

# Machine Learning Applied to Hepatocellular Carcinoma

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## Section 1

### Background HCC

# HepatoCellular Carcinoma (HCC)

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- Data mining approach to tailor evaluation and treatment for HCC are limited in the literature.
- Using the HCC dataset, we undertook the data mining approach to evaluate the patient level factors to identify those who are at risk of one year mortality.

## Section 2

### Data Summary

# Data Summary

- Clinical data of 165 pts with HCC (demographic, risk factors, lab data, and survival features)
- 49 features from HCC clinical practice guidelines (Table 1)
- About 80
- Missing data represents 10.22
- Missing data were imputed using various methods, then `step_nzv` was used to remove the variables that have near zero variability.
- The target variable is the survival at 1 year, coded as 0 (dies) and 1 (lives).

# Data Summary Table

Table 1: HCC Data Summary of Selected Variables

Overall (N=165)	
<b>gender</b>	
female	32 (19.4%)
male	133 (80.6%)
<b>symptom</b>	
no	53 (36.1%)
yes	94 (63.9%)
N-Miss	18
<b>alc</b>	
no	43 (26.1%)
yes	122 (73.9%)



## Section 3

### Random Forest Model

## Description

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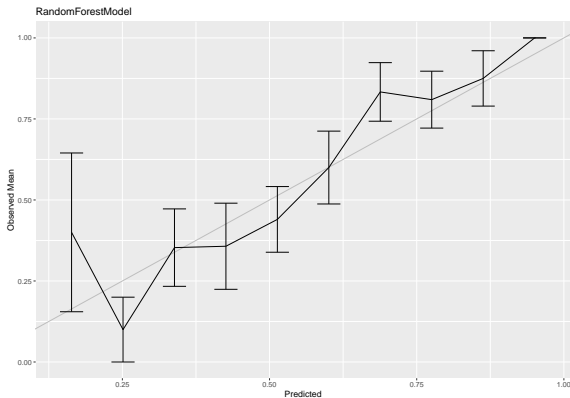
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- 10 fold CV
- Tuned Hyperparameter Parameters with Bayesian Optimization
- Applied and tuned KNN imputation and Correlation Filter

## Estimated Performance

Table 2: RF results with knn imputation and corr filter

Metric	Mean	Median	SD	Min	Max	NA
Brier	0.175	0.175	0.046	0.115	0.245	0
Accuracy	0.720	0.719	0.103	0.562	0.875	0
Kappa	0.390	0.383	0.220	0.143	0.714	0
ROC AUC	0.799	0.808	0.137	0.608	0.983	0
Sensitivity	0.824	0.800	0.146	0.500	1.000	0
Specificity	0.555	0.619	0.184	0.333	0.833	0

