



# Heart Failure Prediction

Super Learner

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# Project

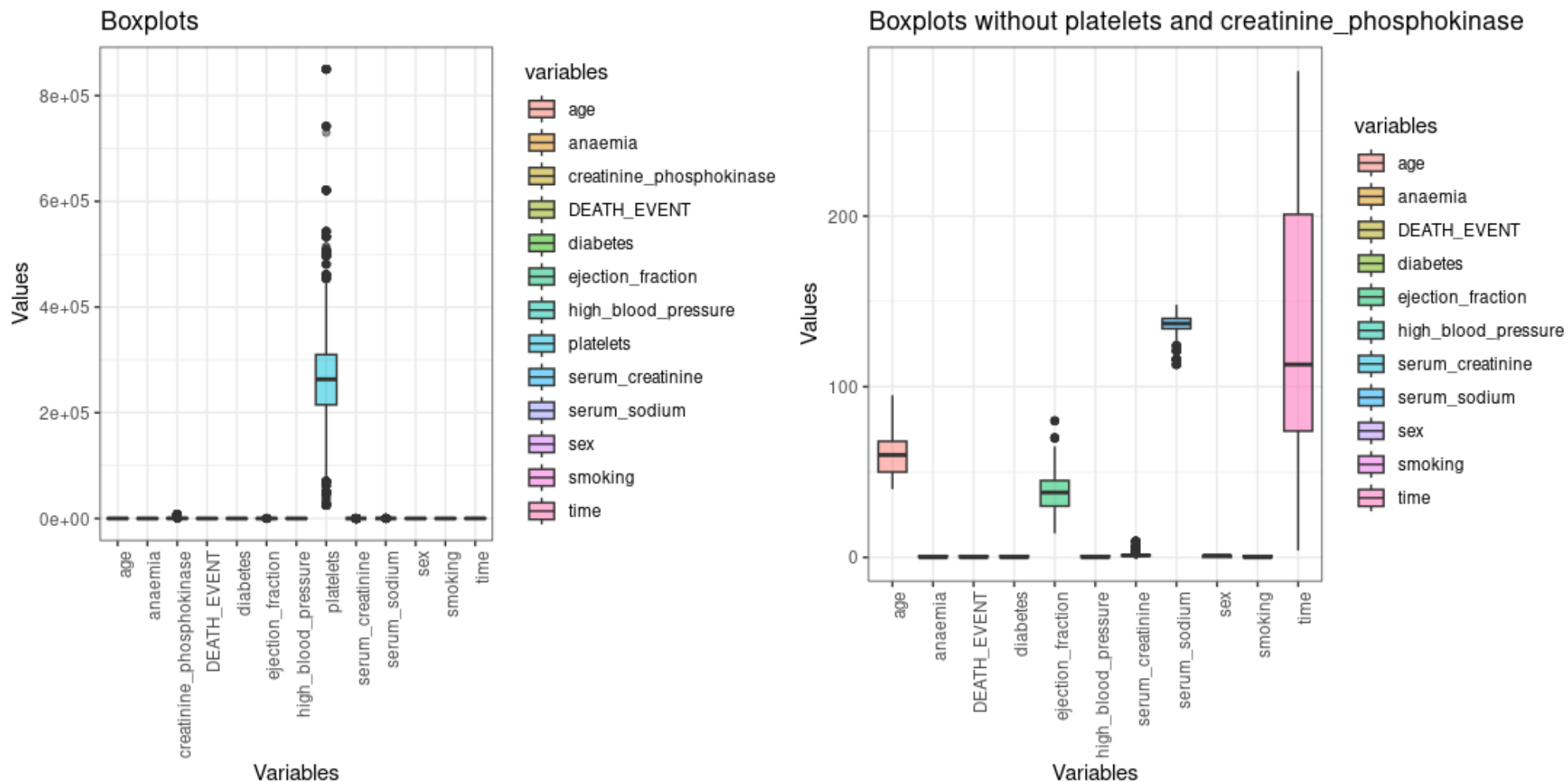
## Overview

### 1. Objective:

Training a super learner model while leveraging the strenghts of multiple predictive algorithms, achieving greater accuracy and a lower RMSE value.

