



# Interpretable Machine Learning Model to Predict and Influence Mortality of Patients with Heart Failure Warded in Intensive Care Unit

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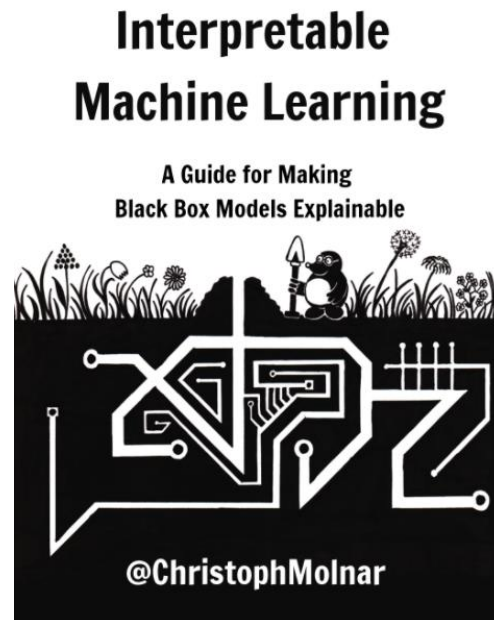
WONG CHI KEONG JOHN  
3 JAN 2022

GENERAL ASSEMBLY  
DATA SCIENCE IMMERSIVE FLEX 2  
CAPSTONE PROJECT

# Data Science Process

- **Objective**
- Collect
- Explore
- Engineer feature
- Build model
- Conclude on objective
- Deployment workflow

- Develop interpretable machine learning model to predict and influence mortality of patients with heart failure warded in intensive care unit
- Interpretable
  - *“Interpretability is the degree to which a human can understand the cause of a decision.”*
  - *“Interpretability is the degree to which a human can consistently predict the model’s result”*





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- Data from Kaggle
  - <https://www.kaggle.com/saurabhshahane/in-hospital-mortality-prediction>
- Paper from journal
  - <https://bmjopen.bmj.com/content/11/7/e044779>

### **BMJ Open Prediction model of in-hospital mortality in intensive care unit patients with heart failure: machine learning-based, retrospective analysis of the MIMIC-III database**

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Fuhai Li,<sup>1,2</sup> Hui Xin,<sup>1</sup> Jidong Zhang,<sup>1</sup> Mingqiang Fu,<sup>2</sup> Jingmin Zhou <sup>2</sup>,  
Zhexun Lian <sup>1</sup>

## Data Science Process

- Objective
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- Data frame size
  - 51 variables × 1,177 cases
- Variable type, category count, encoding
- Missing
- Duplicate
- Imbalance
- Multicollinearity

## Data Science Process

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- Data frame size
- Variable type, category count, encoding
  - 11 features  $\Rightarrow$  Categorical, binary, encoded in numeric
  - 38 features  $\Rightarrow$  Continuous
  - Target  $\Rightarrow$  Categorical, binary, encoded in numeric
  - ID feature  $\Rightarrow$  Drop
- Missing
- Duplicate
- Imbalance
- Multicollinearity

No need one hot encoding

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- Data frame size
- Variable type, category count, encoding
- Missing
  - 8 features have missing values from 12.2% to 25.0%
  - 11 features have missing values from 0.1% to 3.1%
  - Target (“outcome”) has 1 missing value
- Duplicate
- Imbalance
- Multicollinearity

Drop because too much to impute  
without affecting original probability  
distribution

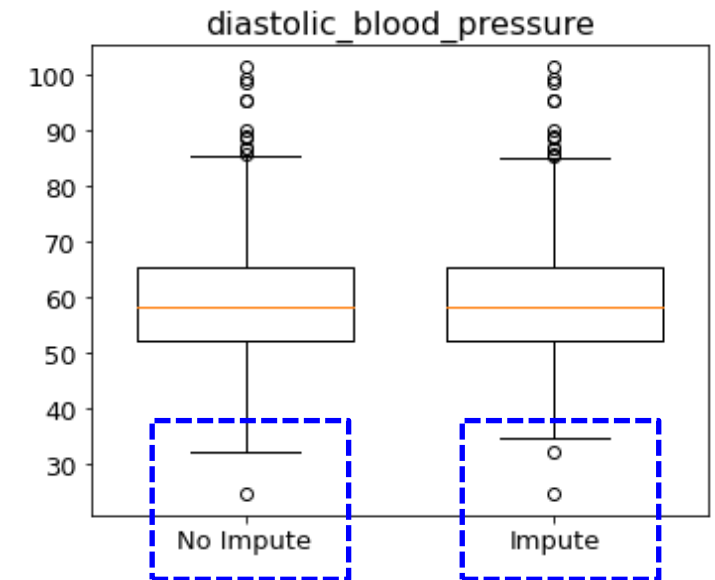
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Impute using KNNImputer with neighbouring samples = 5

