

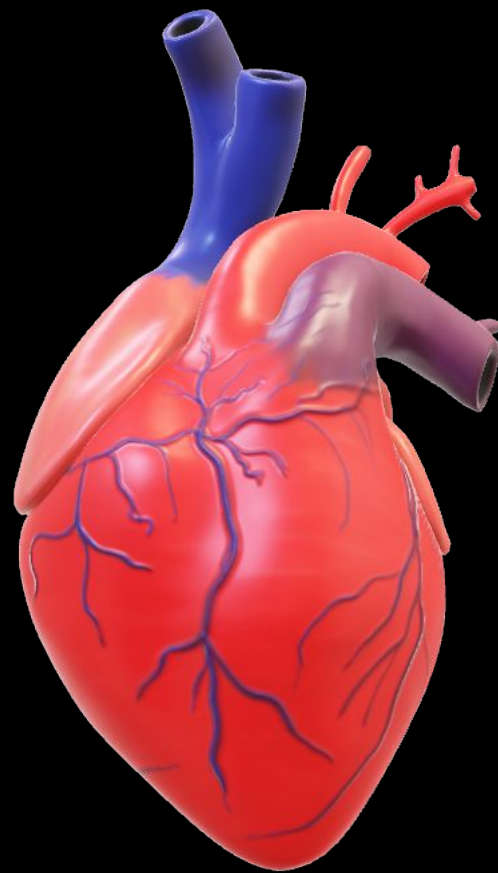
A glowing blue human torso is centered in the background, with light emanating from the neck and chest areas. The torso is semi-transparent, showing internal structures like the heart and lungs in a lighter blue hue. The overall color scheme is dark blue with bright blue highlights.

Predicting Atrial Fibrillation

Bella Falkenberg

Ava Klissouras

Manoj Kambara



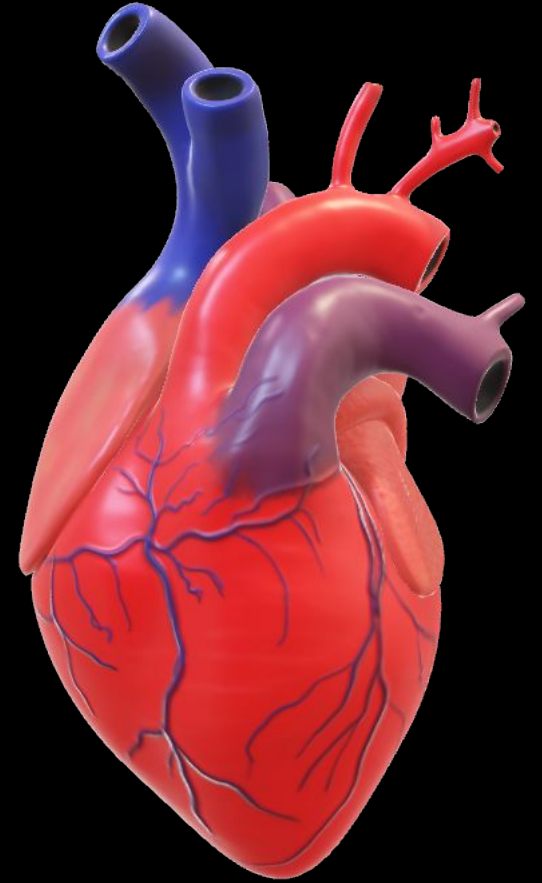
Background

Myocardial Infarction

Muscle of the heart fails to receive enough blood

Major Complications of an MI:

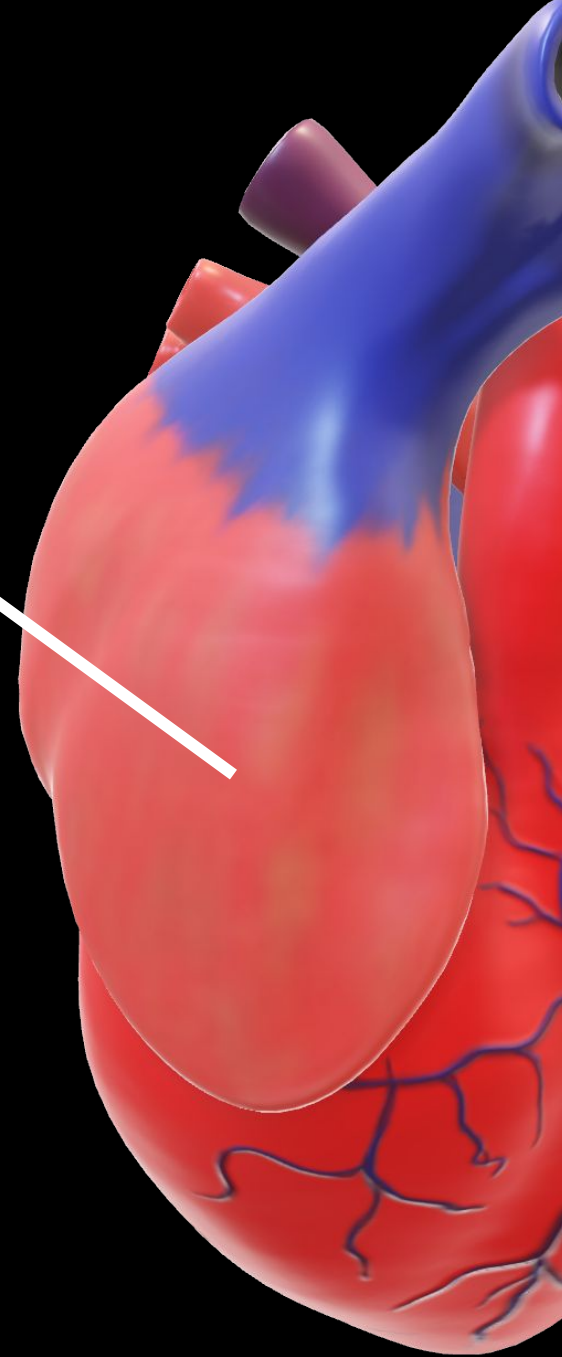
- Atrial Fibrillation
- Pulmonary Edema
- Relapse of MI



Atrial Fibrillation

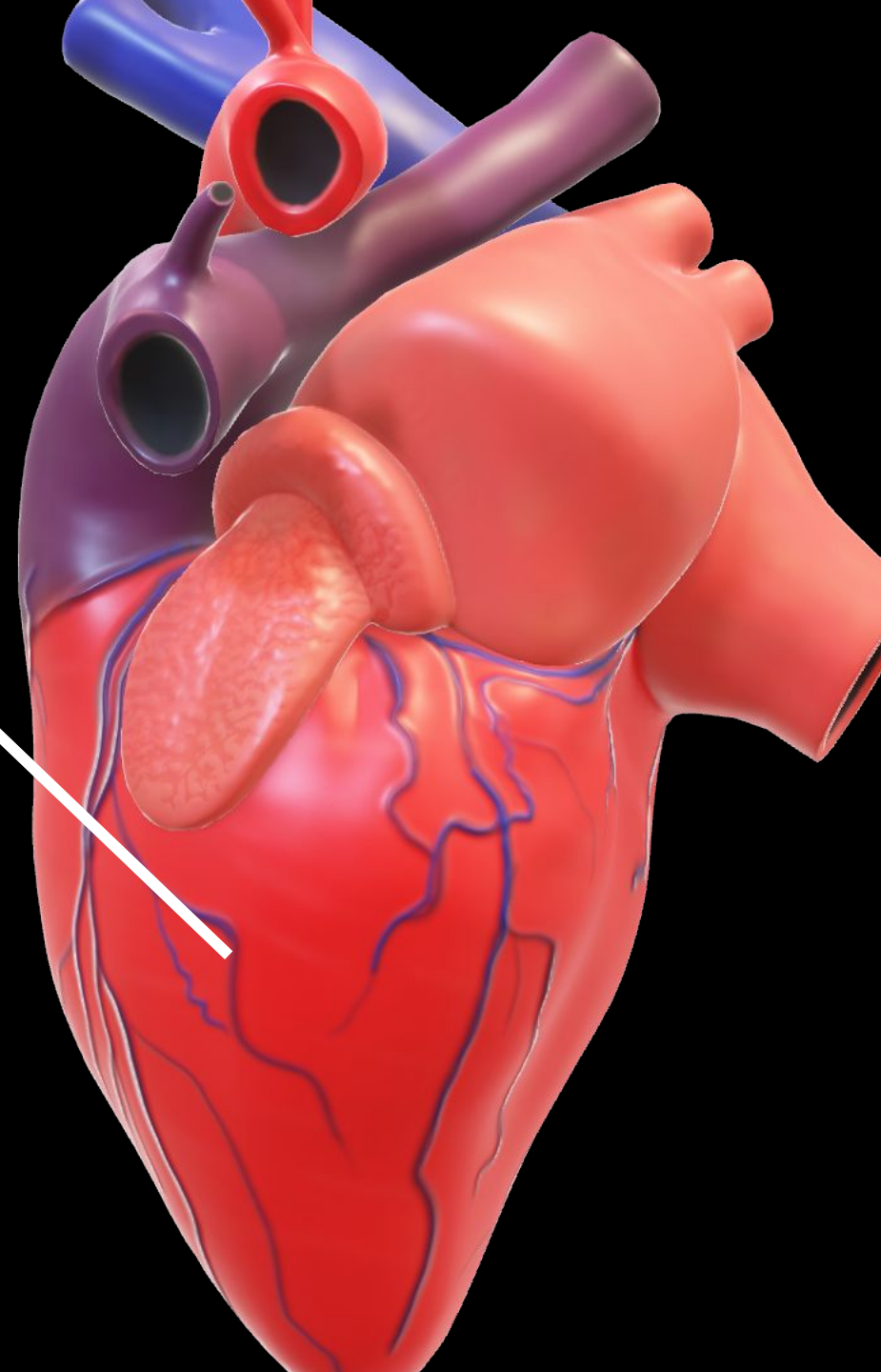
Signals originate from the SA node, located in the right atria of the heart