### 2. Predictive models:

- **RandomForest** and **Ranger**
- **XGBoost**
- **GLMNet** and **Bayesian GLM**
- **SVM**
- **Neural Networks**
- **Polymars** (Piecewise-Polynomial, splines based regression)



Boxplots representations of all the variables. Highlighting the presence of some outliers.

## 10-fold cross-validation (parallel)

```
# cross-validation  
cv_control <- SuperLearner.CV.control(V = 10) # 10-fold cross-validation  
  
# Setting up parallel processing  
cluster <- makeCluster(detectCores() - 1) # use all cores except one  
registerDoParallel(cluster)  
registerDoRNG(seed = 123) # Ensure reproducibility
```

## Model training with 70% of the dataset

```
# Executing the cross-validation phases in parallel thanks to cv_control  
clusterExport(cluster, c("Y_train", "X_train", "learners", "cv_control"))  
super_learner <- SuperLearner(Y = Y_train, X = X_train, family = binomial(),  
                             SL.library = learners, method = "method.NNLS",  
                             cvControl = cv_control, verbose = TRUE)
```



# ROC and AUC

## Evaluation

### ROC (Receiver Operating Characteristic)

- Shows the performance of a binary classification model at different threshold levels
- Trade-Off between specificity and sensitivity

**Sensitivity:** proportion of the true positives correctly identified by the model

$$\text{sensitivity} = \frac{\text{True positives (TP)}}{\text{True positives (TP)} + \text{False negatives (FN)}}$$

**Specificity:** proportion of the true negatives correctly identified by the model

$$\text{specificity} = \frac{\text{True negatives (TN)}}{\text{True negatives (TN)} + \text{False positives (FP)}}$$

# ROC and AUC

## Evaluation

### **AUC (Area Under The Curve):**

- It's the area under the ROC curve
- Summarises the performance of the model and its in the range  $[0,1]$

### AUC = 1:

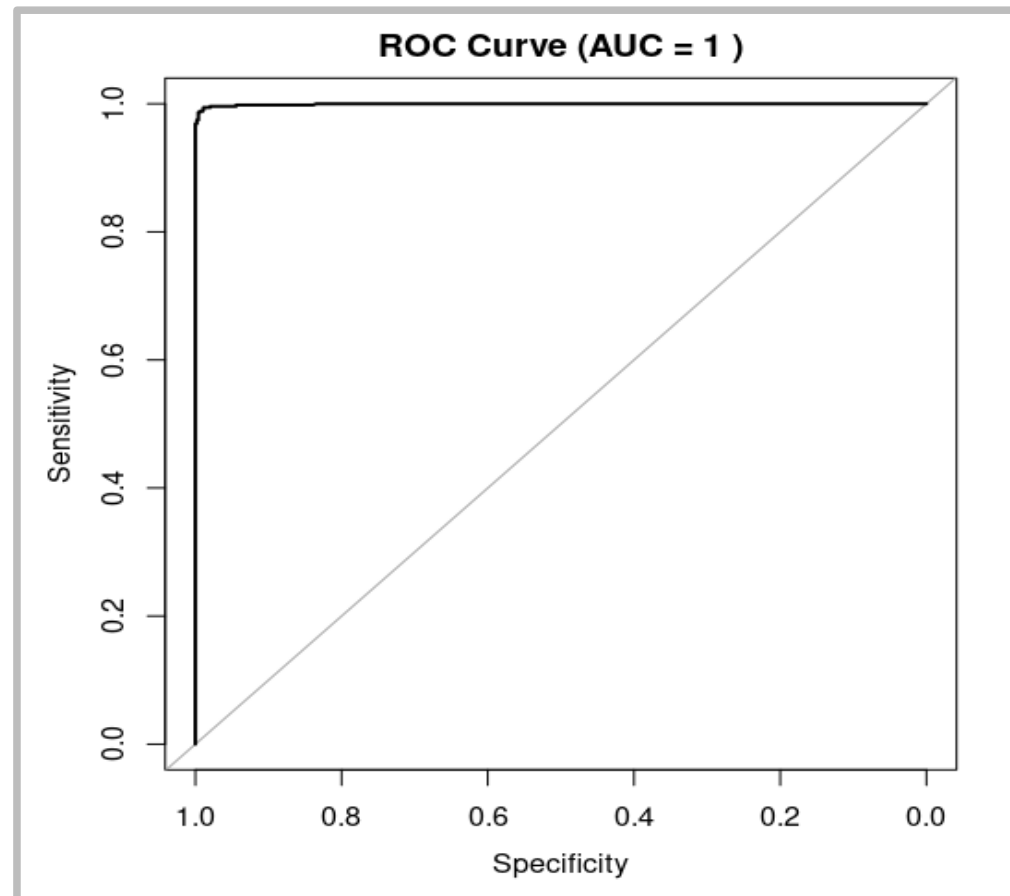
The model is able to correctly distinguish between positive and negative classes