# Calibration Plots



Background HCC  
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Data Summary  
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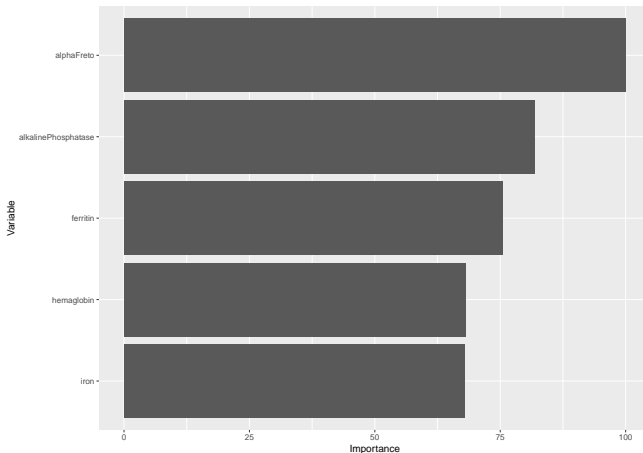
Random Forest Model  
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XGBoost Model  
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Support Vector Machine Model  
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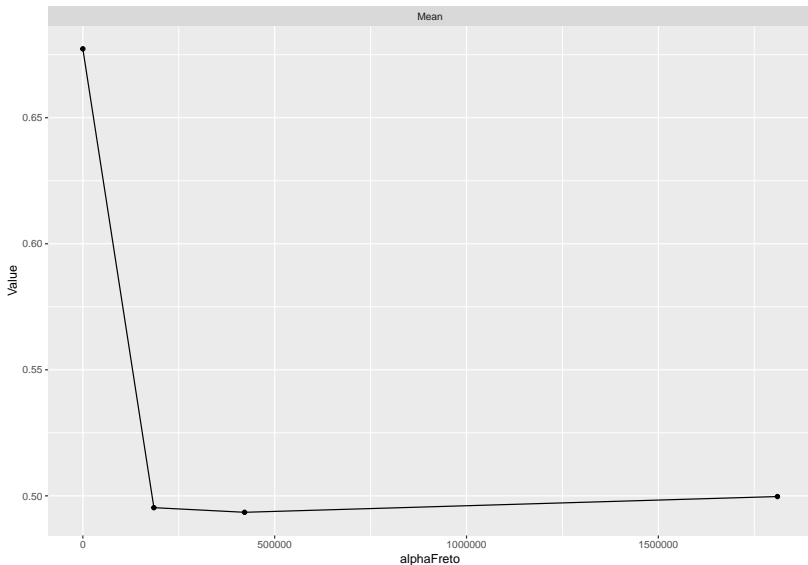
Final Model choice  
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# Variable Importance





# Partial Dependence



## Section 4

### XGBoost Model

# Description

Input: training set  $\{(x_i, y_i)\}_{i=1}^N$ , a differentiable loss function  $L(y, F(x))$ , a number of weak learners  $M$  and a learning rate  $\alpha$ .

- 1 Initialize model with a constant value:

$$\hat{f}_{(0)}(x) = \operatorname{argmin}_{\theta} \sum_{i=1}^N L(y_i, \theta).$$

- 2 For  $m = 1, \dots, M$ :

- a Compute gradients and Hessians:  $\hat{g}_m(x_i) = \left[ \frac{\partial L(y_i, f(x_i))}{\partial f(x_i)} \right]_{f(x)=\hat{f}_{m-1}(x)}$

$$\hat{h}_m(x_i) = \left[ \frac{\partial^2 L(y_i, f(x_i))}{\partial f(x_i)^2} \right]_{f(x)=\hat{f}_{m-1}(x)}.$$

- b Fit a base learner using the training set  $\left\{ x_i, -\frac{\hat{g}_m(x_i)}{\hat{h}_m(x_i)} \right\}_{i=1}^N$  by solving the optimization problem below:

$$\hat{\phi}_m = \operatorname{argmin}_{\phi \in \Phi} \sum_{i=1}^N \frac{1}{2} \hat{h}_m(x_i) \left[ -\frac{\hat{g}_m(x_i)}{\hat{h}_m(x_i)} - \phi(x_i) \right]^2.$$

- c Update the model:

$$\hat{f}_m(x) = \hat{f}_{(m-1)}(x) + \alpha \hat{\phi}_m.$$

- 3 Output  $\hat{f}(x) = \hat{f}_{(M)}(x)$ .

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- Tuning parameters: number of boosting iterations  $M$ , shrinkage of variable weights at each iteration to prevent overfitting  $\eta$ , maximum tree depth.
- Tuning inputs: KNN with the number of neighbors, correlation among variables, principal component analysis proportion of variance to be retained.

## Estimated Performance

Table 3: XGBoost results with knn imputation and feature selection using correlation

Metric	Mean	Median	SD	Min	Max	NA
Brier	0.221	0.219	0.086	0.081	0.344	0
Accuracy	0.701	0.688	0.096	0.562	0.889	0
Kappa	0.360	0.353	0.210	0.097	0.766	0
ROC AUC	0.768	0.775	0.116	0.584	0.961	0
Sensitivity	0.774	0.800	0.116	0.600	0.909	0
Specificity	0.586	0.619	0.187	0.333	0.857	0

- Tried six models with different combinations of imputation methods and feature selection methods. All models are not doing well for specificity.

