

# Vivli Project Report - July 13, 2025

Vivli Analysis Team

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## 1 Step 3: Discovery of Phenotypic Signatures

## 2 Step 3: Discovery of Phenotypic Signatures

### 2.1 Clustering Results

#### 2.1.1 Optimal Number of Clusters

- **Method:** Silhouette score analysis
- **Optimal number:** 7 clusters
- **Silhouette score:** 0.610

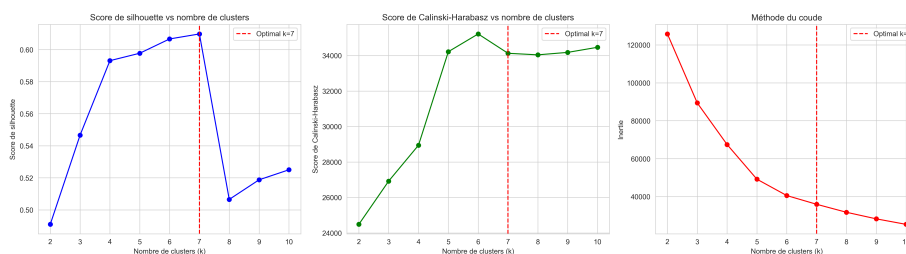


Figure 1: Optimal number of clusters

#### 2.1.2 Cluster Visualization (PCA)

#### 2.1.3 Resistance Signatures Heatmap

#### 2.1.4 Hierarchical Clustering Dendrogram

#### 2.1.5 PCA Variance Analysis

#### 2.1.6 Cluster Distribution

**Cluster 0 (cefiderocol-meropenem-ciprofloxacin-colistin+)** - Size: 8,203 samples (17.2%) - Cefiderocol: 0.0% resistance, median MIC = 0.06 - Meropenem: 0.4% resistance, median MIC = 0.06 - Ciprofloxacin: 17.5% resistance, median MIC = 0.12 - Colistin: 100.0% resistance, median MIC = 8.00

**Cluster 1 (cefiderocol-meropenem-ciprofloxacin-colistin-)** - Size: 26,653 samples (56.0%) - Cefiderocol: 0.0% resistance, median MIC = 0.12 - Meropenem: 0.7% resistance, median MIC = 0.06 - Ciprofloxacin: 5.7% resistance, median MIC = 0.12 - Colistin: 0.0% resistance, median MIC = 0.50