Check original probability distribution not significantly affected (change in outliers, percentile and whiskers values)



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Drop missing case before train test split



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- Data frame size
- Variable type, category count, encoding
- Missing
- Duplicate
  - No duplicate cases
- Imbalance
- Multicollinearity

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- Data frame size
- Variable type, category count, encoding
- Missing
- Duplicate
- Imbalance
  - Die (“1”) at 13.5%
  - Live (“0”) at 86.5%
- Multicollinearity

Need to oversample minority category using SMOTE

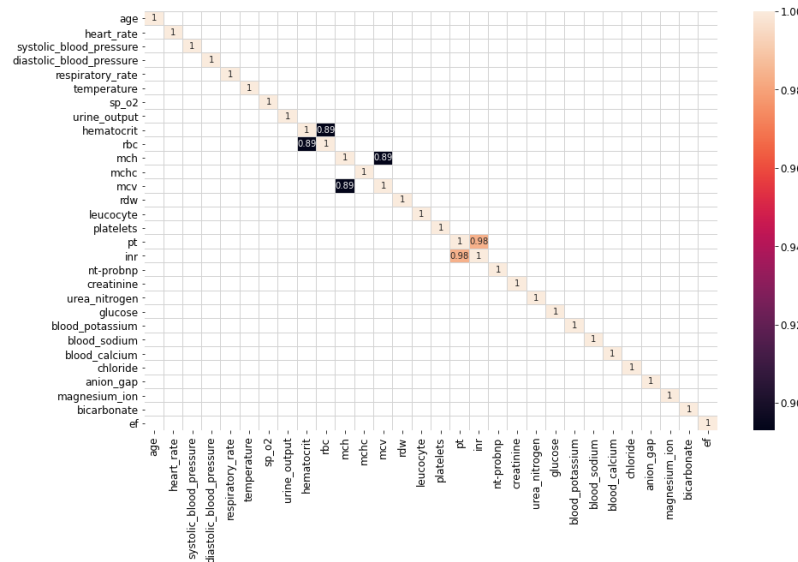
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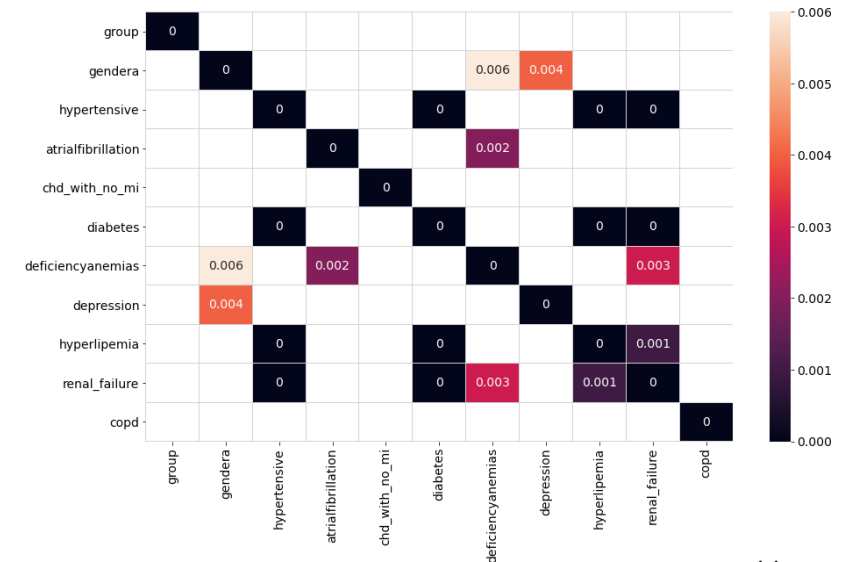
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- Multicollinearity

Need to regularise with L1 (LASSO) as many features are strongly correlated with each other