## Estimated Performance

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- KNN\_corr appears to do better among models for its highest accuracy and kappa, second highest sensitivity and specificity, reasonable brier and roc auc.



## Estimated Performance

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- P-values are not significant.

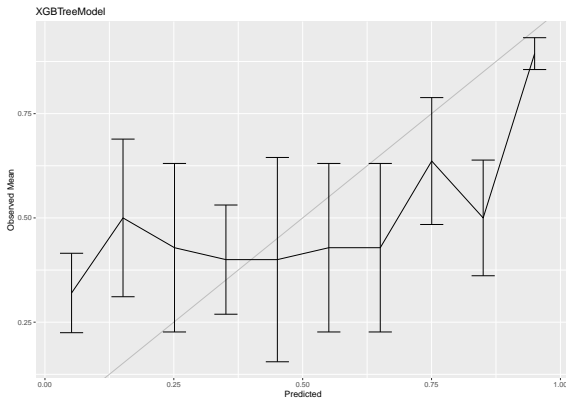
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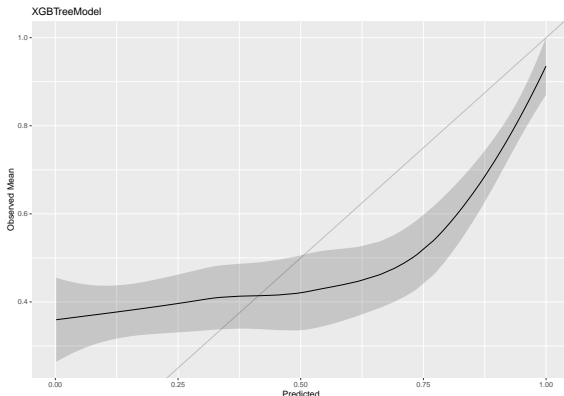
- P-values are not significant.
- Bayesian optimization selected  $\eta = 0.1$ , maximum tree depth = 7, number of boosting iterations = 106, number of features = 44.

# Calibration Plot



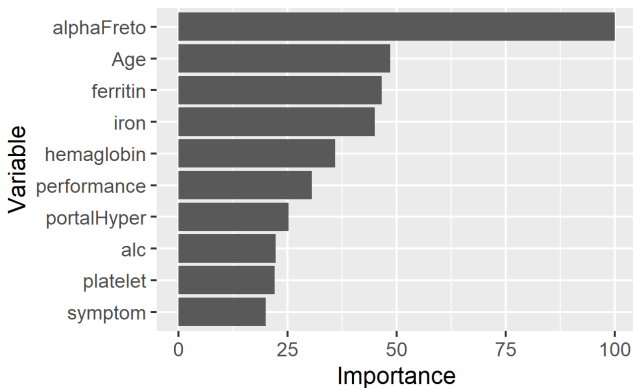
- Some fluctuations.