Irregular contractions of the atria caused by imbalanced signals



Atrial Fibrillation

This causes mistimed contractions of the left ventricle, which pumps blood into the aorta

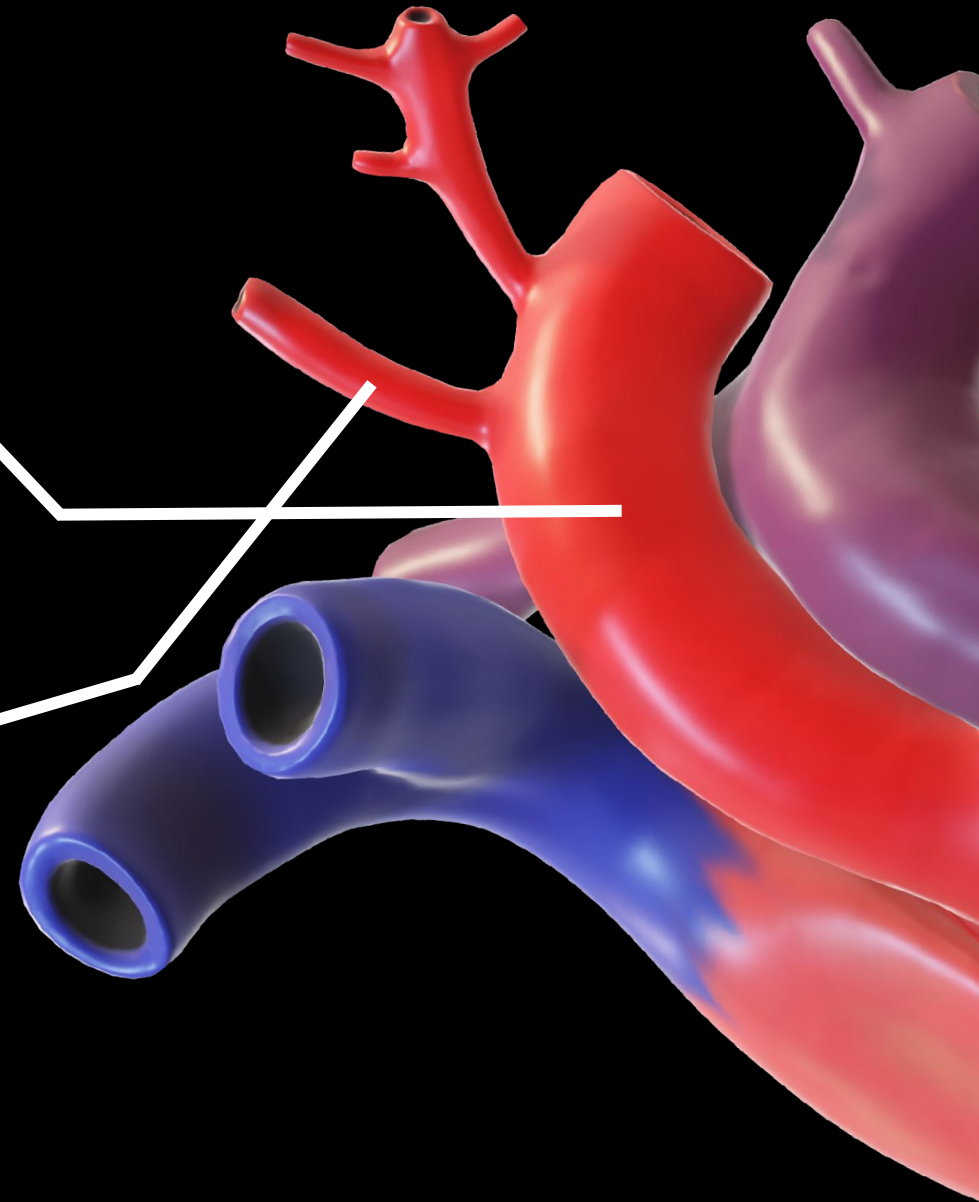


Atrial Fibrillation

AFib can lead to blocks in the aorta

Blocks in the aorta can keep blood from the brain, causing a stroke¹

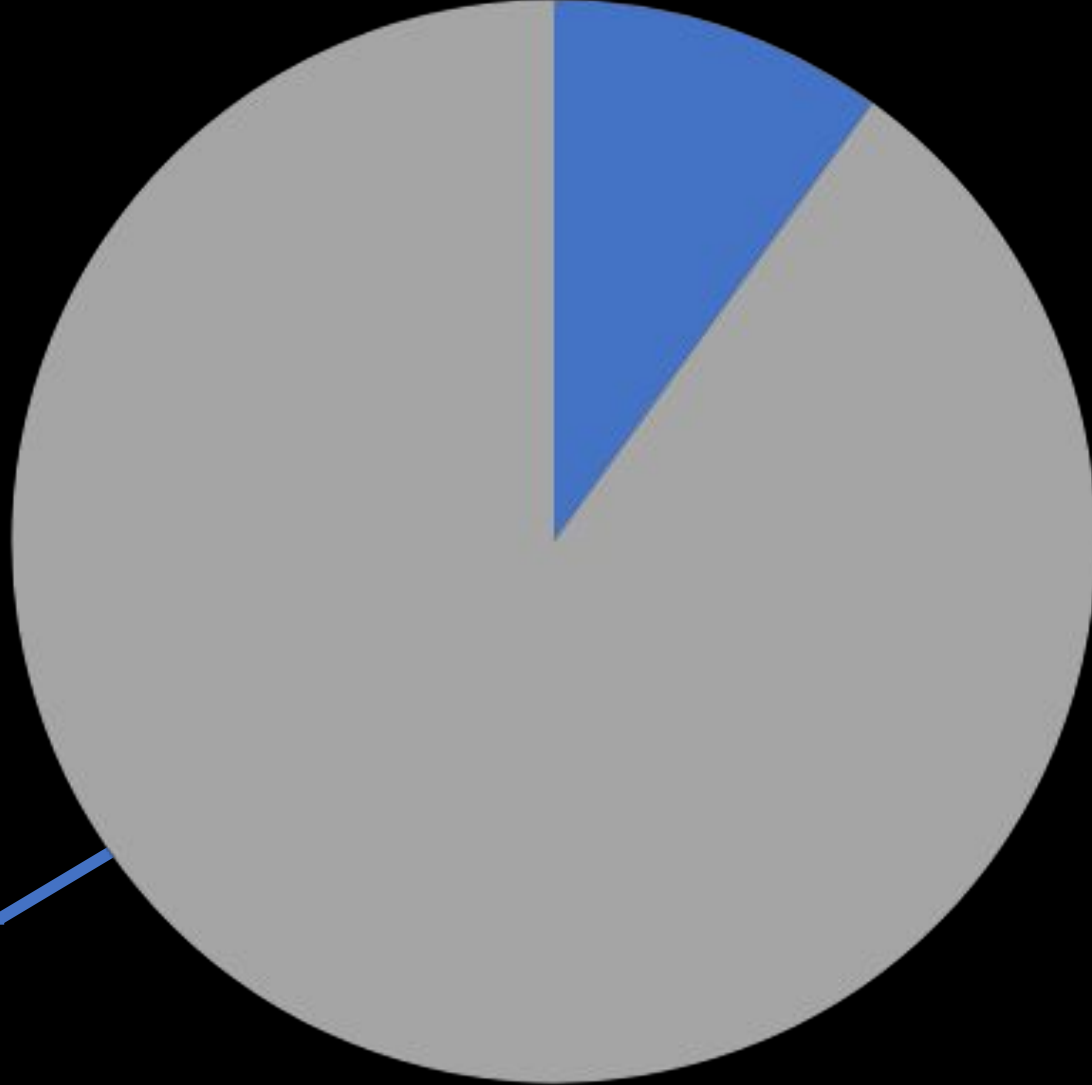
