Summary of preprocessing selection (MAE ± 95% CI)
Method: COMBINATORIC | Averaging: Weighted

Preprocessing (y-axis) categories from top to bottom:
- RandomForestImputer_MaxAbsScaler
- RandomForestImputer_RobustScaler
- RandomForestImputer_MinMaxScaler
- RandomForestImputer_StandardScaler
- KNNImputer_MaxAbsScaler
- KNNImputer_RobustScaler
- KNNImputer_MinMaxScaler
- KNNImputer_StandardScaler
- MedianImputer_MaxAbsScaler
- MedianImputer_RobustScaler
- MedianImputer_MinMaxScaler
- MedianImputer_StandardScaler
- MeanImputer_MaxAbsScaler
- MeanImputer_RobustScaler
- MeanImputer_MinMaxScaler
- MeanImputer_StandardScaler

MAE (Lower is better) — x-axis values: 0.125, 0.150, 0.175, 0.200, 0.225, 0.250

Evaluation type
- MAE : 0.158
- Overall
- Bacillus
- Pathogen
- Interaction