# Smoothed Calibration Plot

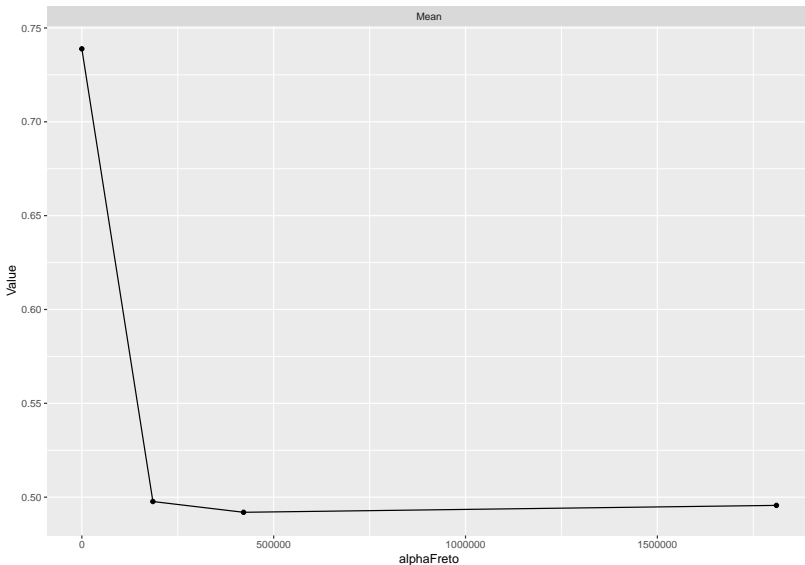


- Smoothed calibration curve shows false negatives on the lower left part and some false positives on the upper right part. This is consistent to the low specificity and relatively high sensitivity of the model.