AFib can lead to heart failure¹

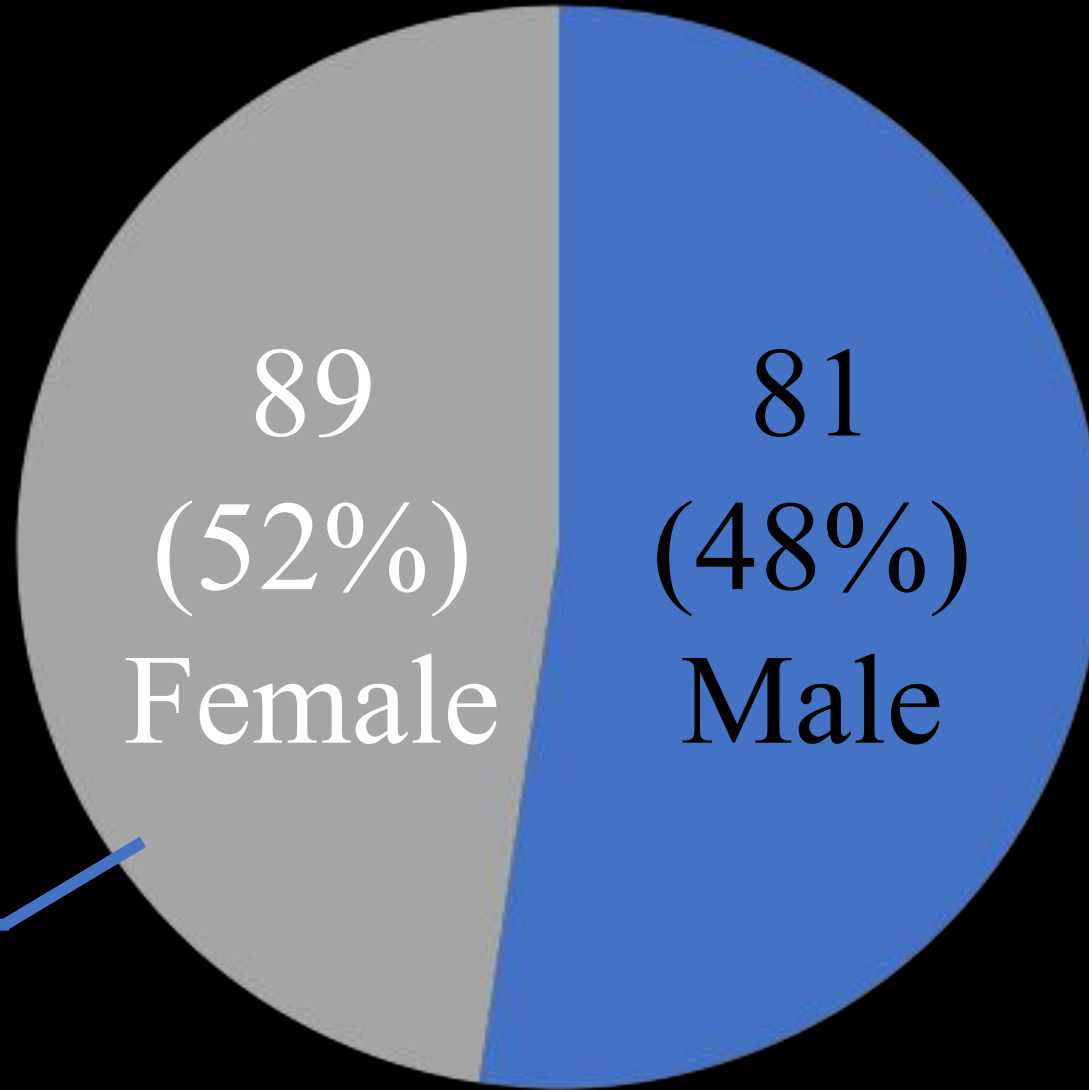




1700
Patients

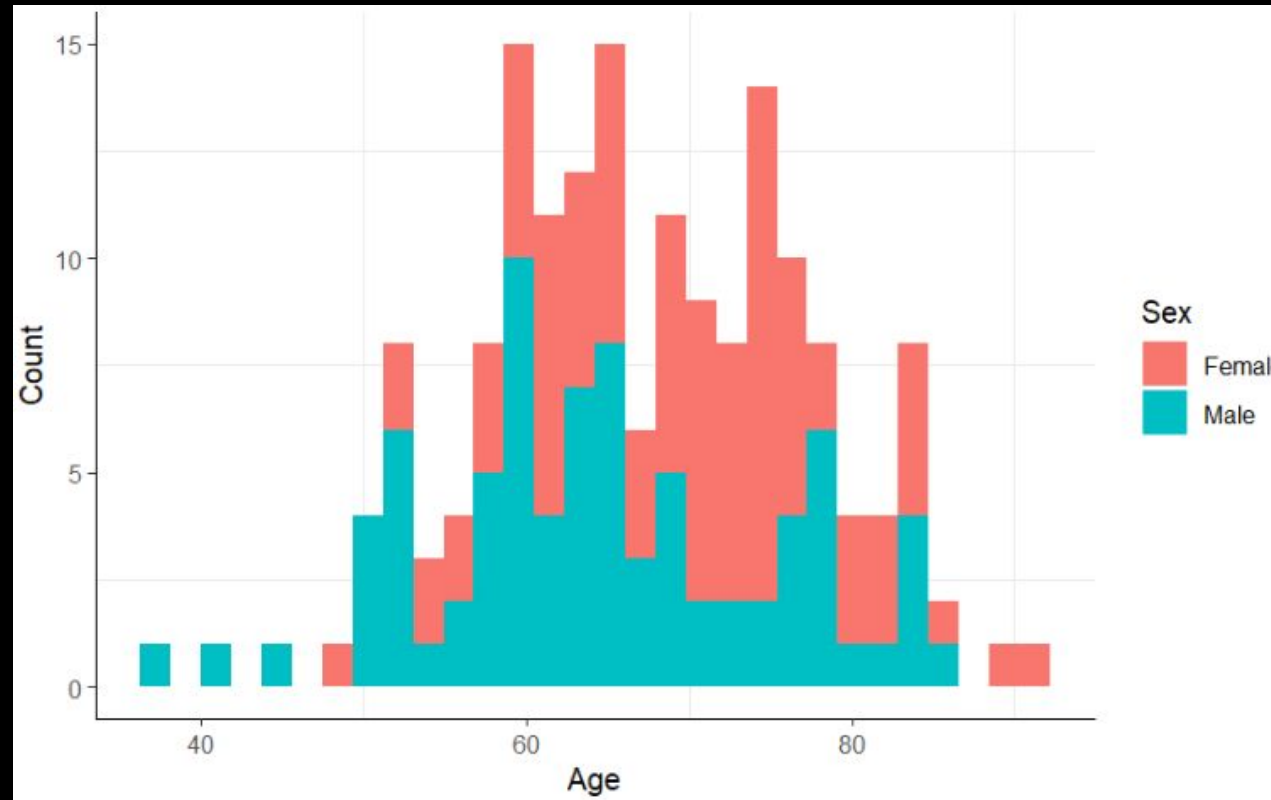
170 (10%)
Patients with
AFib





170 (10%)
Patients with
AFib

Mean: 67.124



Distribution of ages of patients that developed AFib as a complication of their MI.

Aim

Create models to predict probability of AFib as a complication following treatment for an MI.