### AUC = 0.5:

The model is unable to distinguish the classes

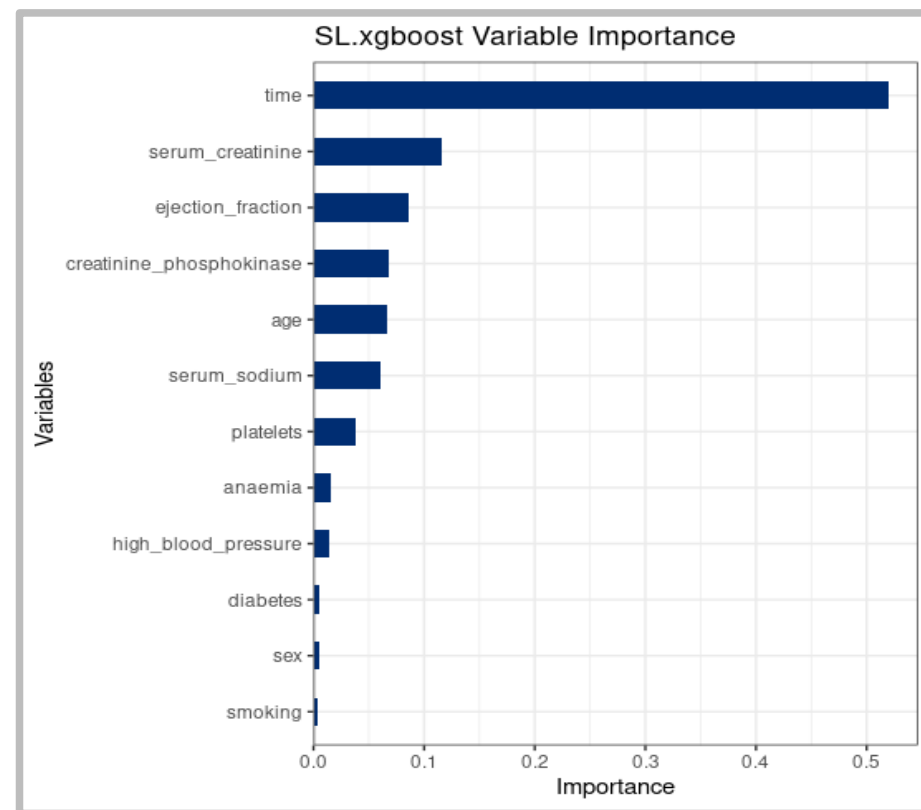
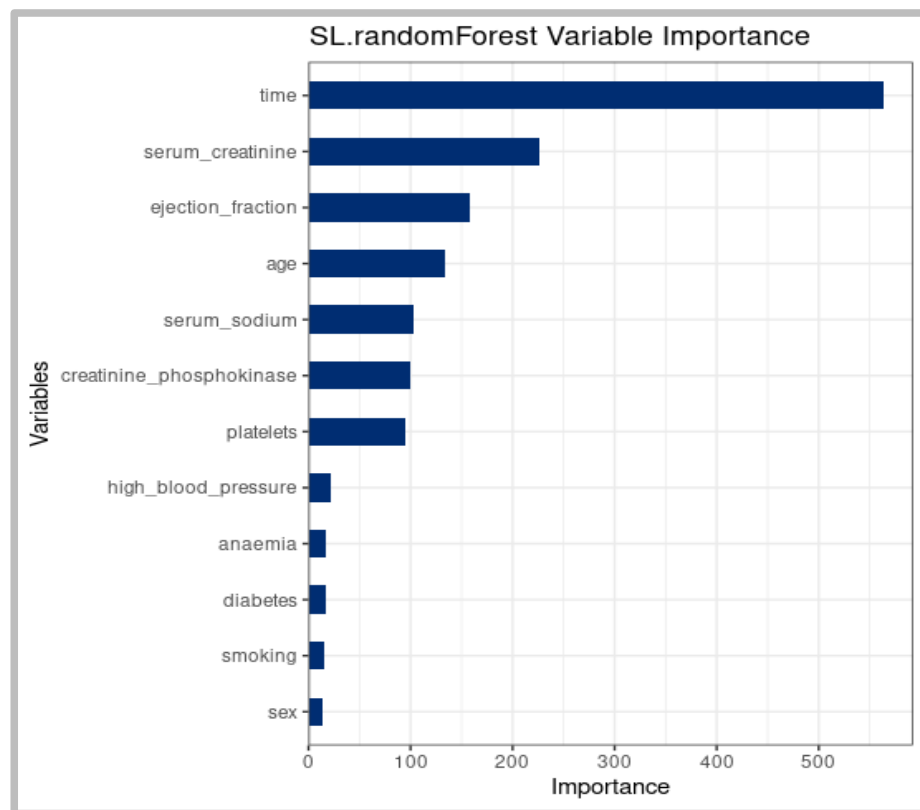
### AUC < 0.5:

The model performs worse than a casual classification



The graph shows that the model is able to correctly identify true positives and avoid false positives.

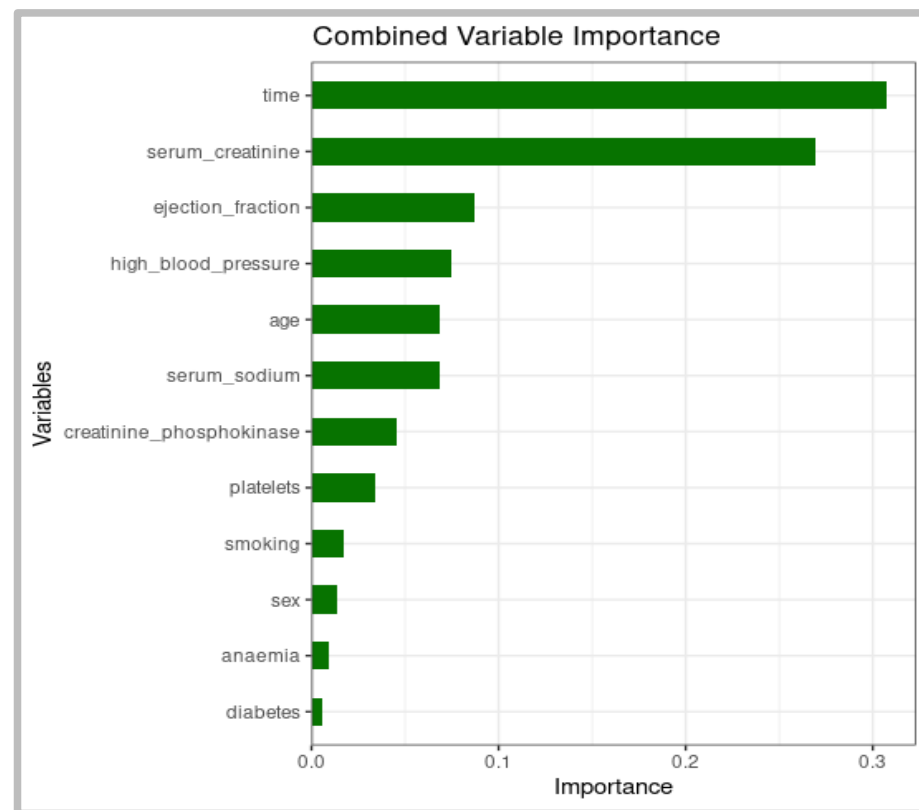