Pearson (continuous, more than  $R^2 \geq 0.80$ )



Chi Square (categorical, less than  $p < 0.01$ )



## Data Science Process

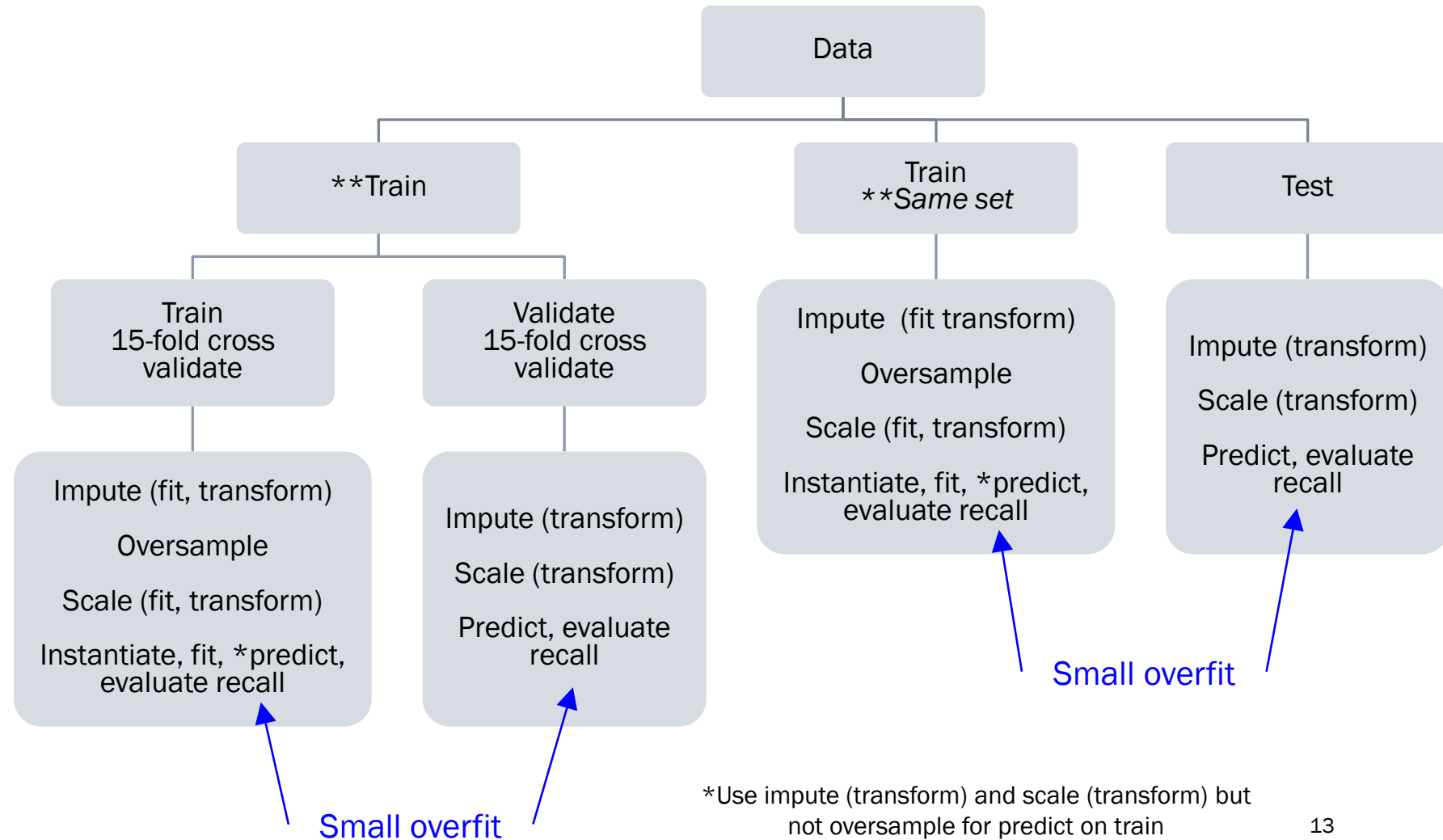
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- Model requirement
  - Want high recall  $\Rightarrow TP \div (TP + FN)$ 
    - FN  $\Rightarrow$  Cannot save patient who will actually die
  - Balance with precision  $\Rightarrow TP \div (TP + FP)$ 
    - FP  $\Rightarrow$  Unnecessary extra resources spent on patient who will actually live
  - Few important features to facilitate interventions to change mortality
  - Smallest overfit

