RES-129



Explainable Machine Learning for Predicting CLABSI in Severe Burn Patients : A 10—Year Time-Series Analysis

Kibum Jeon¹, Seri Jeong¹, Wonkeun Song¹, Jae–Seok Kim¹, Han–Sung Kim¹, Kyu Sung Shin¹, Hyun Soo Kim¹, Seung Beom Han² ¹Department of Laboratory Medicine, Hallym University College of Medicine, Republic of Korea ²Department of Pediatrics, Catholic University College of Medicine, Republic of Korea

Backgrounds

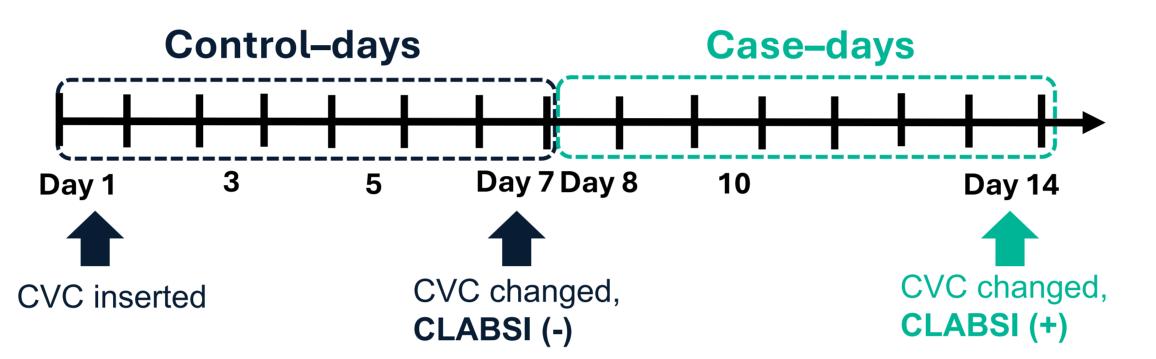
- Patients with severe burns are at a high risk for central line-associated bloodstream infections (CLABSI)
- Aim of Study: Identify risk factors for CLABSI in severe burn patients using time—series data , analyze via a machine learning (ML) model with explainable AI (XAI)



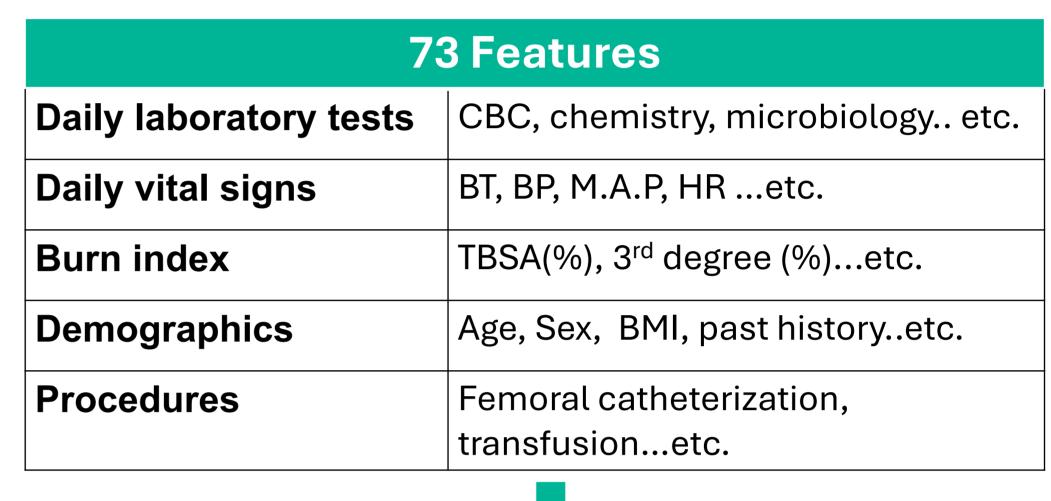
2 Methods

Burn center in Seoul **Daily data: Jan 2015 – Dec 2024**

Inclusion Criteria : ICU patients who underwent routine CVC changes



- Training Set: Test Set = 8:2, 5-fold cross validation
- ML method: XGBoost
- **SHAP**: ensured ML model interpretability