1. Find and apply a method to clean the data
2. Use R methods to make a model
3. Evaluate the models for calibration and discrimination

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Missing Data Imputation

Multiple Imputation by Chained Equations (MICE)

MICE Imputation²: Data Preparation

| | | |
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| | | |
| | | |
| | | |

Removed columns in which greater than 50% of values were missing

- KFK_Blood
- Serum PCK content
- IBS_NASL
- Heredity on CHD
- S_AD_KBRIG/D_AD_KBRIG
- Systolic/Diastolic by emergency cardiology team

MICE Imputation²: Data Preparation

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Removed a highly correlated column

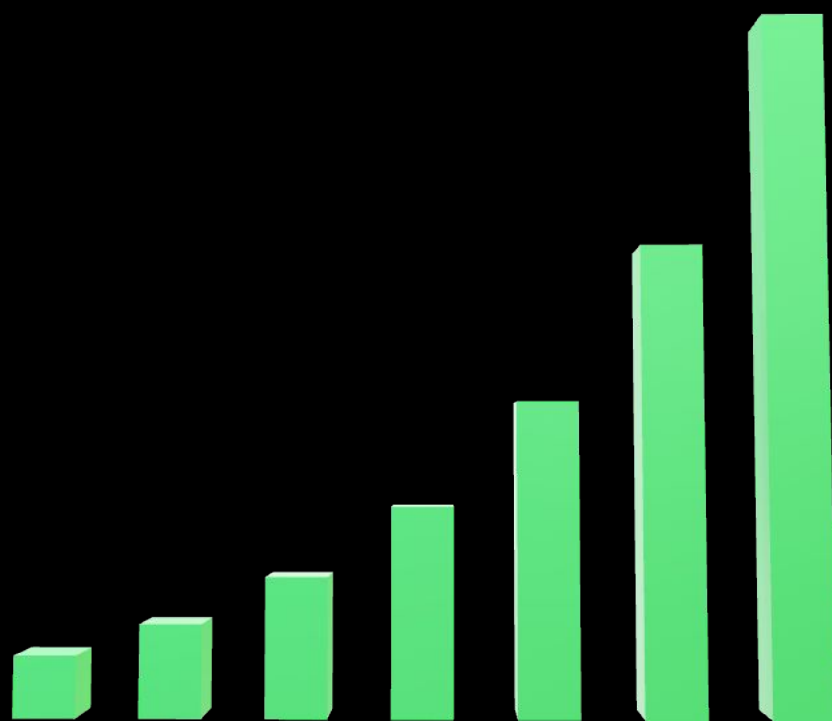
- S_AD_ORIT
- Systolic BP by ICU
- Used `findCorrelation()`
- Used a cutoff of 0.8
- Assumed that data is Missing at Random (MAR)
- Applied the function `mice()` to the data set

MICE Imputation²: Data Preparation

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Removed a highly correlated column

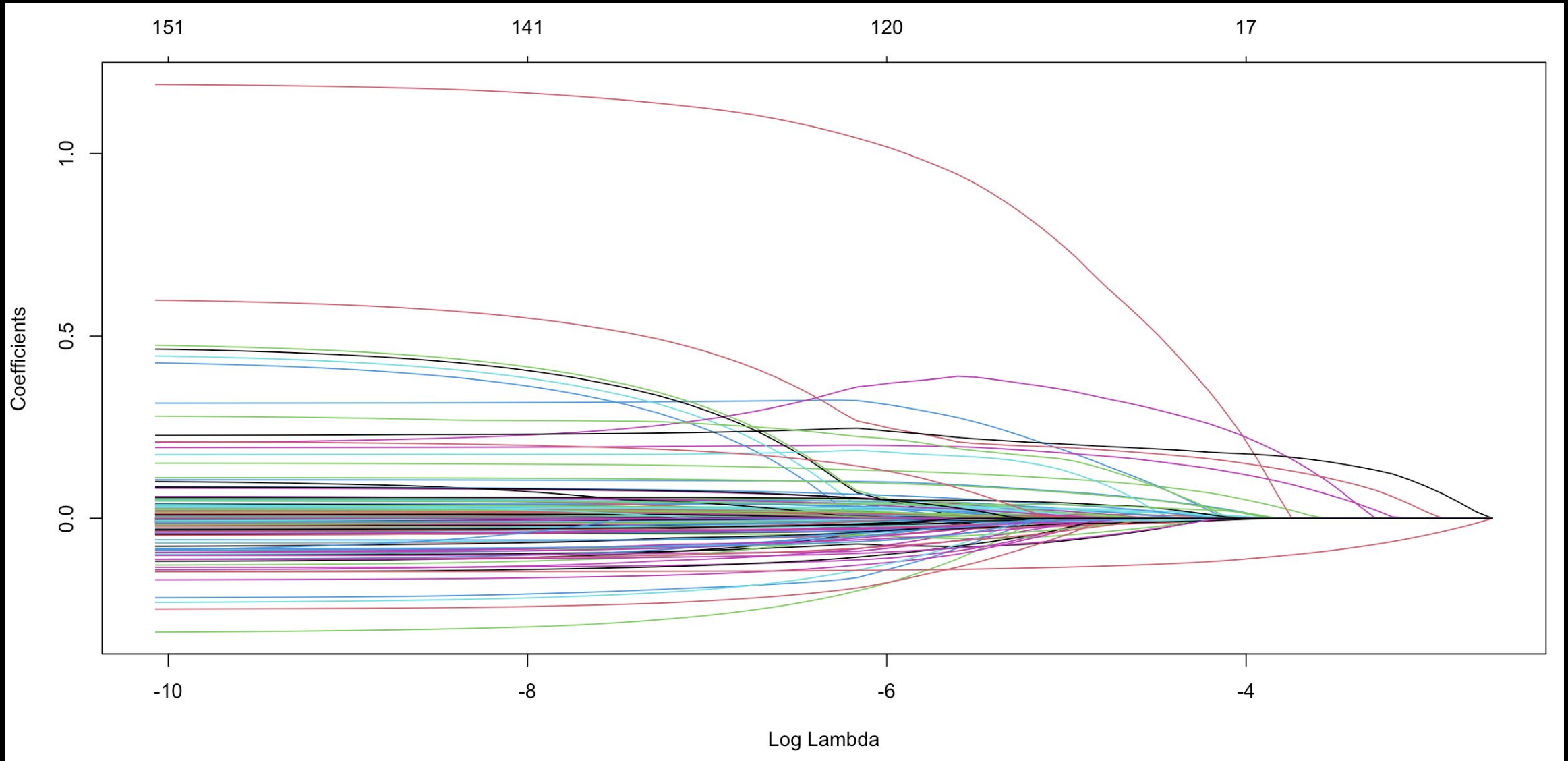
- S_AD_ORIT
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Modeling