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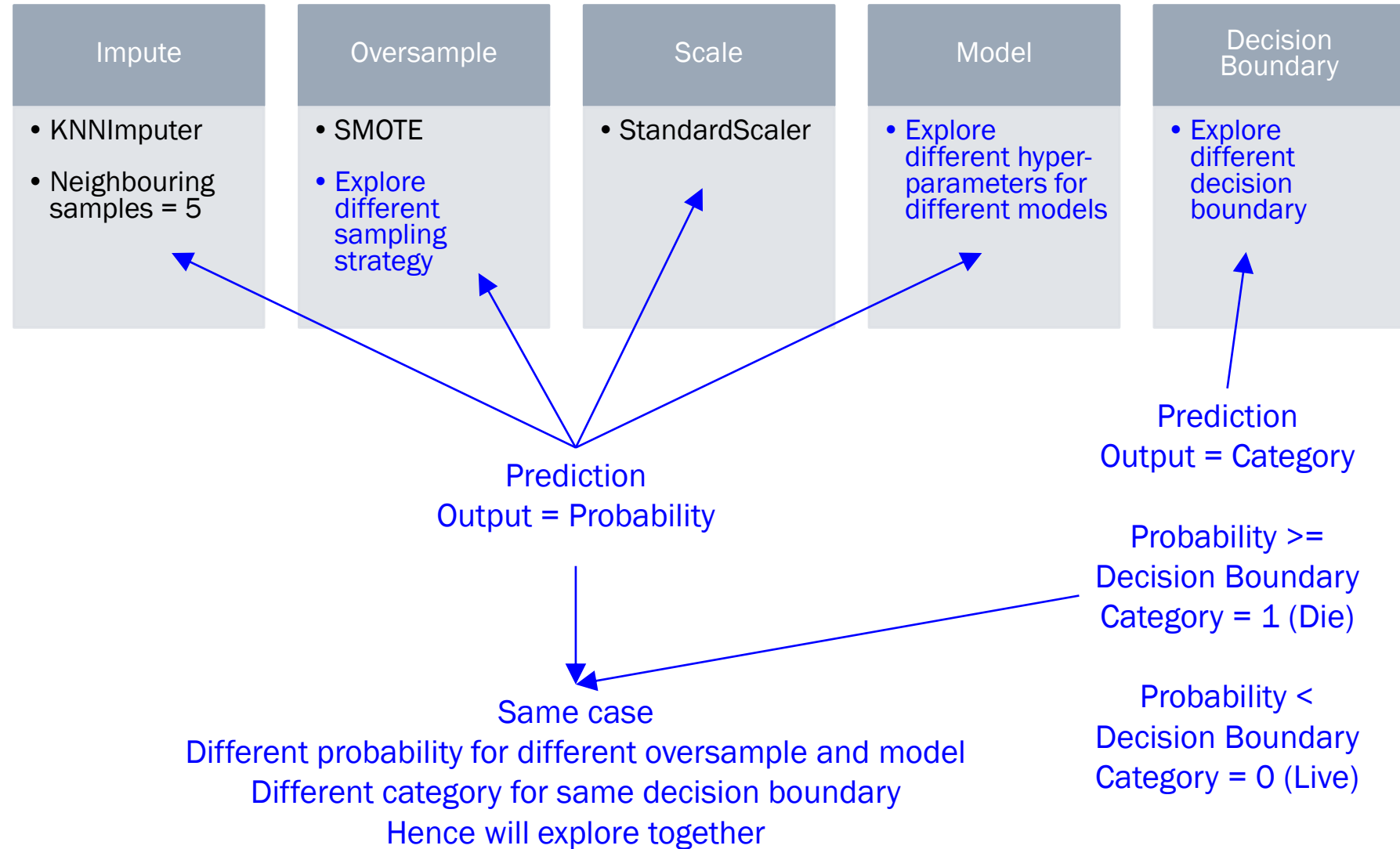
- Model workflow
  - Mechanics of Pipeline and GridSearchCV to do manually for model in statsmodels.api