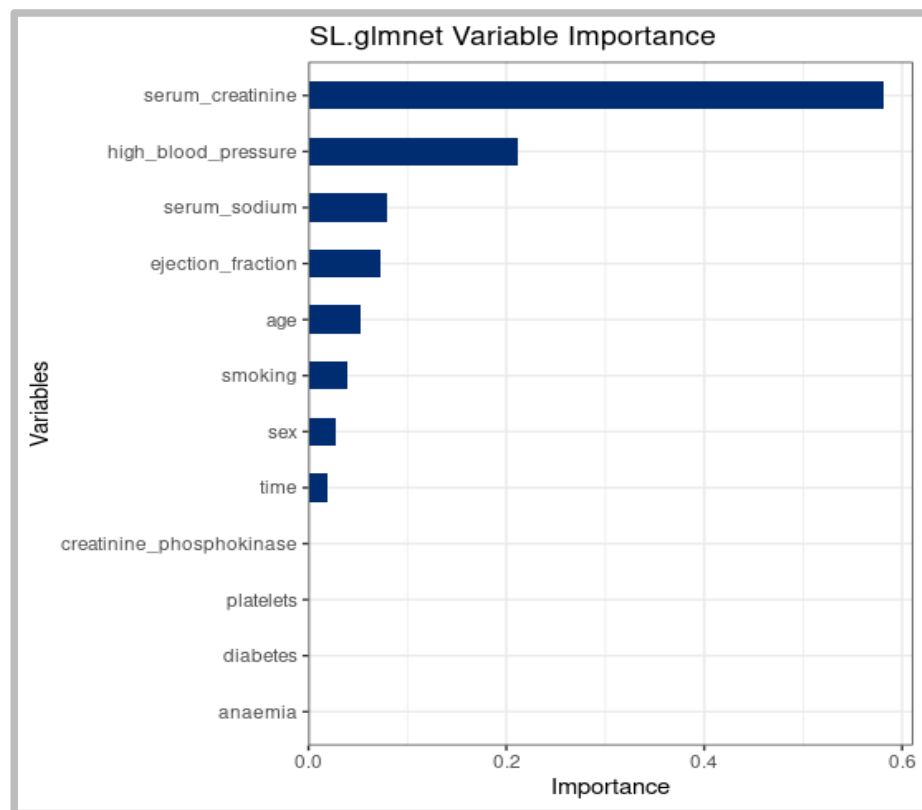
## Variable importance



Graphs showing the variable importance identified by the Random Forest and the XGBoost models