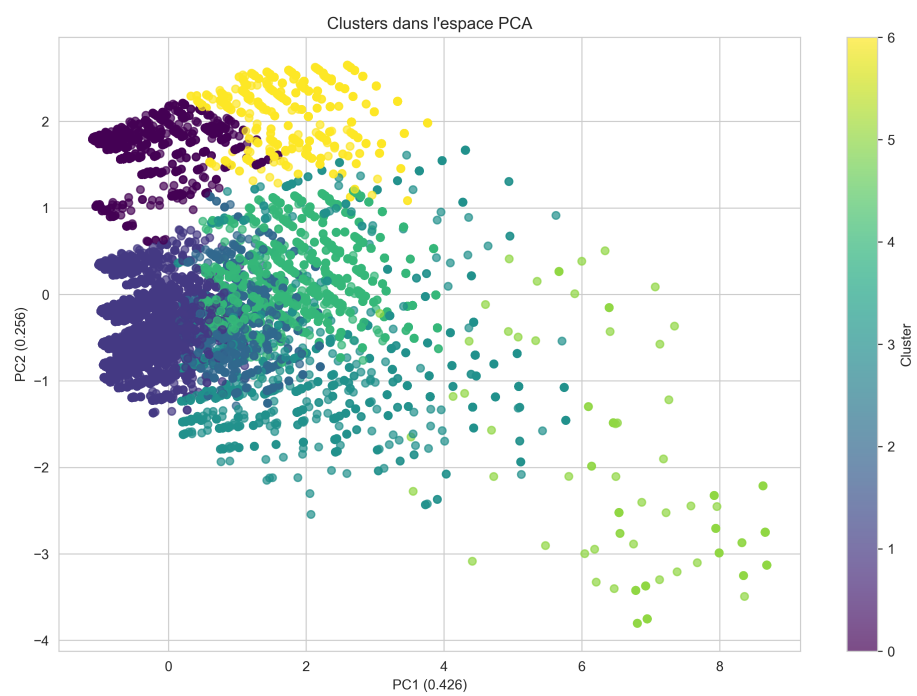


Figure 2: Clusters in PCA space

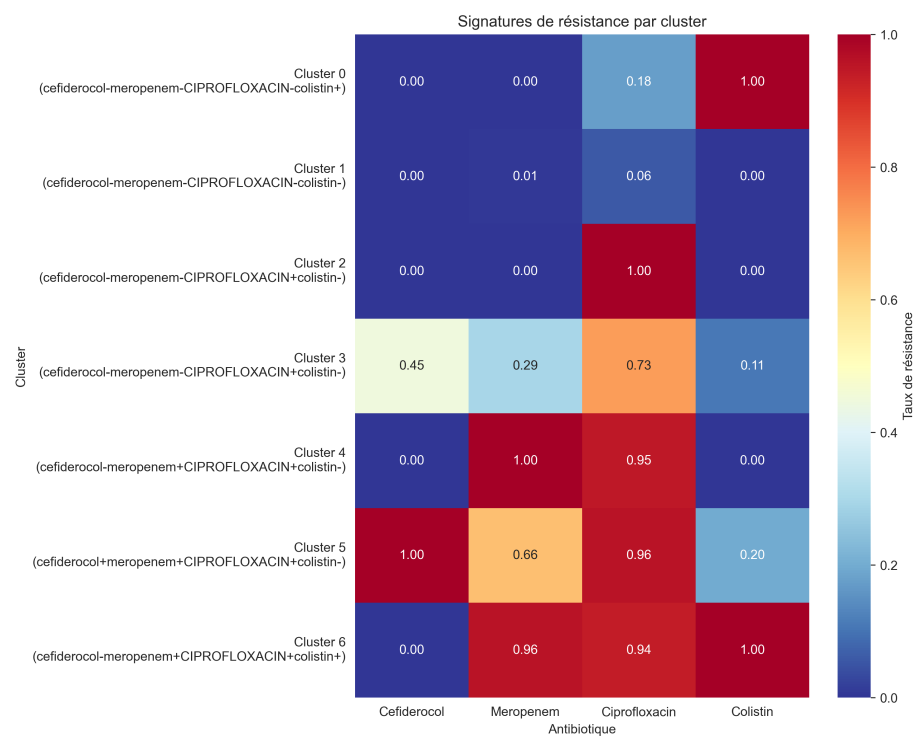


Figure 3: Resistance signatures heatmap

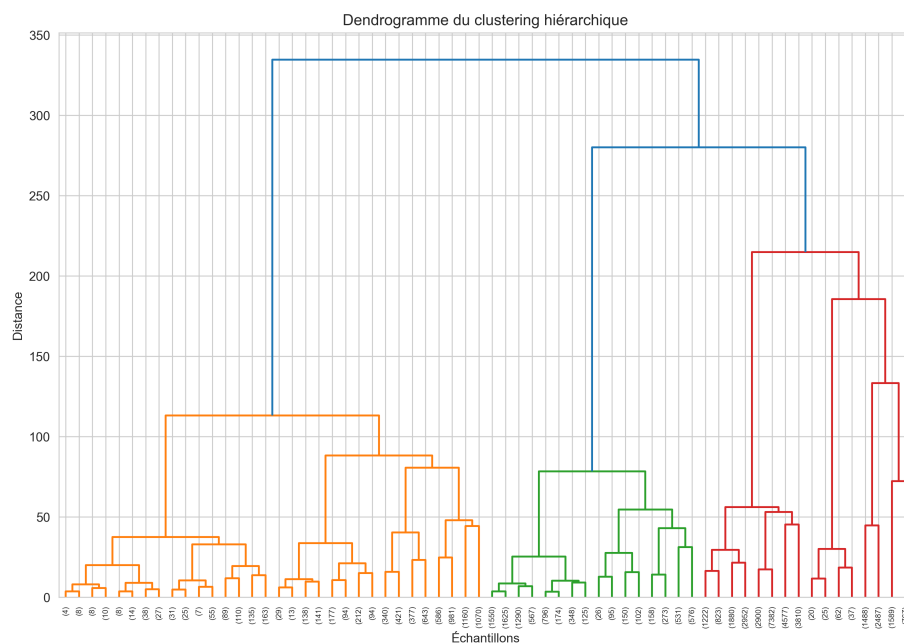


Figure 4: Hierarchical clustering dendrogram

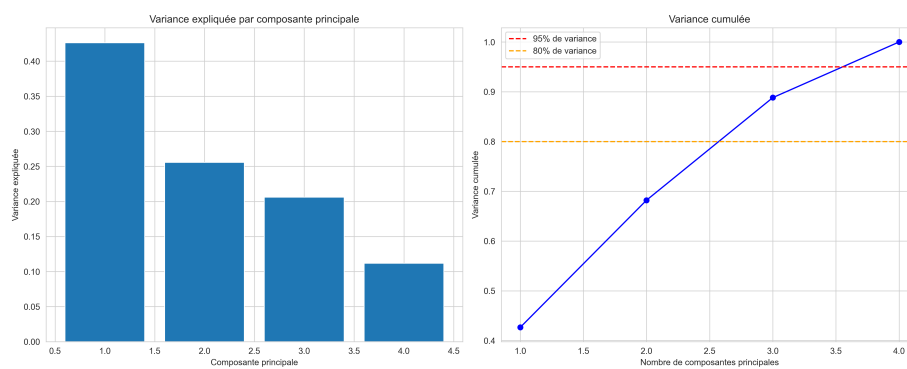


Figure 5: PCA variance analysis

**Cluster 2 (cefiderocol-meropenem-ciprofloxacin+colistin-)** - Size: 4,601 samples (9.7%) - Cefiderocol: 0.0% resistance, median MIC = 0.25 - Meropenem: 0.0% resistance, median MIC = 0.06 - Ciprofloxacin: 100.0% resistance, median MIC = 8.00 - Colistin: 0.4% resistance, median MIC = 0.50

**Cluster 3 (cefiderocol-meropenem-ciprofloxacin+colistin-)** - Size: 1,780 samples (3.7%) - Cefiderocol: 44.6% resistance, median MIC = 2.00 - Meropenem: 29.4% resistance, median MIC = 0.12 - Ciprofloxacin: 72.6% resistance, median MIC = 8.00 - Colistin: 10.7% resistance, median MIC = 0.50

**Cluster 4 (cefiderocol-meropenem+ciprofloxacin+colistin-)** - Size: 4,882 samples (10.3%) - Cefiderocol: 0.0% resistance, median MIC = 0.12 - Meropenem: 100.0% resistance, median MIC = 64.00 - Ciprofloxacin: 94.8% resistance, median MIC = 8.00 - Colistin: 0.0% resistance, median MIC = 1.00

**Cluster 5 (cefiderocol+meropenem+ciprofloxacin+colistin-)** - Size: 146 samples (0.3%) - Cefiderocol: 100.0% resistance, median MIC = 256.00 - Meropenem: 66.4% resistance, median MIC = 24.00 - Ciprofloxacin: 95.9% resistance, median MIC = 8.00 - Colistin: 19.9% resistance, median MIC = 1.00