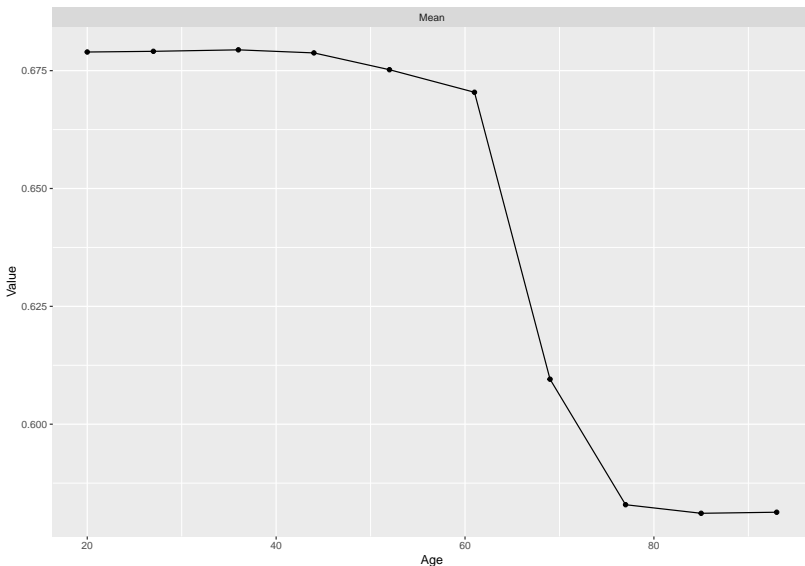
# Variable Importance



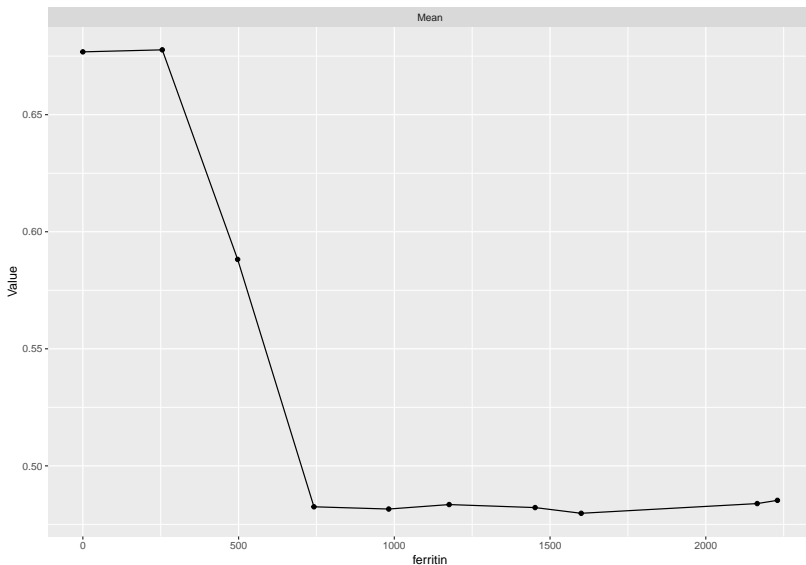
# Partial Dependence: alphaFreto



# Partial Dependence: Age

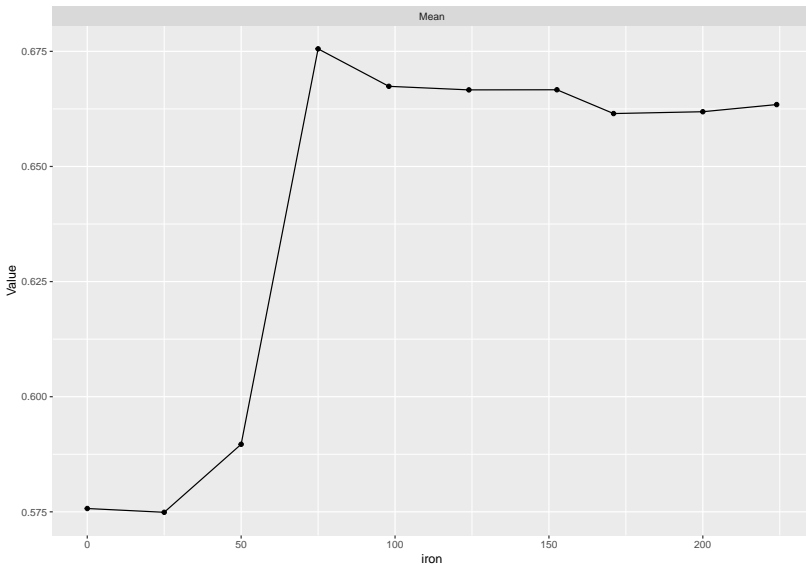


# Partial Dependence: ferritin

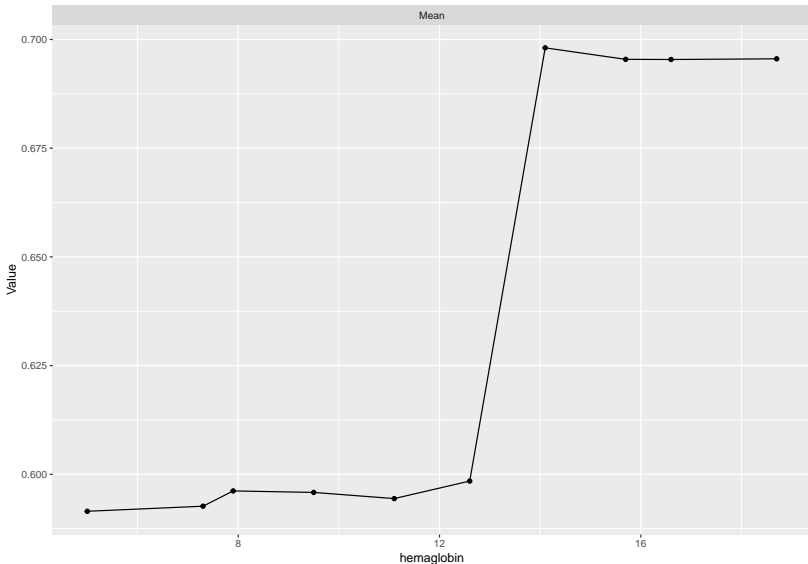




## Partial Dependence: iron



## Partial Dependence: hemaglobin



## Section 5

# Support Vector Machine Model

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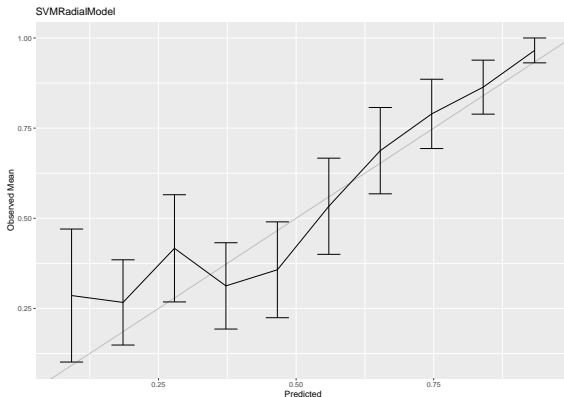
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- Cost of misclassification  $C = 0.8956493$
- imputation using KNN with neighbors = 4