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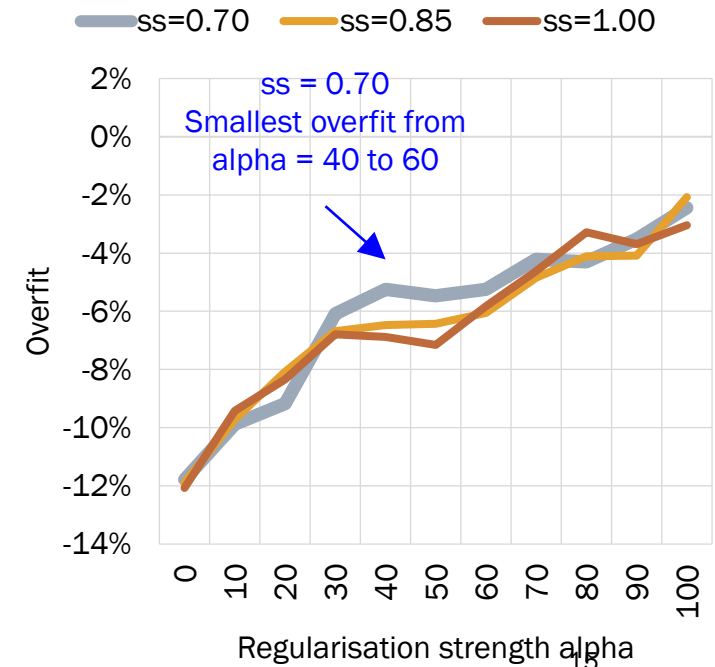
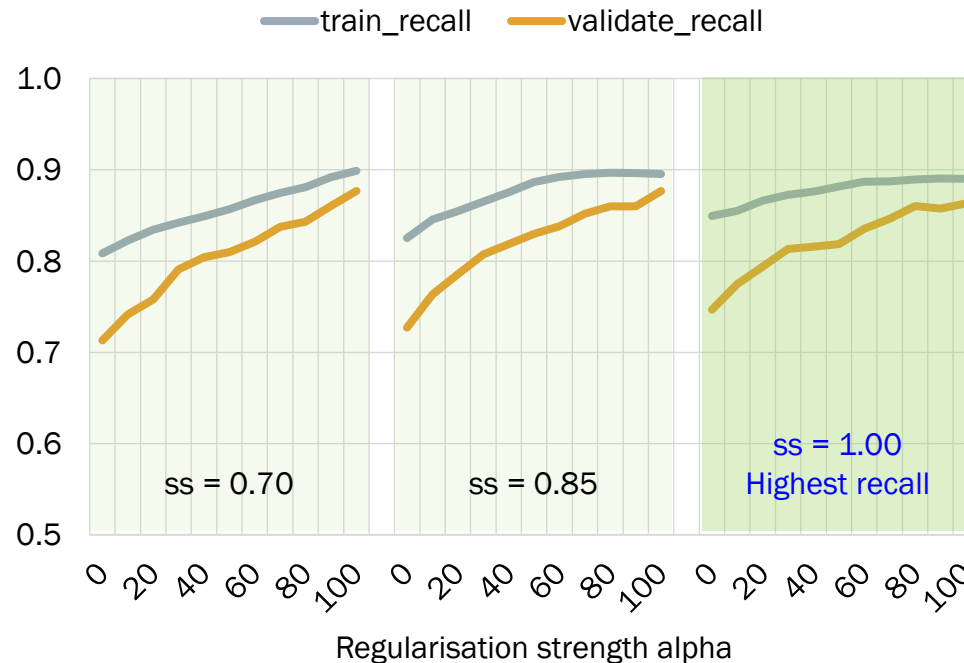
- Model performance



# Data Science Process

- Objective
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- **Build model**
  - **Logistic Regression**
  - K Nearest Neighbours
  - Neural Network
- Conclude on objective
- Deployment workflow