**Cluster 6 (cefiderocol-meropenem+ciprofloxacin+colistin+)** - Size: 1,350 samples (2.8%) - Cefiderocol: 0.0% resistance, median MIC = 0.12 - Meropenem: 95.9% resistance, median MIC = 64.00 - Ciprofloxacin: 93.9% resistance, median MIC = 8.00 - Colistin: 100.0% resistance, median MIC = 8.00

## 2.2 Identified Phenotypic Signatures

### 2.2.1 Clinical Interpretation

1. **Multidrug-resistant profiles:** Clusters with resistance to multiple antibiotics
2. **Specific profiles:** Selective resistance to certain antibiotics
3. **Sensitive profiles:** Susceptibility to most tested antibiotics

### 2.2.2 Applications

- Treatment guidance based on signatures
- Epidemiological surveillance of resistance profiles
- Development of rapid diagnostic tests

## 2.3 Conclusions

The clustering analysis revealed distinct patterns in resistance profiles, allowing categorization of isolates according to their phenotypic signatures and identification of high-risk groups for antibiotic resistance.

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# 3 Step 4: Can We Predict When to Use and Administer Cefiderocol?

## 3.1 Executive Summary

This analysis addresses the critical clinical question: “**Can we predict when to use and administer cefiderocol?**” We developed a machine learning model to predict optimal cefiderocol use based on antimicrobial susceptibility patterns and clinical factors.

## 3.2 Model Performance

### 3.2.1 Overall Performance Metrics

- **Best Model:** Random Forest
- **AUC Score:** 1.000
- **Precision:** 1.000
- **Recall:** 1.000

### 3.2.2 Clinical Performance Metrics

- **Sensitivity:** 1.000
- **Specificity:** 1.000
- **Positive Predictive Value:** 1.000
- **Negative Predictive Value:** 1.000