## Estimated Performance

Table 6: SVMRad results with knn imputation

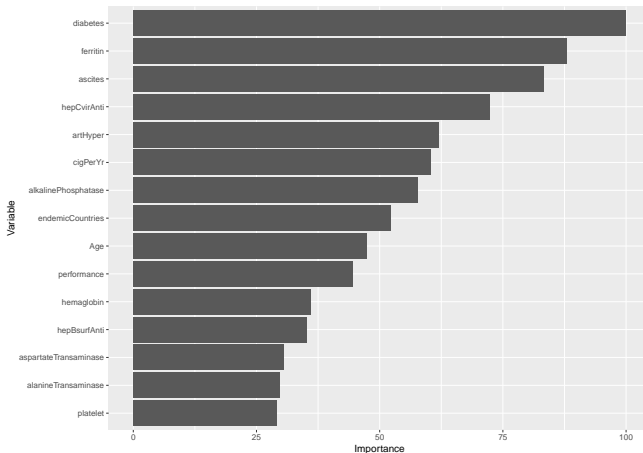
Metric	Mean	Median	SD	Min	Max	NA
Brier	0.176	0.166	0.057	0.101	0.287	0
Accuracy	0.756	0.764	0.119	0.562	0.938	0
Kappa	0.482	0.503	0.254	0.097	0.871	0
ROC AUC	0.818	0.829	0.101	0.600	0.967	0
Sensitivity	0.803	0.809	0.107	0.600	0.909	0
Specificity	0.681	0.643	0.190	0.500	1.000	0

# Calibration Plots



- Decently calibrated. Low probabilities have many false negatives.