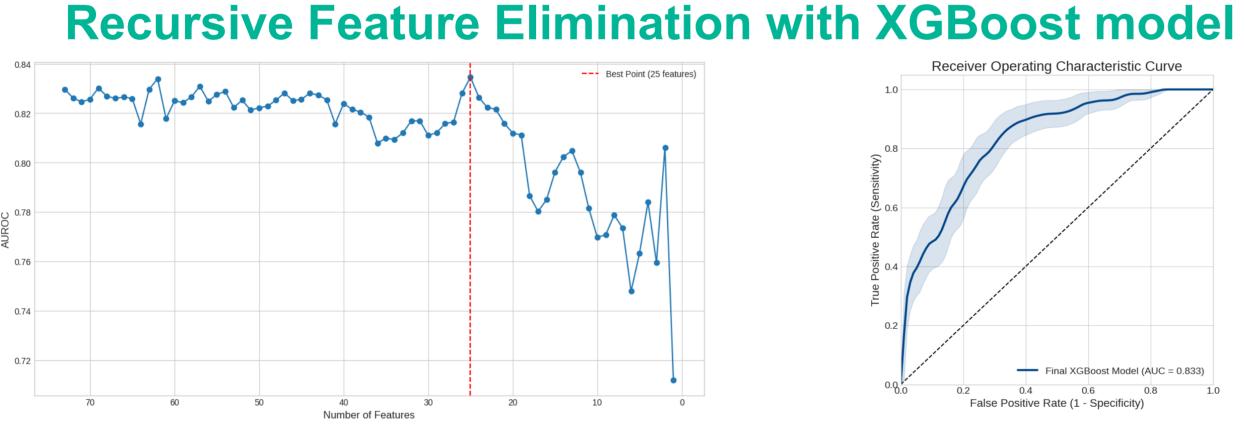
Recursive Feature Elimination: Select key features

Results - CLABSI

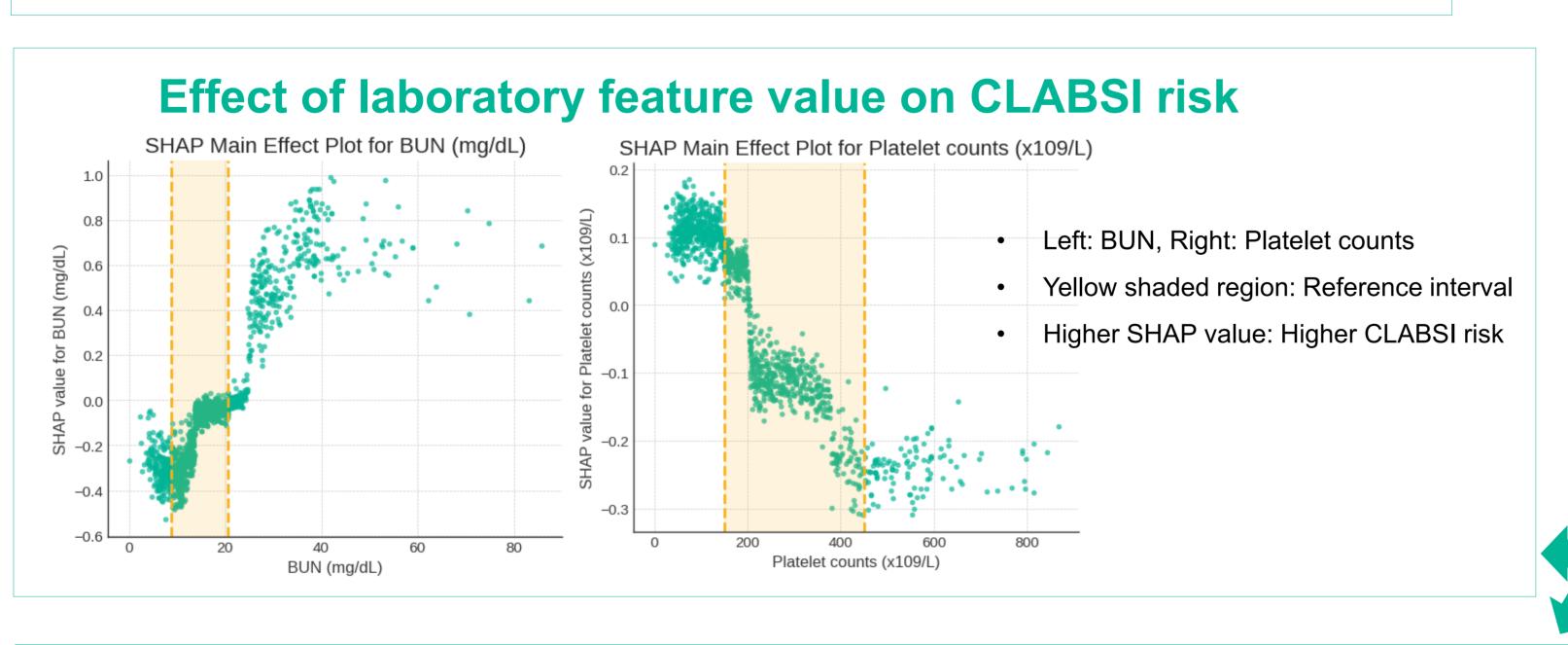
- 6,378 line-days from 860 CVCs in 286 patients were were analyzed
- CLABSI rate: 15.99 per 1,000 line-days
- **Most common** pathogens: *A.baumanii* (35.3%), *K.pneumoniae* (25.4%), and *Candida* spp. (17.6%)