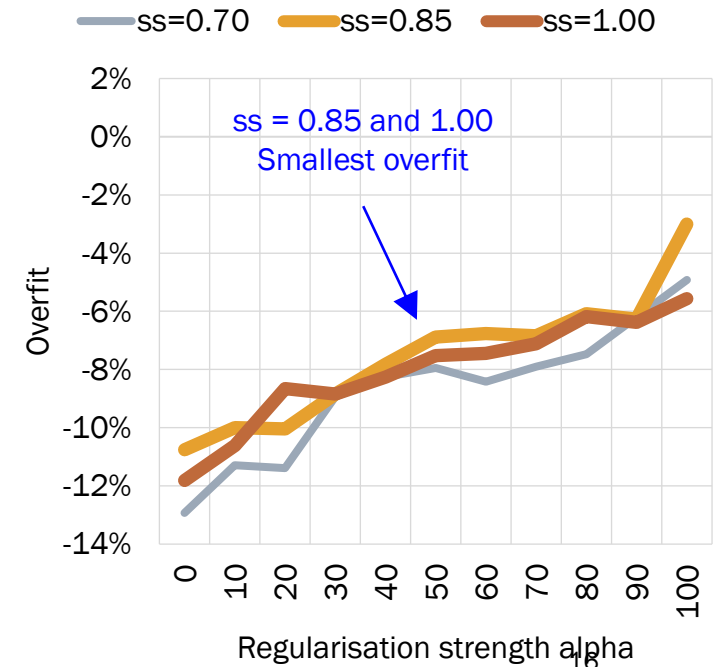
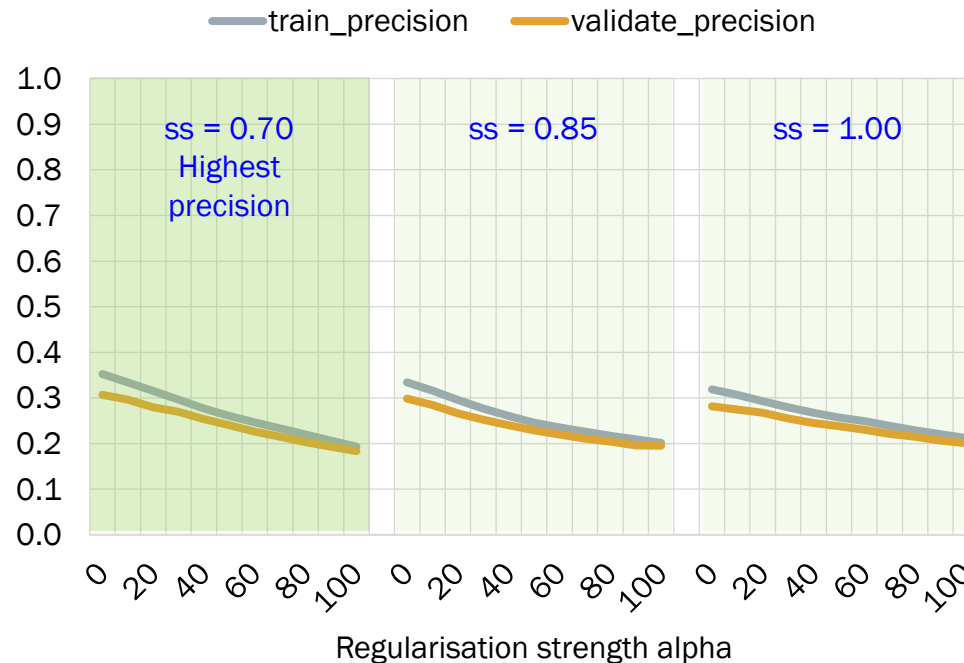
- Logistic Regression with Regularisation (statsmodels.api)