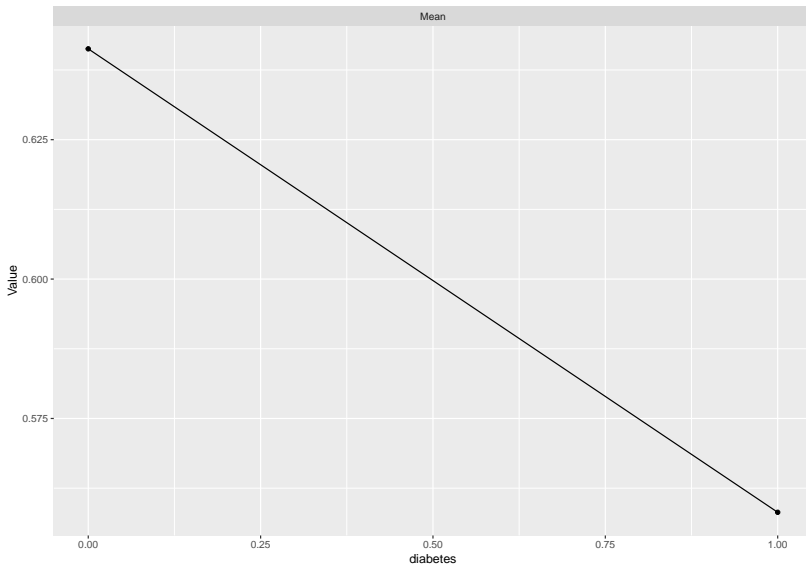
# Variable Importance



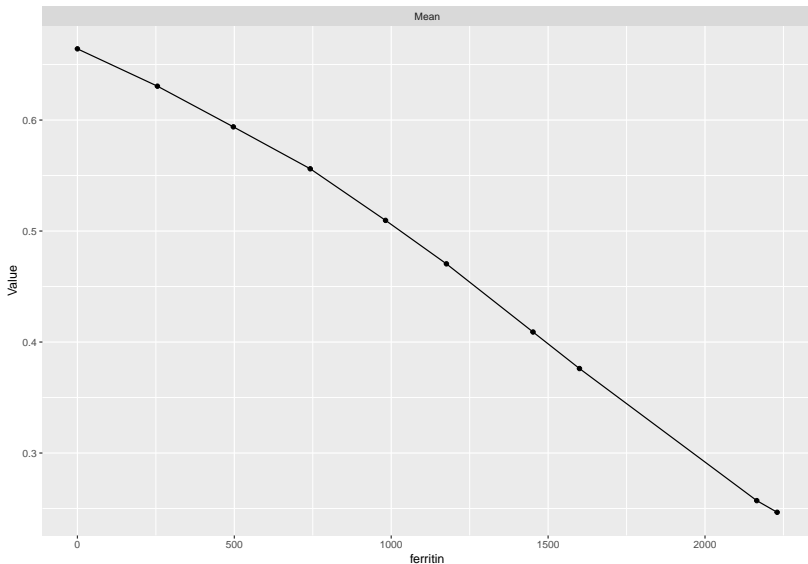
# Variable Importance

- Symptoms (performance)
- Indicators of liver injury/disease/infection (hepBsurfAnti, hepCvirAnti, aspartateTransaminase, alkalinePhosphatase, ferritin, totalProteins, ascites, hemoglobin)
- Biological Characteristics (age, portalVeinThromb)
- Risk factors (diabetes, artHyper)
- behavioral/demographic (endemicCountries, cigPerYr)

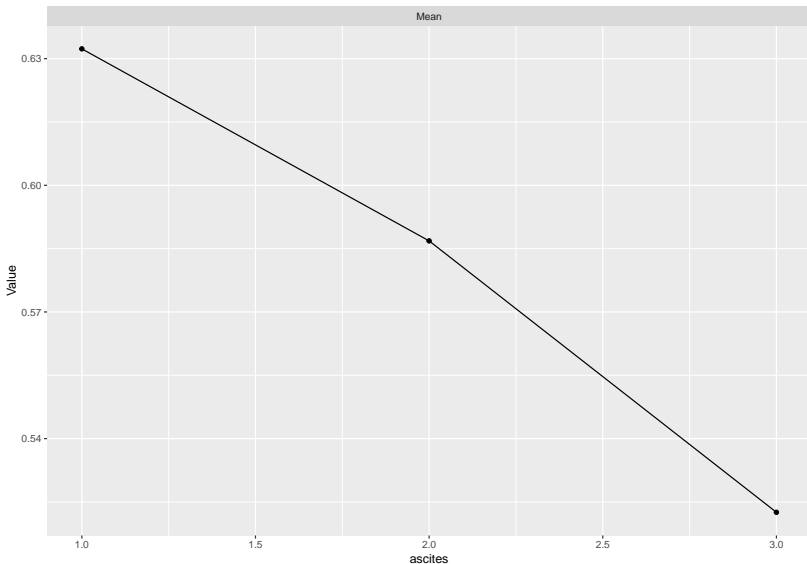
## Partial Dependence: diabetes



# Partial Dependence: ferritin



## Partial Dependence ascites





## Section 6

Final Model choice

## Criteria

- Final model was chosen by the model with the highest Sensitivity

Table 7: Comparison of Sensitivity

	Metric	Mean	Median	SD	Min	Max
RandomForest	Sensitivity	0.824	0.800	0.146	0.5	1.000
XGBTree	Sensitivity	0.774	0.800	0.116	0.6	0.909
SVMRadial	Sensitivity	0.803	0.809	0.107	0.6	0.909

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- Minimizes incorrectly telling a patient they will die

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### References

# References I

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