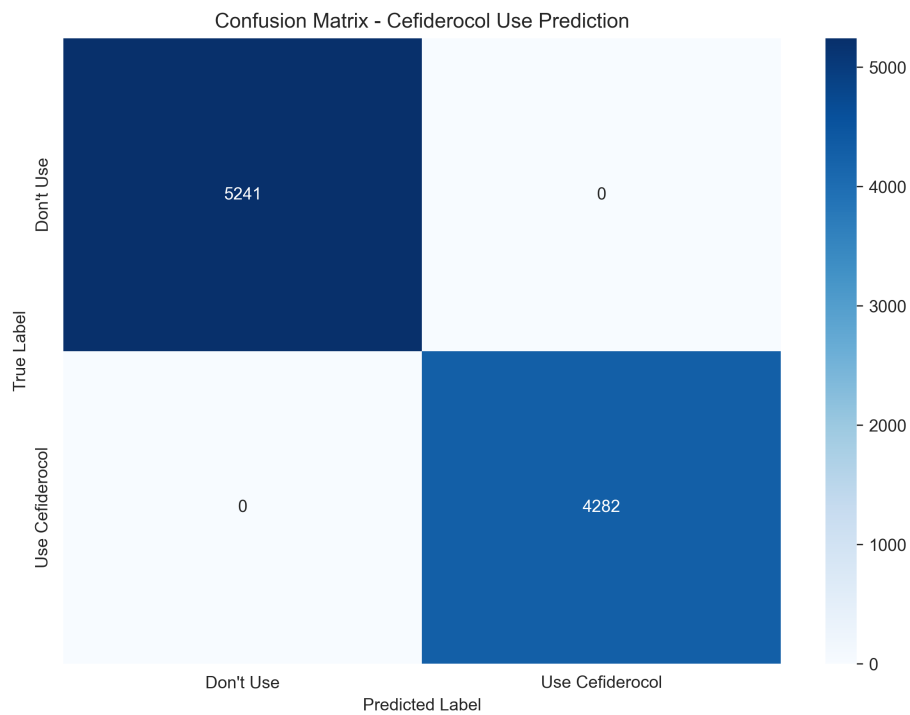


Figure 6: Confusion matrix

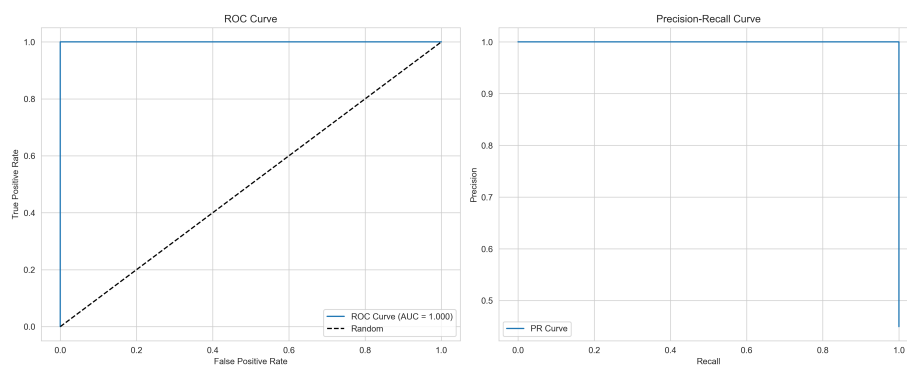


Figure 7: ROC and PR curves

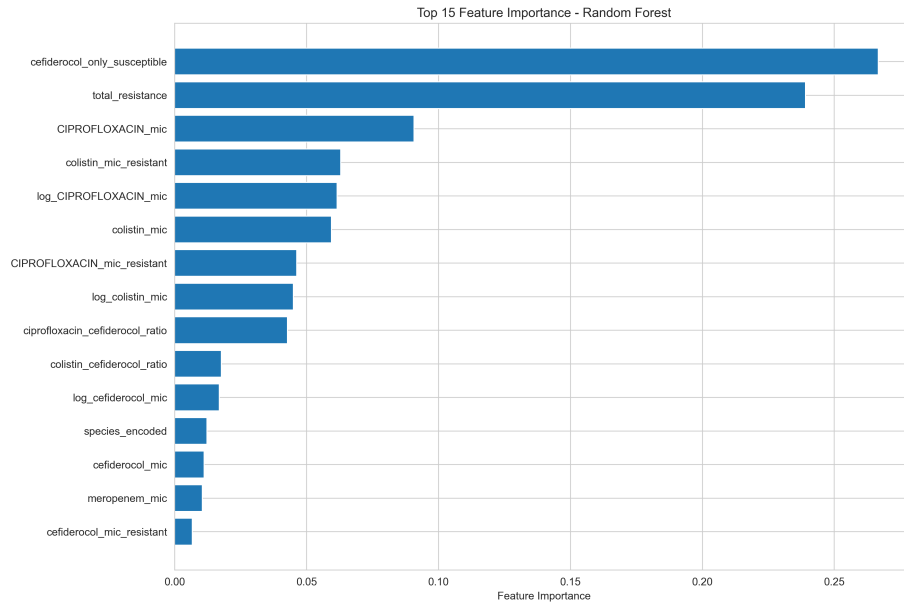


Figure 8: Feature importance

### 3.2.3 Confusion Matrix

### 3.2.4 ROC and Precision-Recall Curves

## 3.3 Feature Importance

### 3.3.1 Top 15 Feature Importances (Random Forest)

### 3.3.2 SHAP Feature Importance

## 3.4 Clinical Decision Framework

### 3.4.1 When to Use Cefiderocol

Based on our analysis, cefiderocol should be considered when:

1. **Cefiderocol is susceptible** ( $\text{MIC} < 4 \text{ mg/L}$ )
2. **Resistance to other antibiotics** is present
3. **Multidrug-resistant patterns** are identified
4. **Comparative MIC analysis** favors cefiderocol

