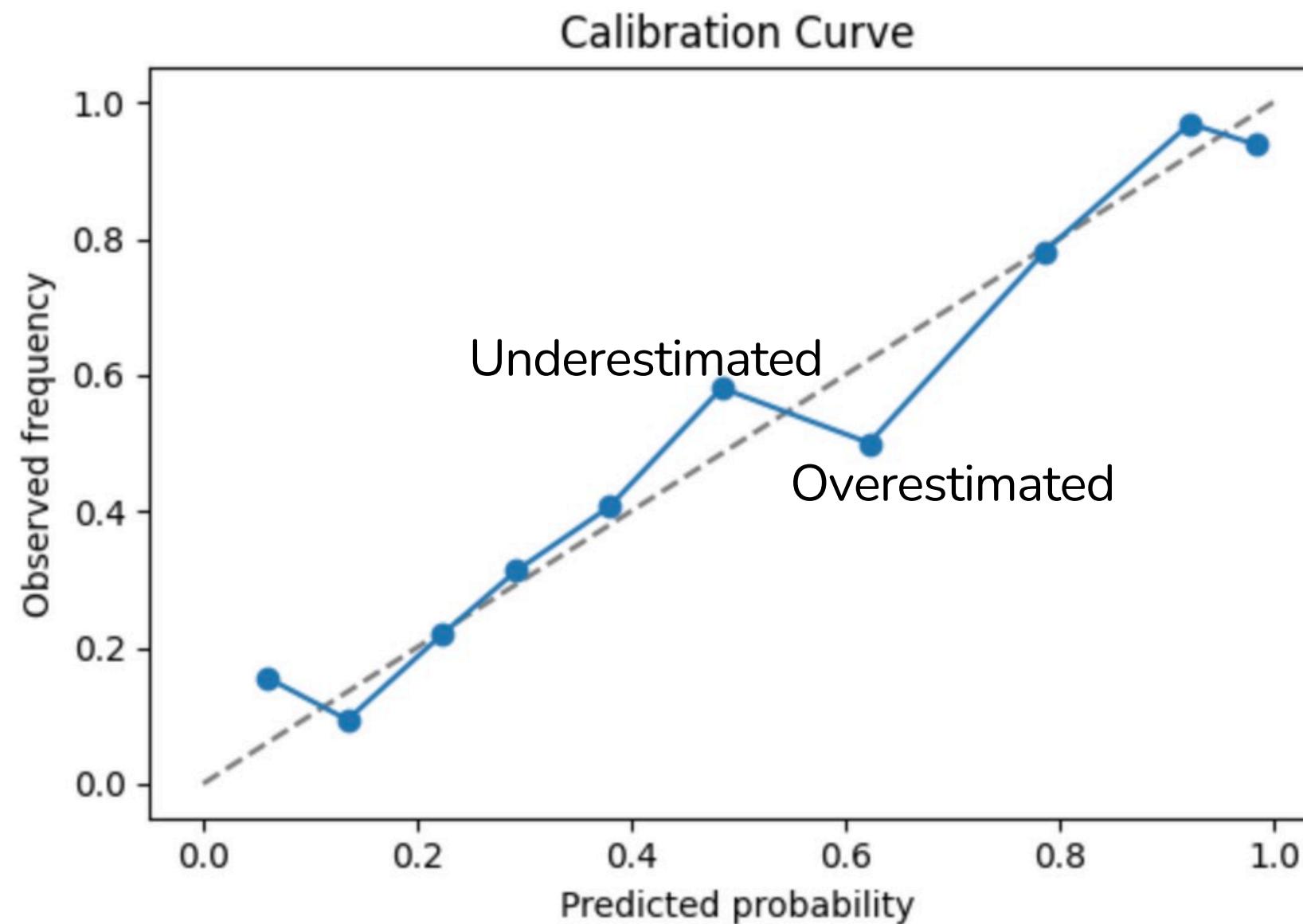
- Clinicians can understand and trust the prediction
- Enables a simple risk calculator for routine use

Deployment

- No complex infrastructure needed
- Easy integration into existing LIS/EMR systems