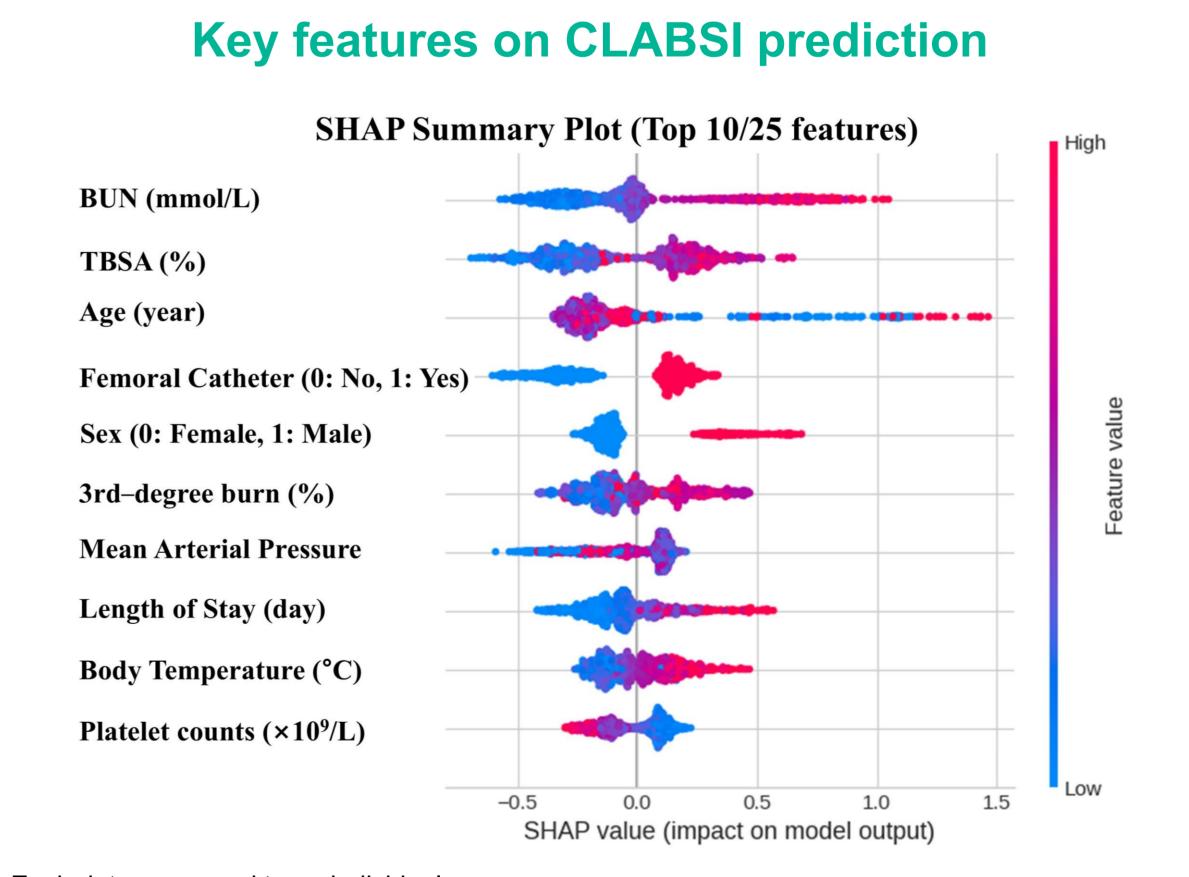
A. baumanii 35% Candida spp. K. pneumoniae