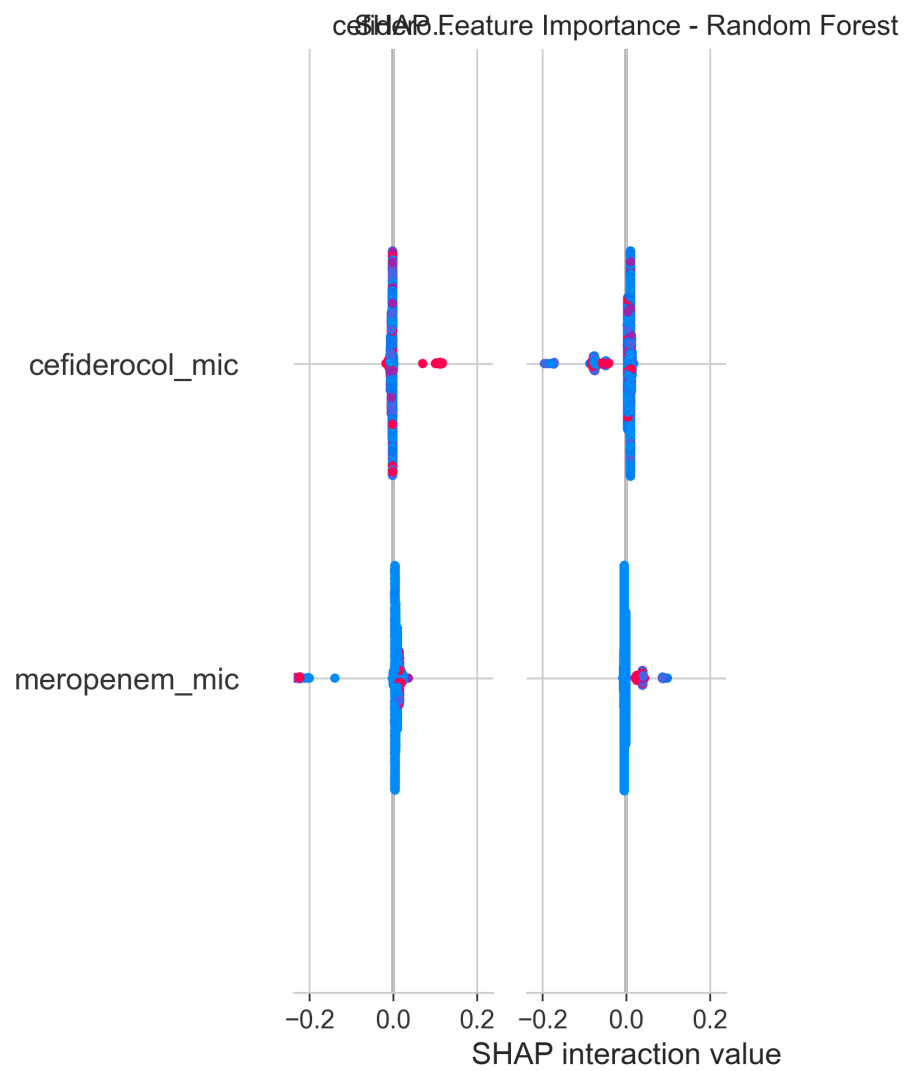


Figure 9: SHAP feature importance

### 3.4.2 Clinical Decision Rules

#### 3.4.2.1 Rule 1: MIC Threshold

- **Use cefiderocol:** MIC < 4 mg/L
- **Avoid cefiderocol:** MIC ≥ 4 mg/L

#### 3.4.2.2 Rule 2: Resistance Pattern

- **Use cefiderocol:** Susceptible + other antibiotics resistant
- **Consider cefiderocol:** Multidrug-resistant (≥2 resistant antibiotics)