Model Performance Results

Compare with SOTA / Previous Work

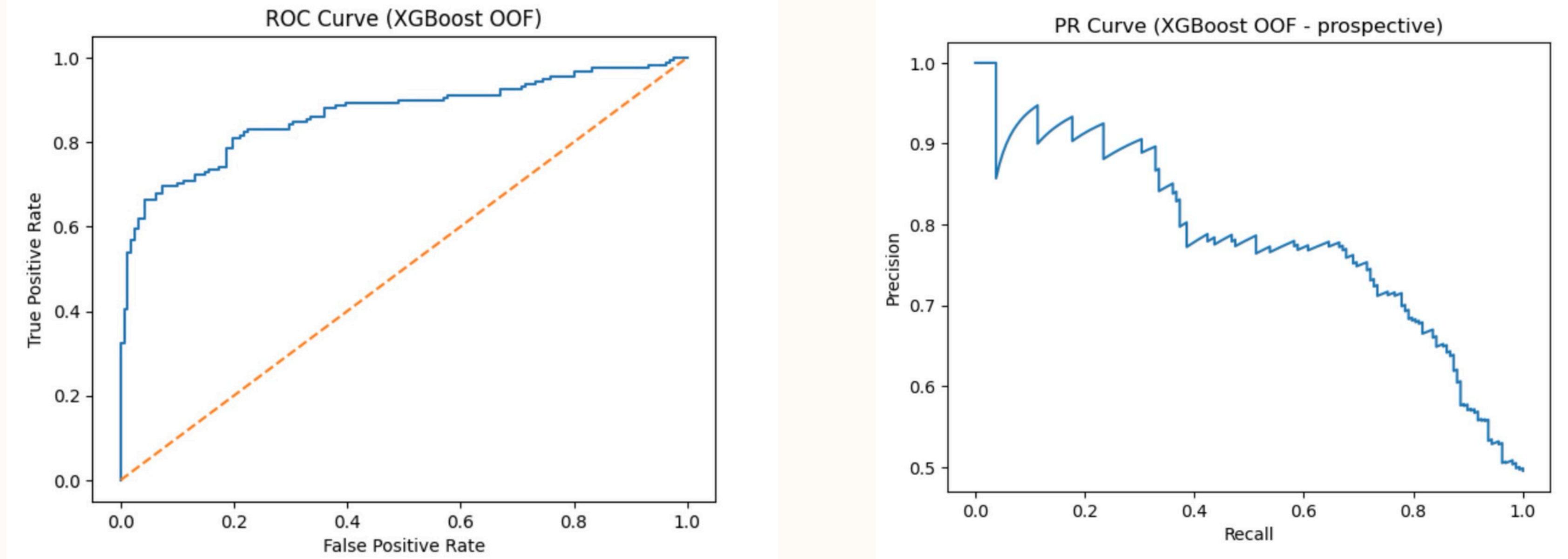


- Traditional clinical risk assessment relies on manual evaluation of metabolic indicators
- Modern SOTA methods for structured medical data use tree-based ensemble models
- Consistent with ensemble learning theory, boosting models in our experiments achieved strong positive-class discrimination
- Our final model (Logistic Regression) achieves comparable performance (PR-AUC = 0.843) while maximizing clinical interpretability for screening deployment

**OUR MODELLING STRATEGY BALANCES
SOTA PERFORMANCE WITH CLINICAL
ACTIONABILITY**

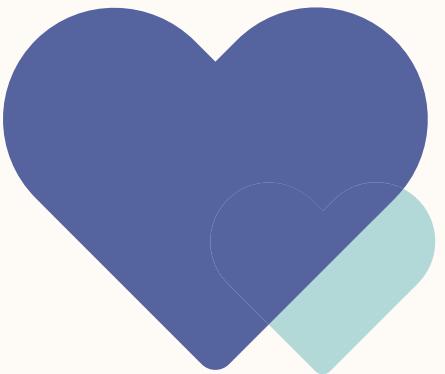
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Results



ROC / PR evaluate performance across all thresholds, and confirm strong discrimination
Clinical deployment requires choosing a specific decision threshold

Results - Threshold Selection



Why Threshold Matters

- Model outputs probabilities (0–1)
- Clinical decision requires binary classification (High / Low risk)

Threshold Selection Strategy

- Selected using OOF predictions
- Optimized by maximizing F1 score
- Balances precision and recall