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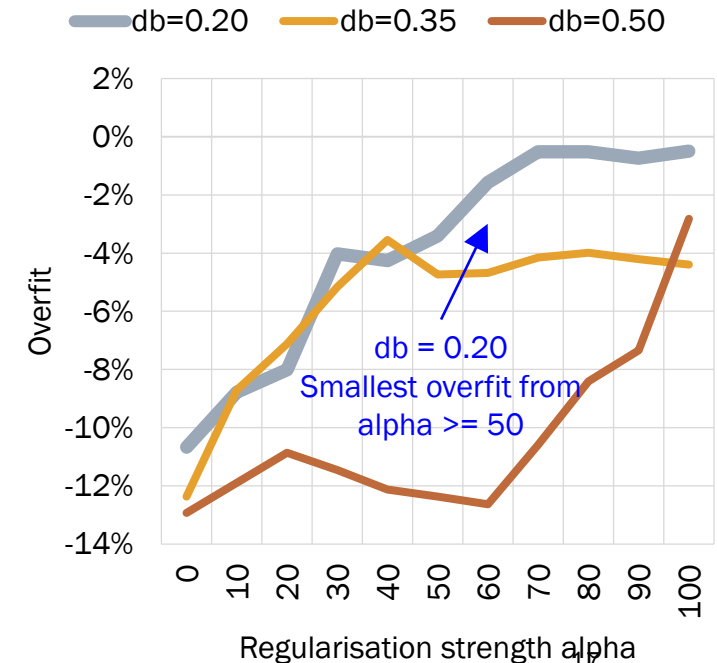
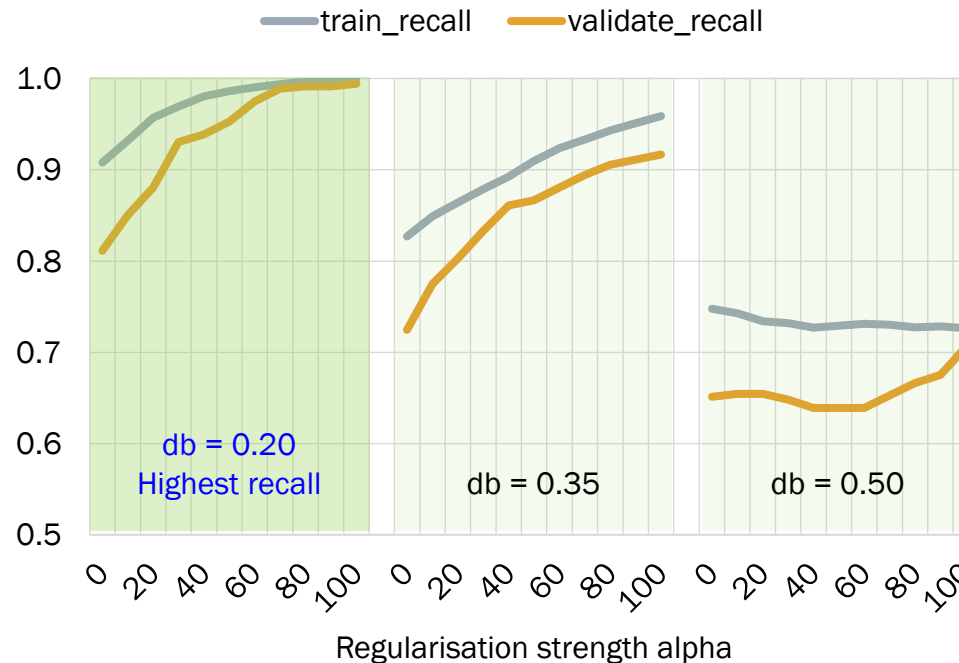


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Impute	Oversample	Scale	Model	Decision Boundary
<ul style="list-style-type: none"><li>• KNNImputer</li><li>• Neighbouring samples = 5</li></ul>	<ul style="list-style-type: none"><li>• SMOTE</li><li>• Sampling strategy = 0.70, 0.85, 1.00</li></ul>	<ul style="list-style-type: none"><li>• StandardScaler</li></ul>	<ul style="list-style-type: none"><li>• Regularisation strength</li><li>• L1 Alpha = 0 to 100</li></ul>	<ul style="list-style-type: none"><li>• Probability &gt; 0.50, 0.35, 0.20 <math>\Rightarrow</math> Category = 1 (Die)</li></ul>