LASSO Regression

Introduction to LASSO

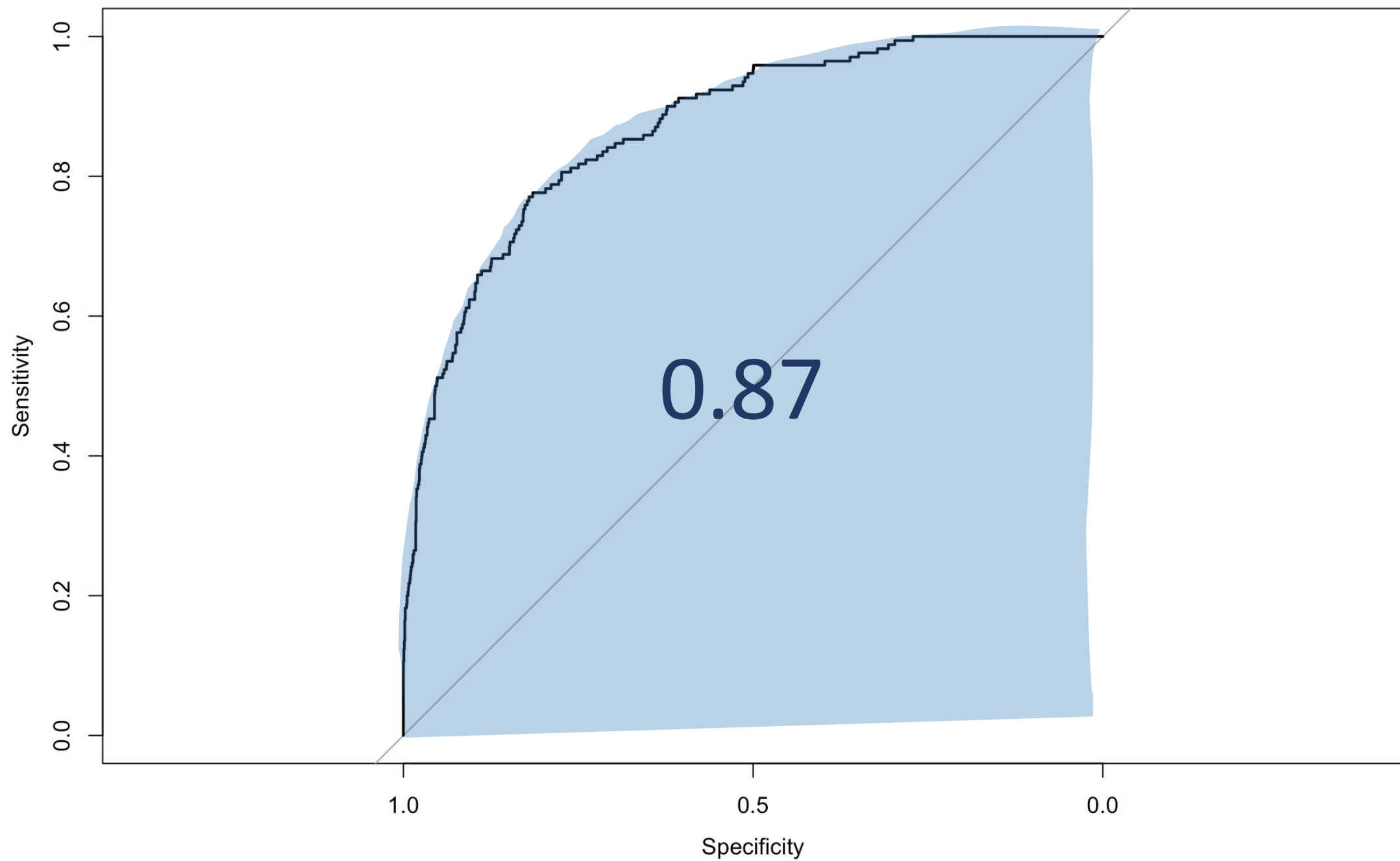


Our LASSO Regression

- Categorical variables --> dummy variables
- p-value = 6.59E-9
- 50 significant covariates

Table 1. Significant covariates for LASSO regression

| Covariate | Log Odds | Probability | p-value |
|--|----------|-------------|---------|
| Age (AGE) | 0.037 | 0.51 | 0.003 |
| Afib in anamnesis (nr03) | 1.73 | 0.85 | <0.001 |
| ECG rhythm @ hospital (bradycardia) (ritm_ecg_p_08) | 2.33 | 0.91 | <0.001 |
| ECG rhythm @ hospital (Afib) (n_r_ecg_p_05) | 1.82 | 0.86 | <0.001 |
| NSAID use by ECT (NOT_NA_KB) | -0.41 | 0.12 | <0.001 |



Evaluating LASSO

- Calibration plot
- Discrimination plot
- AUC of ROC plot
- Confusion Matrix
 - Accuracy = 0.8112
 - Sensitivity = 0.9704
 - Specificity = 0.3181

| | | Actual | |
|---------------|------------------------------------|----------------------|-------------------------|
| | | AFib Complication | No AFib Complication |
| Pred icted | AFib Com plicat ion | 1247 | 283 |
| | No AFib Com plicat ion | 38 | 132 |

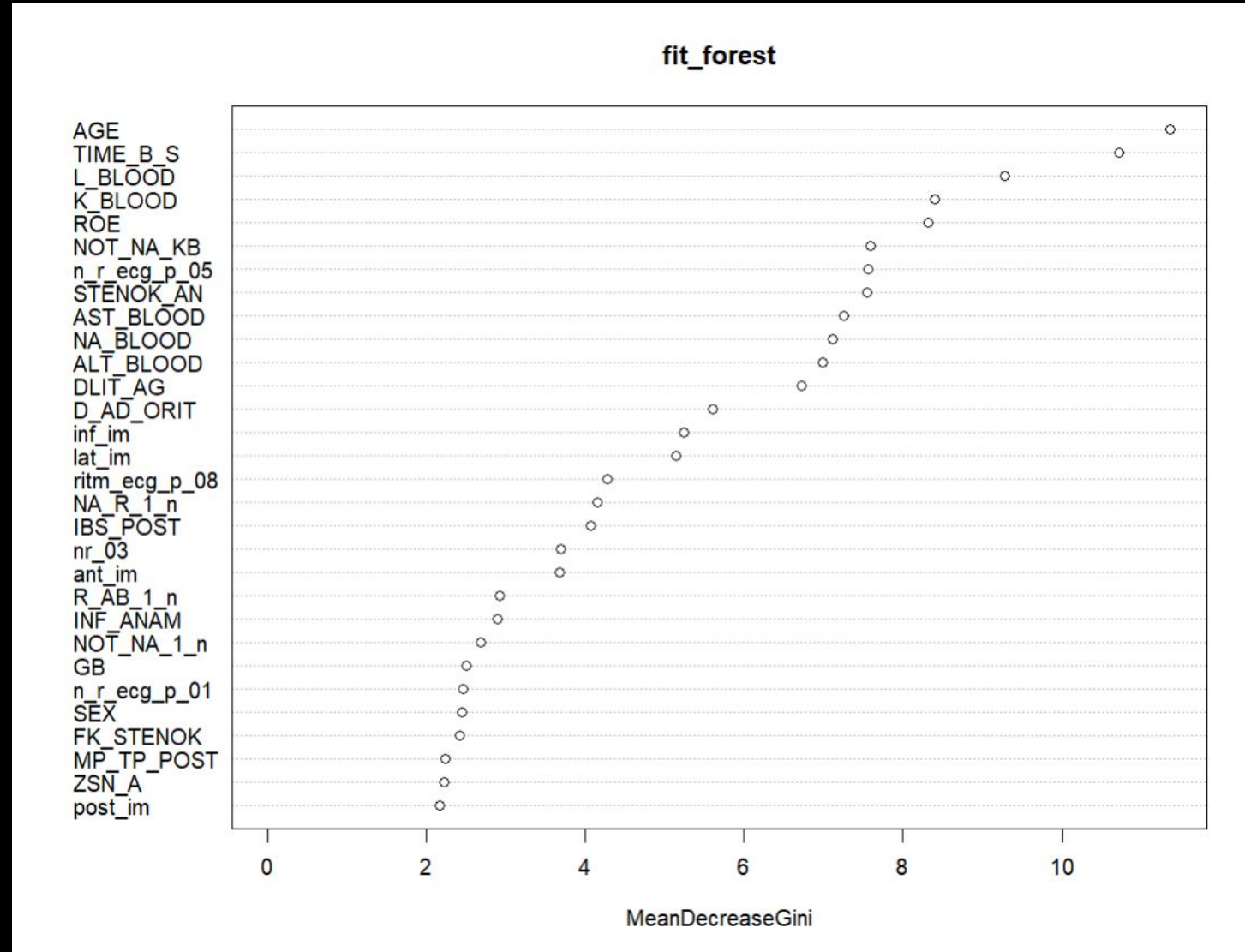
Random Forest

Random Forest

- Supervised machine learning model
- Combines the output of multiple decision trees in order to make a single result
- Can perform both regression and classification
- Random forests can be greedy