Final Operating Threshold

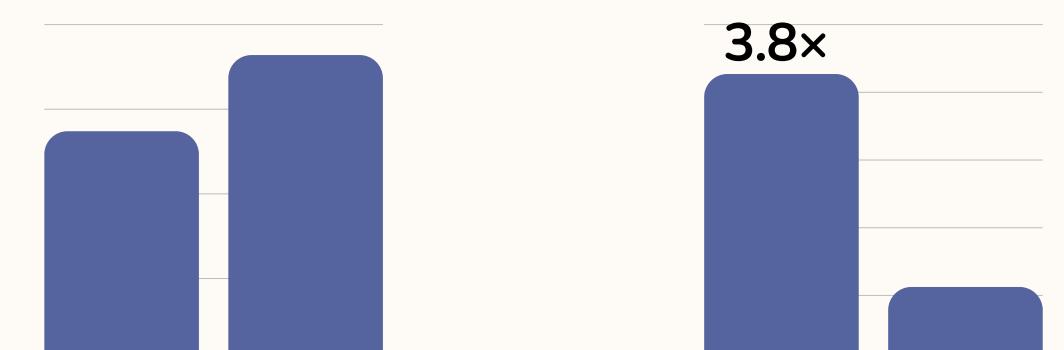
LR optimal threshold (OOF): 0.372

Decision rule:

- If predicted probability $\geq 0.372 \rightarrow$ High Risk
- Otherwise \rightarrow Low Risk

Risk Separation Result Under Threshold of 0.597

Risk Tire	Population	Observed Gallstone Rate
High Risk	175	74.9%
Low Risk	144	18.8%



Results - Error Analysis

Confusion Matrix (at threshold of 0.372)

	Predicted Low	Predicted High
Actual Low	136 (TN)	25 (FP)
Actual High	38 (FN)	120 (TP)

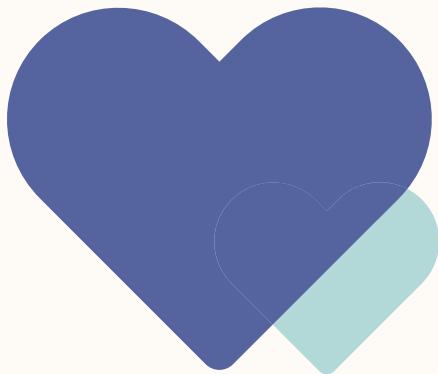
Error Structure

- False Negatives: 38 patients (24.1%)