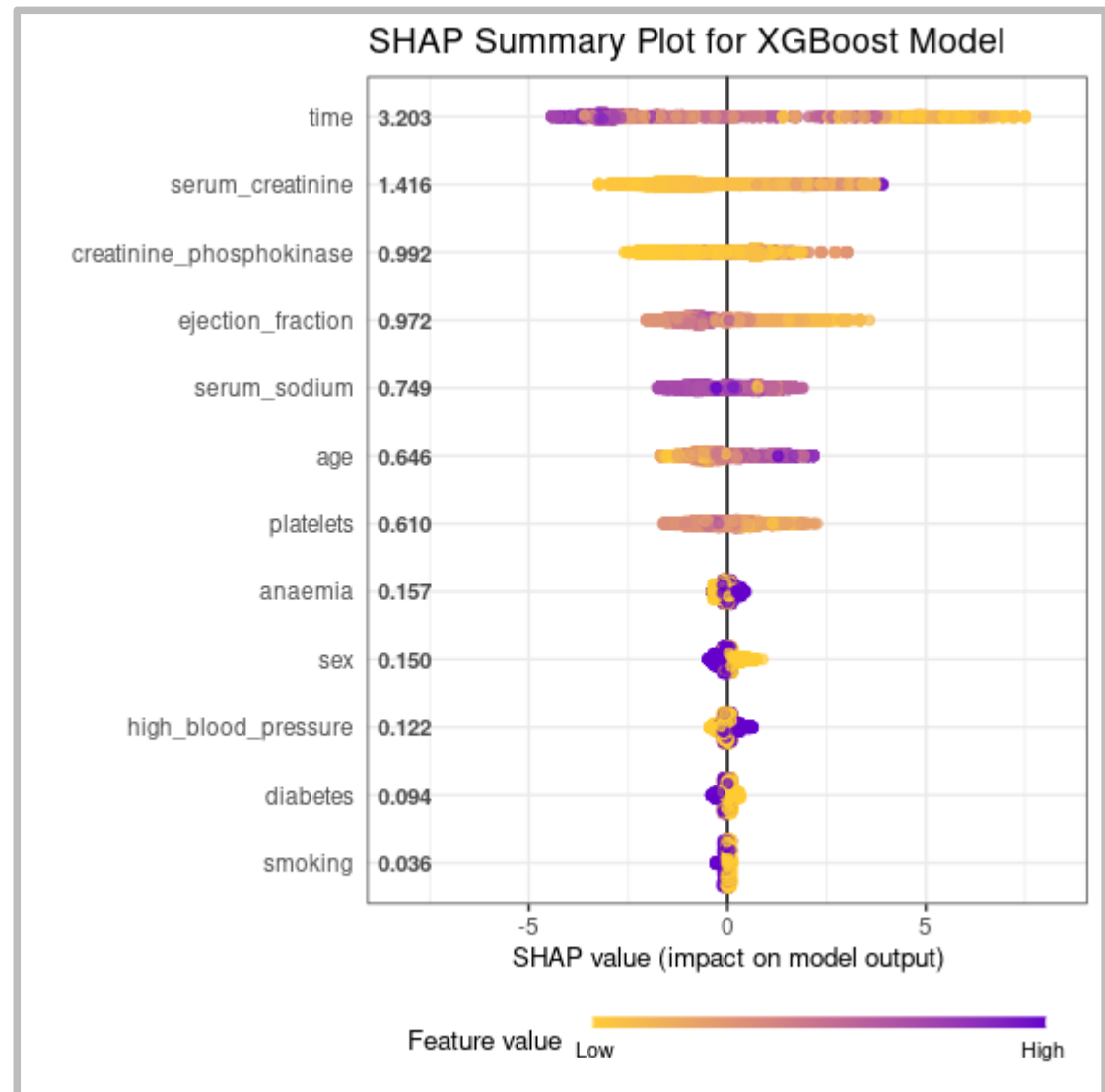
## Variable importance



Graphs showing the variable importance of the GLMNet model alongside the average variable importance of all the models combined.