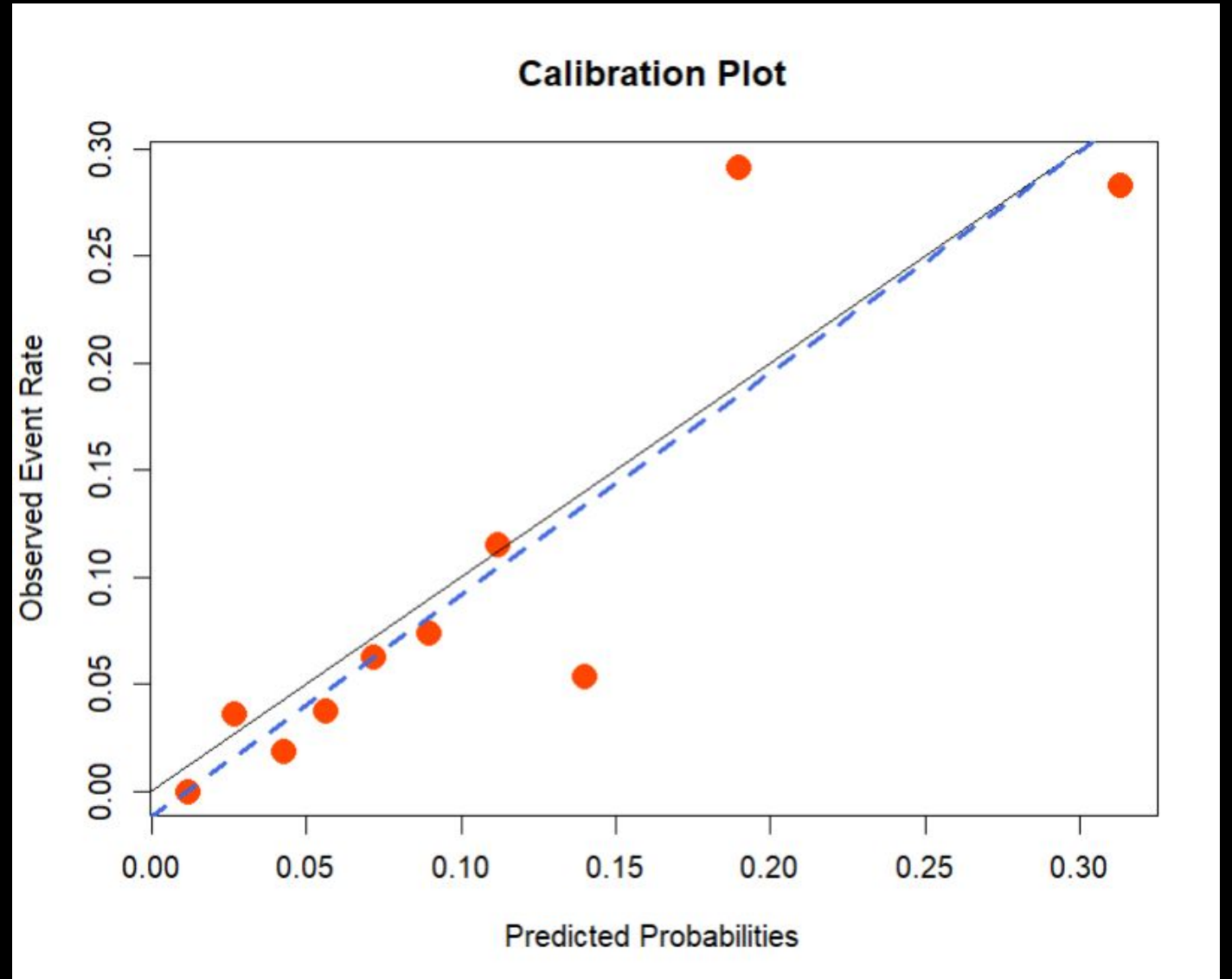
Our Random Forest Model

- We split the data into a training set (70%) and a testing set (30%)
 - Most significant:
 - Age
 - Time to hospital
 - Blood content



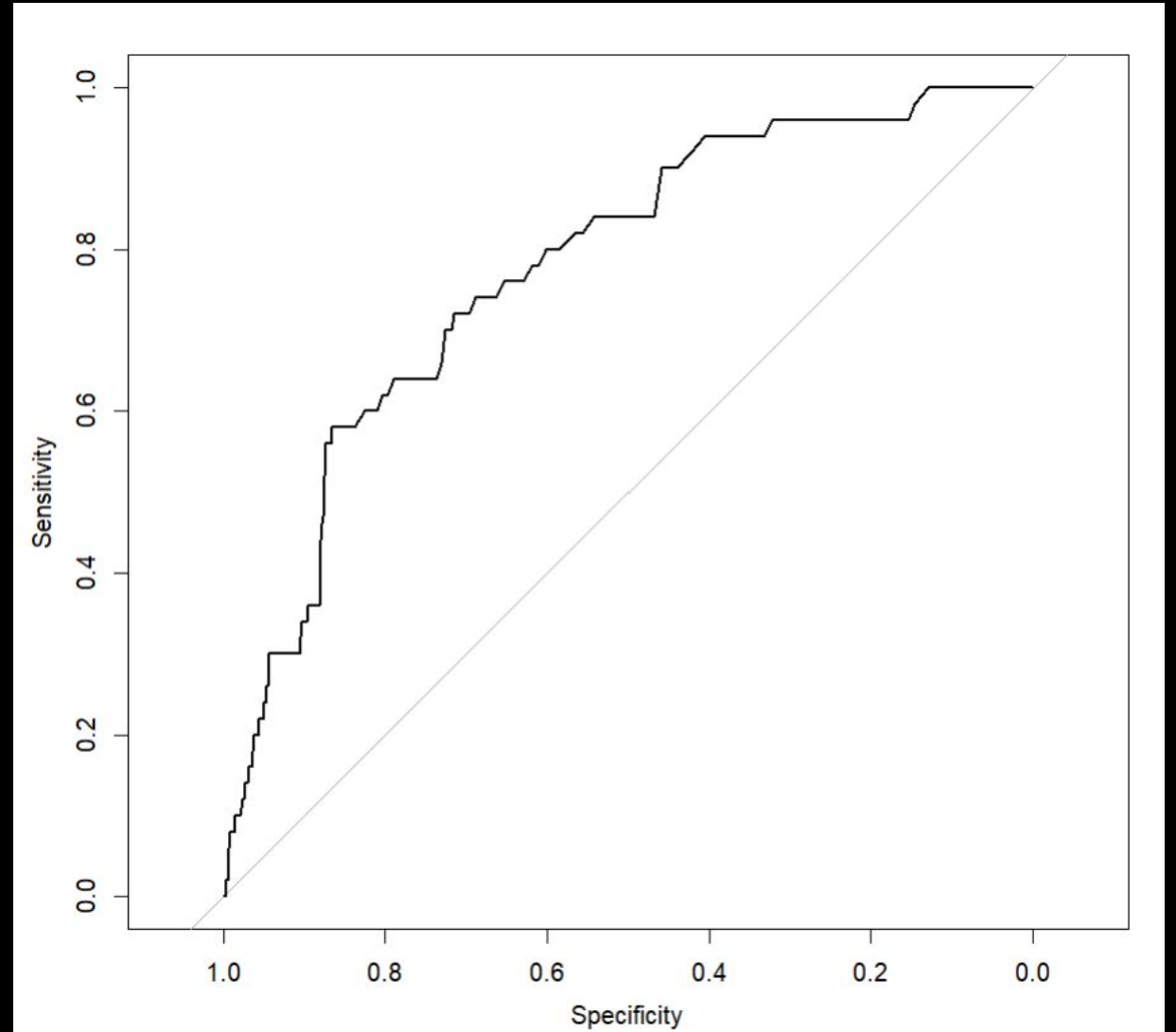
Our Random Forest Model

- We split the data into a training set (70%) and a testing set (30%)
- Calibration Plot
 - Slope
 - 1.04 [0.62,1.45]
 - Y-intercept
 - -0.01 [-0.07,0.05]



Our Random Forest Model

- We split the data into a training set (70%) and a testing set (30%)
- Calibration Plot
- ROC Curve
 - AUC = 0.78



Our Random Forest Model

- We split the data into a training set (70%) and a testing set (30%)
- Calibration Plot
- ROC Curve
 - AUC = 0.78
- Confusion Matrix
 - Accuracy = 0.8381
 - Sensitivity = 0.9514
 - Specificity = 0.3118

| | | Actual | |
|-----------|----------------------|-------------------|----------------------|
| | | AFib Complication | No AFib Complication |
| Predicted | AFib Complication | 411 | 64 |
| | No AFib Complication | 21 | 29 |