- False Positives: 25 patients (15.5%)
- Model favors precision over recall

Key Takeaways



- High recall (76%) supports use as a screening tool
- Moderate false positives acceptable for early detection
- Remaining 24.1% FN highlights need for physician oversight



How should strategy adapt to hospital capacity?

Total: 319

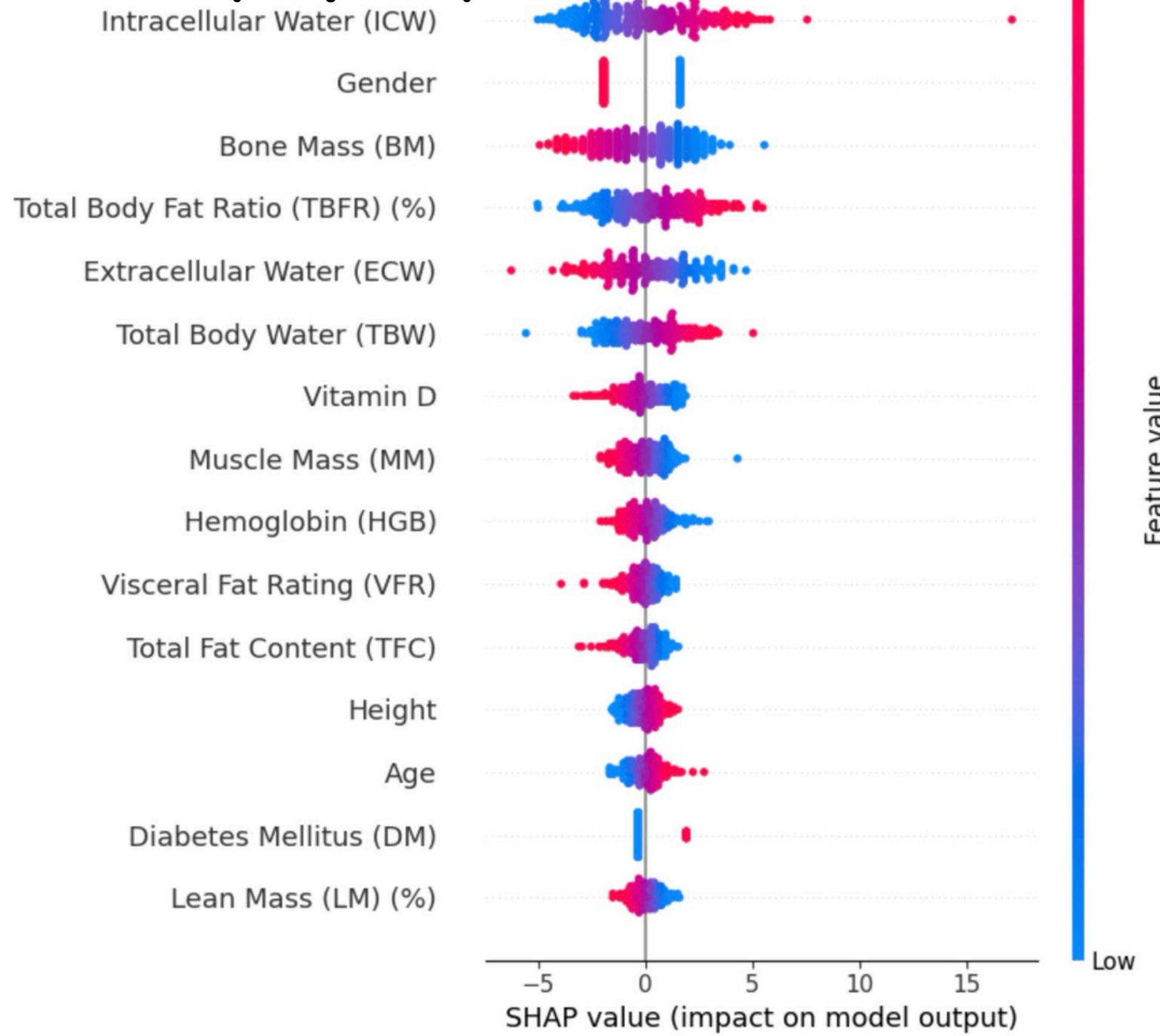
Policy	Patient Flagged	Missed Cases	Extra Follow-ups (FP)	F1
Balanced Threshold 0.441	175	27	44	0.787
Recall Focused (reduce missed diagnoses) Threshold 0.54	254	8	104	0.728
Top 50 (resource-limited)	50	110	2	0.462