4 Results - CLABSI prediction ML model



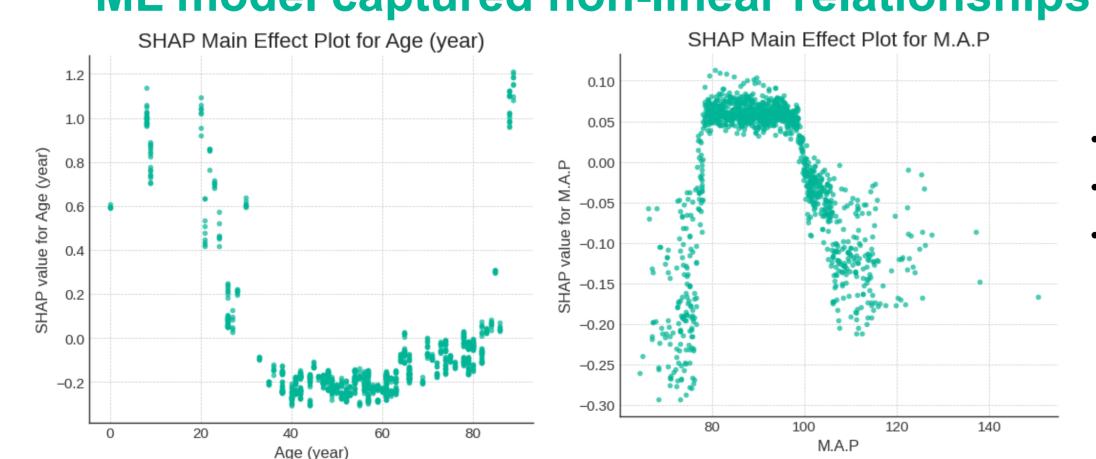
25 Features showed highest AUROC of 0.833 (95% CI: 0.798–0.868)





- Each dot correspond to an individual
- The colors represent feature values for numeric features: red for larger values, and blue for smaller
- · A negative SHAP value (extending to the left) indicates reduced CLABSI risk, while a positive one (extending to the right) indicates increased CLABSI risk

ML model captured non-linear relationships with CLABSI risk



- Left: Age, Right: mean arterial pressure
- Higher SHAP value: Higher CLABSI risk
- Age exhibited a U-shaped relationship and
 - M.A.P an inverted U-shaped relationship with CLABSI prediction





Conclusions

- ML model demonstrated excellent performance, and XAI revealed complex relationships between CLABSI risk and features
- Daily laboratory tests and vital signs were critical, showing that integrating time-series data with baseline characteristics enhances predictive accuracy