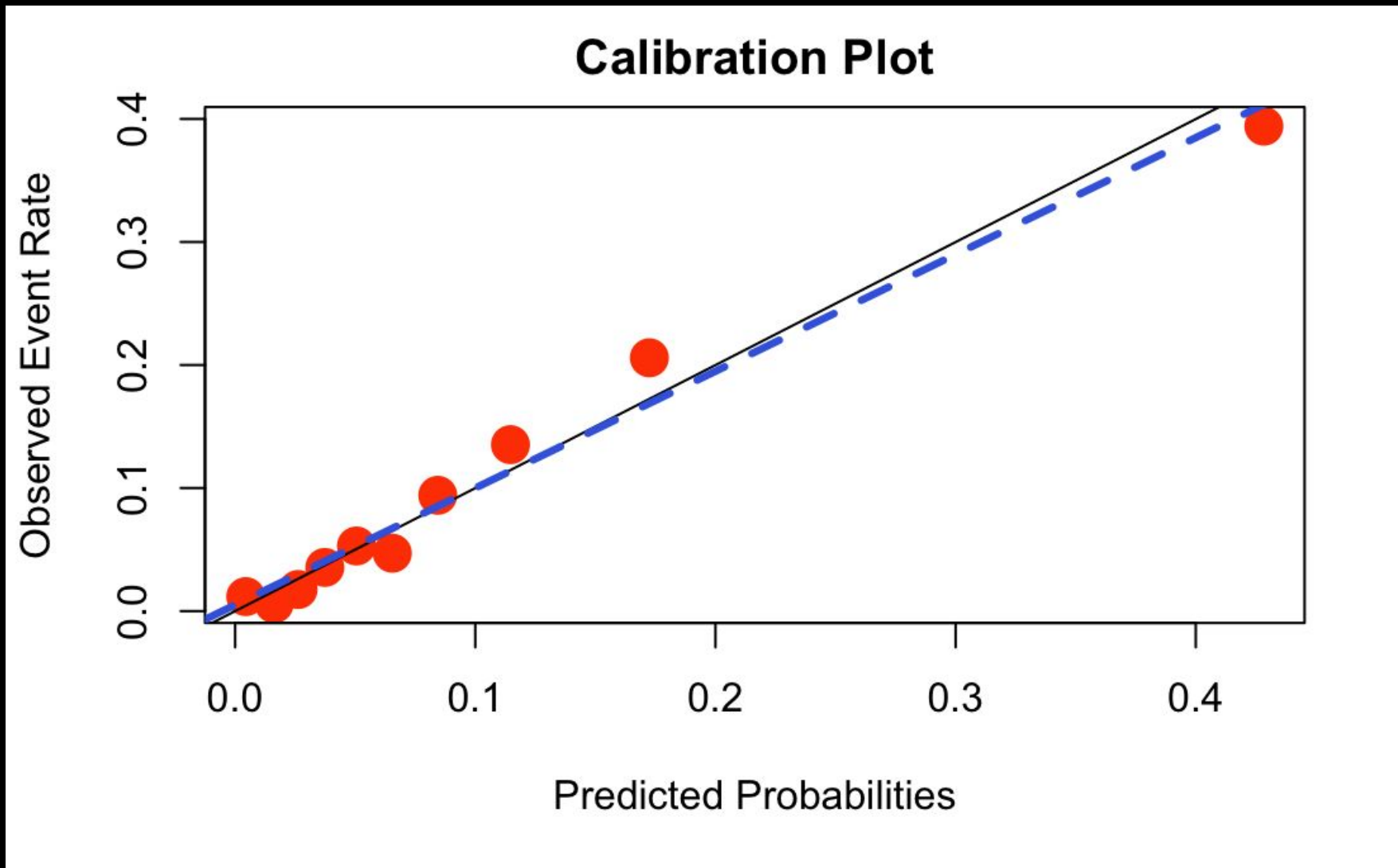
Forward Selection

Our Forward Selection

- We chose forward selection because:
 - Step and backwards selection were computationally intensive
- Used $\alpha = 0.05$
- Final model had 25 predictors

Evaluating Forward Selection

- Calibration Plot

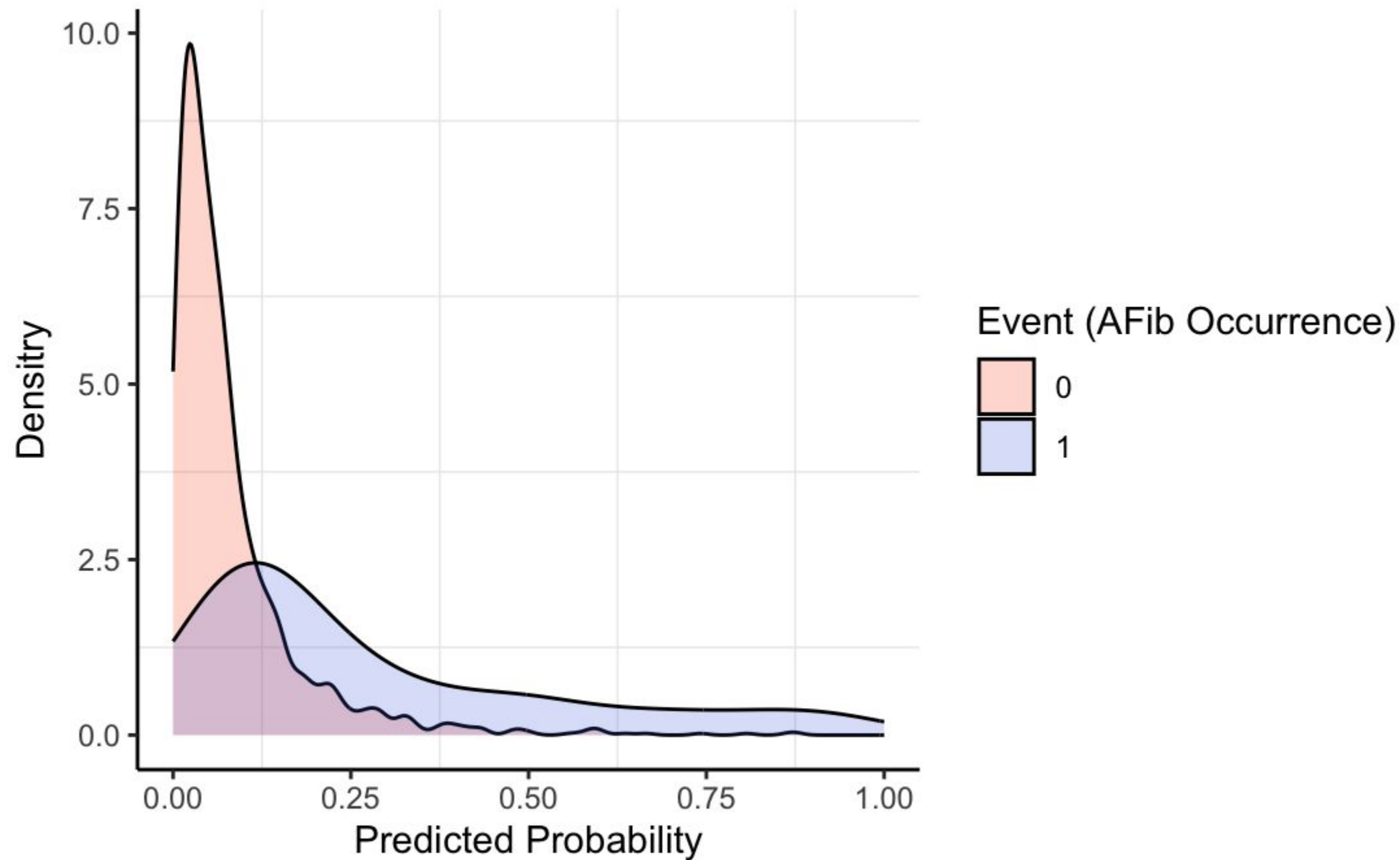


Slope: point estimate = 0.949
confidence interval [0.831, 1.068]

Intercept: point estimate = 0.005
confidence interval [-0.013, 0.024]