## SHapley Additive exPlanations:

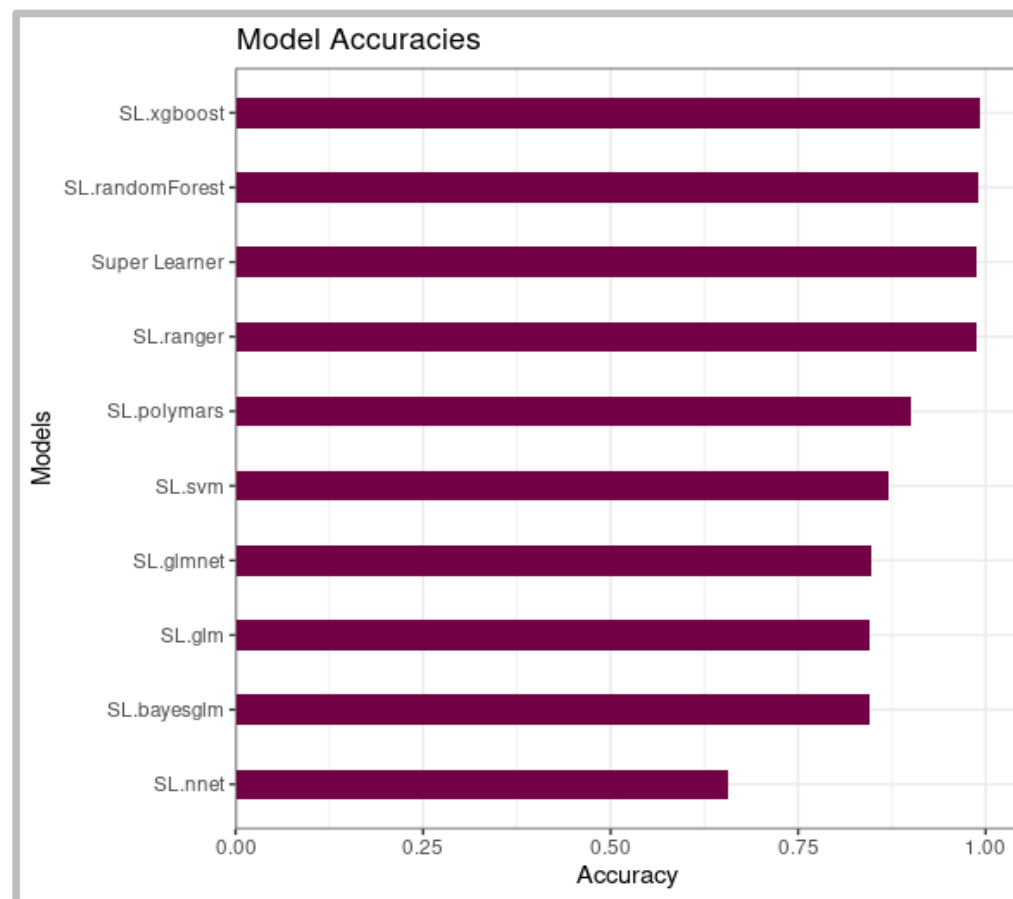
- Help identifying the most important features and their effects on the predictions by quantifying their impact
- Show how and how much the feature shifts the prediction in relation to the average



Descriptive SHAP summary showing the impact of each feature over the target variable

## Performance evaluation

Model	Accuracy	RMSE
SL.xgboost	0.9927	0.0845
SL.randomForest	0.9893	0.0877
SL.ranger	0.9887	0.0866
Super Learner	0.9887	0.0848
SL.polymars	0.8993	0.2688
SL.svm	0.8713	0.2986
SL.glmnet	0.8480	0.3467
SL.glm	0.8460	0.3474
SL.bayesglm	0.8460	0.3474
SL.nnet	0.6573	0.4764



Descriptive bar plot showing the level of accuracy for each base model, compared to the SuperLearner

## Resources

Dataset:

<https://www.kaggle.com/datasets/aadarshvelu/heart-failure-prediction-clinical-records>

Project source code:

[https://github.com/Scrayil/Heart\\_Failure\\_Prediction](https://github.com/Scrayil/Heart_Failure_Prediction)