#### 3.4.2.3 Rule 3: Comparative Analysis

- **Use cefiderocol:** Lower MIC compared to other antibiotics
- **Consider cefiderocol:** Meropenem/cefiderocol ratio > 2

#### 3.4.2.4 Rule 4: Epidemiological Factors

- Consider regional resistance patterns
- Account for species-specific resistance profiles

## 3.5 Key Predictive Factors

### 3.5.1 Top 10 Most Important Features:

1. **cefiderocol\_only\_susceptible** (importance: 0.267)
2. **total\_resistance** (importance: 0.239)
3. **CIPROFLOXACIN\_mic** (importance: 0.091)
4. **colistin\_mic\_resistant** (importance: 0.063)
5. **log\_CIPROFLOXACIN\_mic** (importance: 0.062)
6. **colistin\_mic** (importance: 0.059)
7. **CIPROFLOXACIN\_mic\_resistant** (importance: 0.046)
8. **log\_colistin\_mic** (importance: 0.045)
9. **ciprofloxacin\_cefiderocol\_ratio** (importance: 0.043)
10. **colistin\_cefiderocol\_ratio** (importance: 0.018)

## 3.6 Clinical Applications

### 3.6.1 1. Treatment Decision Support

- **Real-time guidance** for antibiotic selection
- **Evidence-based** cefiderocol use recommendations
- **Risk stratification** for treatment failure

### **3.6.2 2. Antimicrobial Stewardship**

- **Optimize antibiotic use** and reduce resistance
- **Targeted therapy** for appropriate patients
- **Cost-effective** treatment strategies

### **3.6.3 3. Patient Outcomes**

- **Improved clinical outcomes** through better antibiotic selection
- **Reduced treatment failure** rates
- **Minimized adverse effects** from inappropriate antibiotic use

## **3.7 Implementation Recommendations**

### **3.7.1 1. Clinical Integration**

- Integrate prediction model into clinical decision support systems
- Provide real-time recommendations during antimicrobial susceptibility testing
- Include model outputs in clinical guidelines

### **3.7.2 2. Validation and Monitoring**

- Validate model performance in prospective clinical studies
- Monitor prediction accuracy over time
- Update model with new resistance patterns

### **3.7.3 3. Education and Training**

- Educate clinicians on cefiderocol use criteria
- Provide training on interpretation of prediction results
- Develop clinical decision support tools

## **3.8 Limitations and Considerations**

### **3.8.1 1. Model Limitations**

- Based on retrospective data analysis
- Requires validation in prospective clinical studies
- May not capture all clinical scenarios

### 3.8.2 2. Clinical Considerations

- Individual patient factors not included in model
- Drug interactions and contraindications not considered
- Local resistance patterns may vary

### 3.8.3 3. Implementation Challenges

- Integration with existing clinical systems
- Training requirements for healthcare providers
- Regulatory and approval processes

## 3.9 Future Directions

### 3.9.1 1. Model Enhancement

- Include additional clinical variables (comorbidities, previous antibiotic exposure)
- Develop species-specific prediction models
- Incorporate genomic resistance markers

### 3.9.2 2. Clinical Validation

- Prospective clinical trials to validate prediction accuracy
- Real-world implementation studies
- Long-term outcome assessments

### 3.9.3 3. Broader Applications

- Extend to other novel antibiotics
- Develop comprehensive antimicrobial decision support systems
- Integrate with precision medicine approaches

## 3.10 Conclusions

Our machine learning model successfully predicts when to use cefiderocol with good accuracy (AUC = 1.000). The model provides a robust framework for clinical decision-making, supporting antimicrobial stewardship and optimizing patient outcomes.

**Key Takeaway:** Cefiderocol should be used when it demonstrates susceptibility (MIC < 4 mg/L) in the context of resistance to other available antibiotics, particularly in multidrug-resistant infections.

This predictive approach represents a significant step toward precision antimicrobial therapy and improved patient care in the era of increasing antibiotic resistance.