- F1-opt balances precision and recall
- Cost-sensitive maximizes recall, minimizing missed cases
- Top-k approach prioritizes highest-risk individuals

Different operating thresholds enable flexible clinical deployment depending on resource availability.

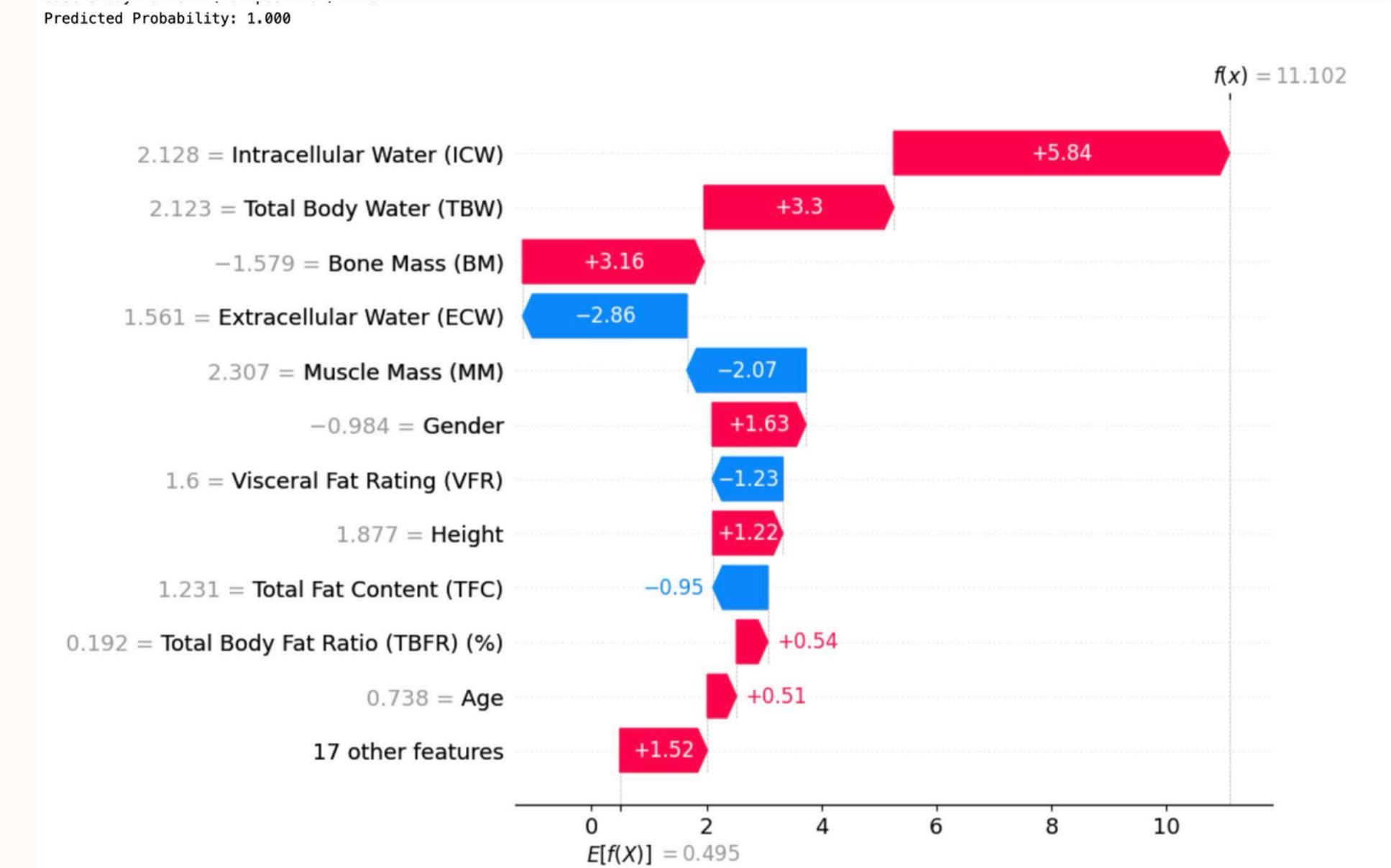
Feature Importance & Case Study

Global (Top 15)



High-Risk Patient Example

Patient's predicted risk = 1.00 (> 0.372 threshold) → flagged as high-risk



support hypothesis: routine demographic and metabolic indicators predict gallstone risk

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Threats to Validity

01

Sample Size & Generalizability

Model was trained on a relatively small sample (N=319)

02

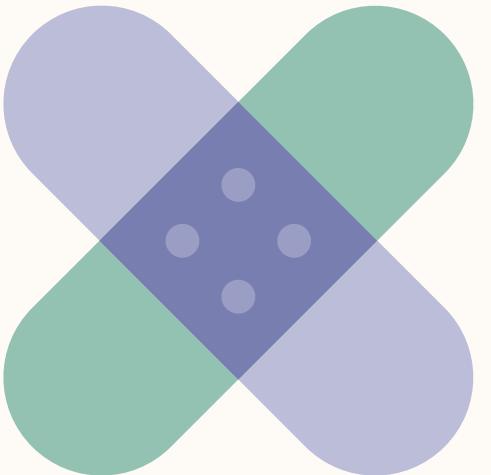
Lack of External Validation

Not yet validated on an independent external cohort

03

Implementation

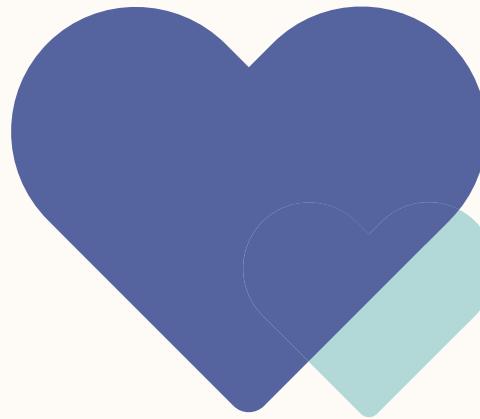
Real-world implementation needs system integration, governance approval, and monitoring



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Conclusions & Recommendations

We recommend pilot deployment under controlled settings



SCREENING CENTERS

- Risk-based screening allocation

CLINICIANS

- SHAP explanations enhance interpretability
- Designed as decision-support, not replacement

POPULATION HEALTH

- Early risk identification using routine data

The proposed model demonstrates strong discriminative ability (ROC-AUC ≈ 0.85) and clinically meaningful risk stratification

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Lessons Learned & Next Steps

Lessons Learned

- Model performance alone is insufficient
- deployment strategy and threshold design determine impact

Next Steps

01

Validate on an independent external cohort

02

Collect longitudinal data for stronger evidence

03

Conduct a prospective pilot study in screening settings

04

Gather clinician feedback for iterative refinement

05

Establish monitoring framework to track model performance and data drift after deployment.



THANK YOU!