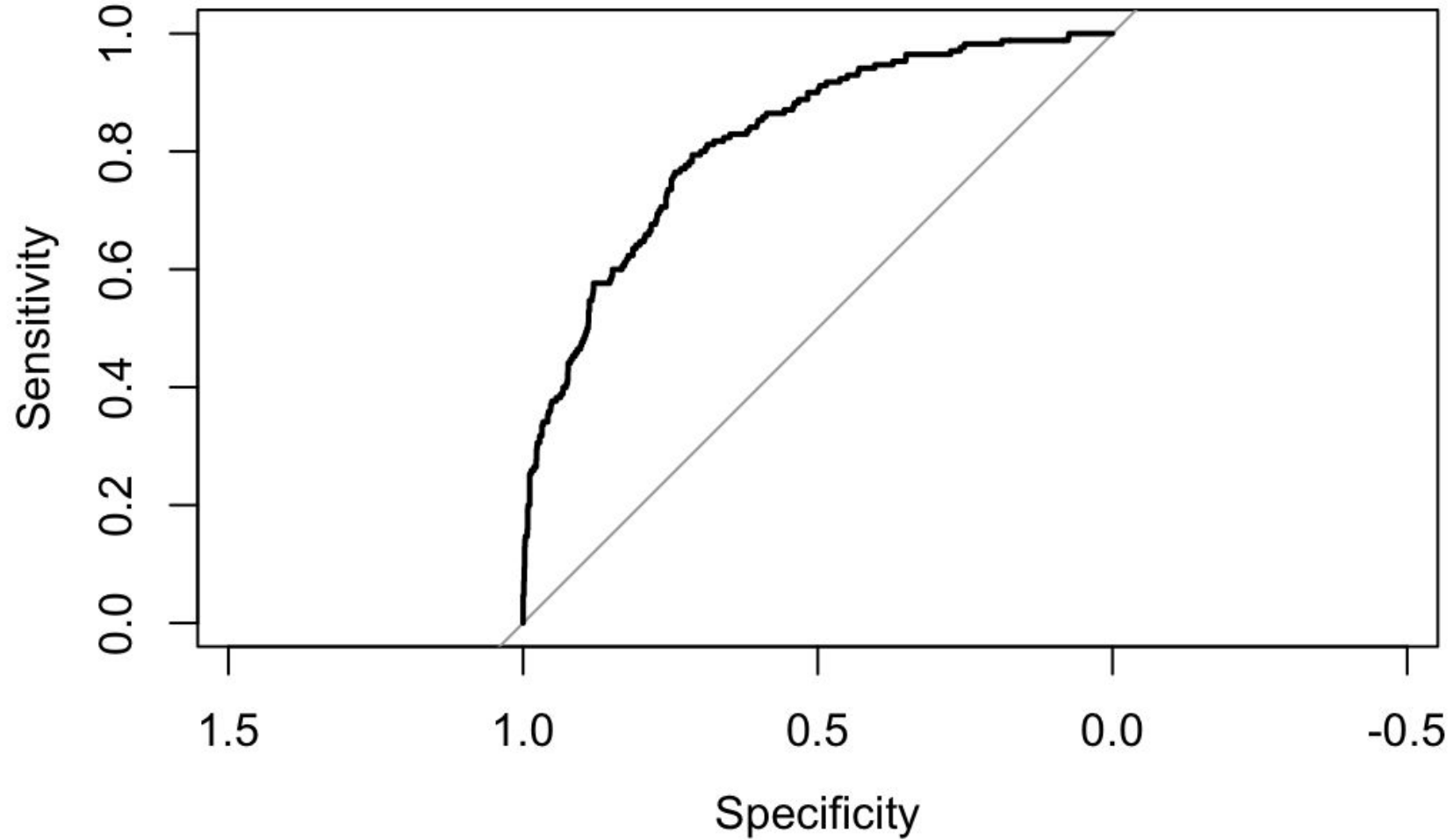
Evaluating Forward Selection

- Calibration Plot
- Discrimination Plot



Evaluating Forward Selection

- Calibration Plot
- Discrimination Plot
- AUC of ROC Plot



AUC: point estimate = 0.8207
Confidence Interval [0.7880, 0.8534]

Evaluating Forward Selection

- Calibration Plot
- Discrimination Plot
- AUC of ROC Plot
- Confusion Matrix
 - Accuracy = 0.8229
 - Sensitivity = 0.9443
 - Specificity = 0.2934

| | | Actual | |
|-------------------|--------------------------------|----------------------|-------------------------|
| | | AFib Complication | No AFib Complication |
| Pre dict ed | AFib Compli cation | 1306 | 224 |
| | No AFib Compli cation | 77 | 93 |

Best Model

Forward Selection

Why Forward Selection?

- Well-calibrated
- AUC under ROC = 0.82
- High accuracy (0.823)

Covariates: Anamnesis

Table 2. Significant covariates of anamnesis from the forward selection model

| Covariate | Log Odds | Probability | p-value |
|---|----------|-------------|---------|
| Atrial Fibrillation (nr03) | 1.75 | 0.85 | <0.001 |
| Persistent Atrial Fibrillation (nr04) | -2.26 | 0.09 | 0.048 |
| Ventricular Fibrillation (nr07) | 17.14 | 0.99 | 0.974 |
| Arrhythmia (nr11) | 1.12 | 0.75 | 0.011 |
| Chronic Bronchitis (zab_leg_01) | -0.88 | 0.29 | 0.055 |
| Bronchial Asthma (zab_leg_03) | 1.21 | 0.77 | 0.001 |
| Exertional Angina Pectoris (IBS_POST1) | 0.32 | 0.58 | 0.193 |
| Unstable Angina Pectoris (IBS_POST2) | -0.45 | 0.39 | 0.082 |

Covariates: Demographic & Testing

Table 3. Significant covariates of demographics and testing from the forward selection model

| Covariate | Log Odds | Probability | p-value |
|------------------------------|----------|-------------|---------|
| Sex (SEX) | -0.45 | 0.39 | <0.001 |
| Age (AGE) | 0.04 | 0.51 | 0.03 |
| Hypokalemia (GIPO_K) | -0.79 | 0.31 | 0.006 |
| Blood Potassium (K_BLOOD) | -0.78 | 0.31 | <0.001 |

Covariates: Treatments

Table 4. Significant covariates of treatments from the forward selection model

| Covariate | Log Odds | Probability | p-value |
|--|----------|-------------|---------|
| Opioids Used on Day 3 Once (NA_R_3_n1) | 1.13 | 0.76 | 0.001 |
| Opioids Used on Day 3 Twice (NA_R_3_n2) | -0.16 | 0.46 | 0.880 |
| NSAID Use by ECT (NOT_NA_KB) | -1.89 | 0.13 | <0.001 |
| Lidocaine Use in ICU (LID_S_n) | 0.70 | 0.67 | <0.001 |
| Heparin use in ICU (GEPAR_S_n) | 0.45 | 0.61 | 0.041 |

Covariates: ECG Measurements

Table 5. Significant covariates of ECG measurements from the forward selection model

| Covariate | Log Odds | Probability | p-value |
|--|----------|-------------|---------|
| Paroxysms of Atrial Fibrillation on ECG (n_r_ecg_p_05) | 1.29 | 0.78 | <0.001 |
| Premature Atrial Contractions on ECG (n_r_ecg_p_01) | 1.12 | 0.75 | 0.002 |
| Idioventricular on ECG (ritm_ecg_p_06) | 2.99 | 0.95 | 0.047 |
| Bradycardia on ECG (ritm_ecg_p_08) | 2.05 | 0.89 | <0.001 |



Conclusions

Conclusions

- The model developed from forward selection was the best model
- This model included many different factors, from patient history, patient biometrics, treatments, and demographics
- Is well calibrated, discriminates well, and had a high accuracy
- Limitations
 - This model is done solely on a Russian cohort which is not fully representative of a wide range of patient populations
 - The data set was missing 7.6% of values. Using imputation adds more variance to the model

Acknowledgements

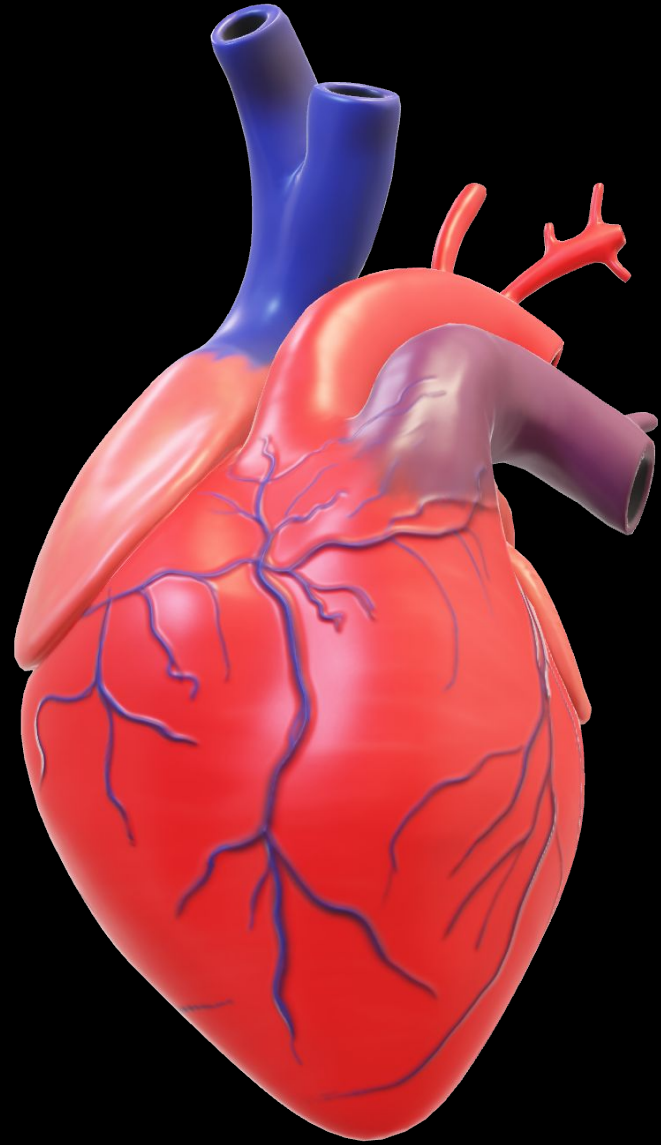
SIBS Faculty

SIBS Mentors

Dr. Sujit Ghosh

Dr. Herle McGowan

Dr. Megan Neely



References

1. Atrial fibrillation. Heart and Stroke Foundation of Canada. Accessed July 14, 2023.
<https://www.heartandstroke.ca/heart-disease/conditions/atrial-fibrillation#:~:text=People%20with%20atrial%20fibrillation%20have,get%20pumped%20to%20the%20brain>.
2. Azur MJ, Stuart EA, Frangakis C, Leaf PJ. Multiple imputation by chained equations: what is it and how does it work?. *Int J Methods Psychiatr Res.* 2011;20(1):40-49. doi:10.1002/mpr.329