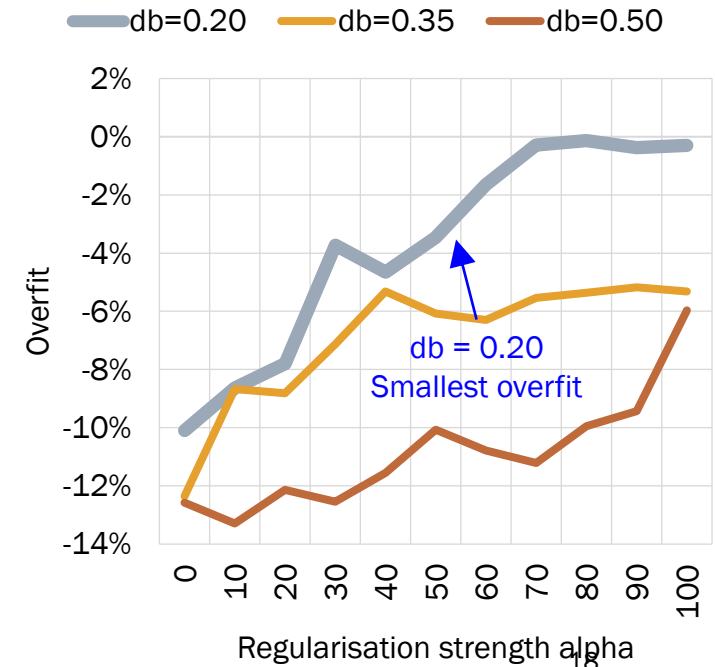
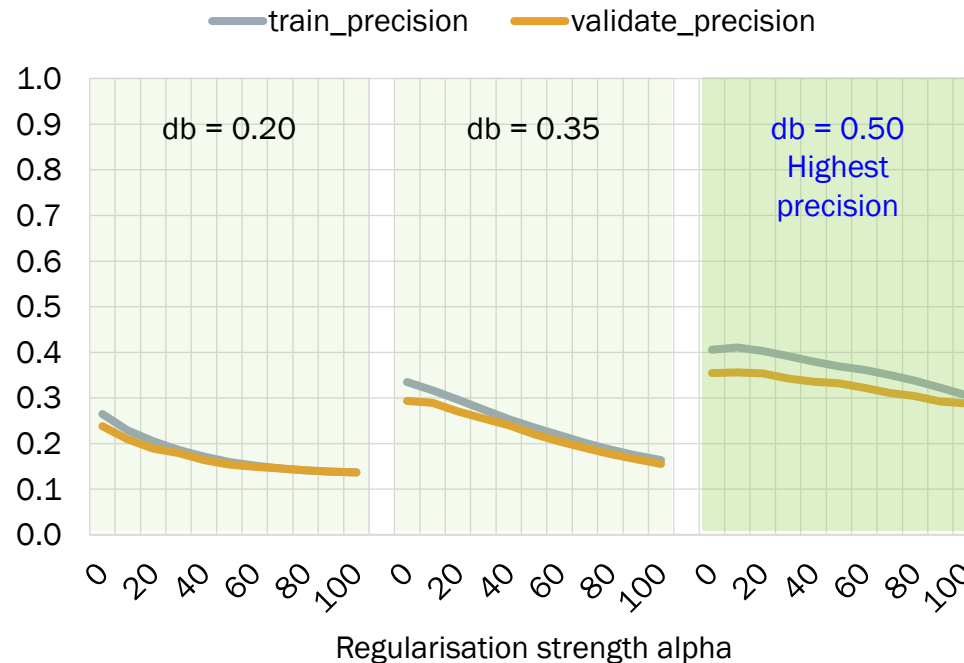


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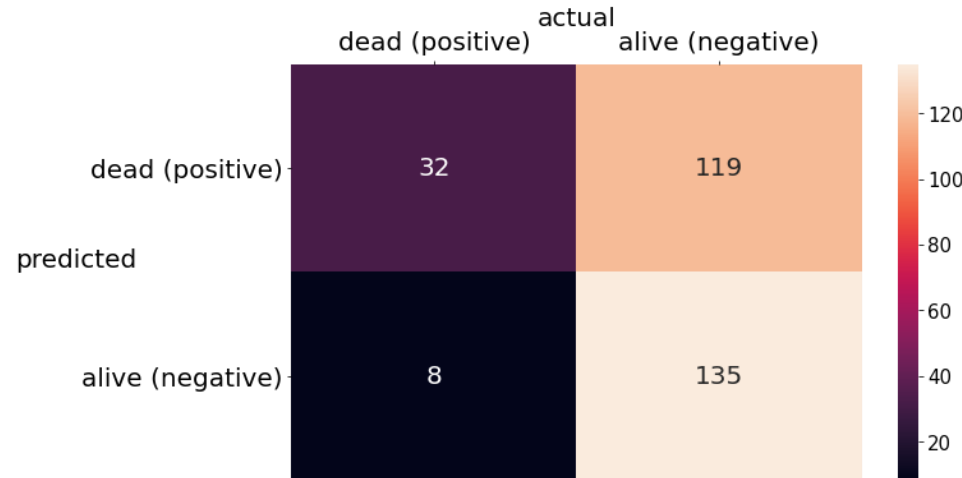


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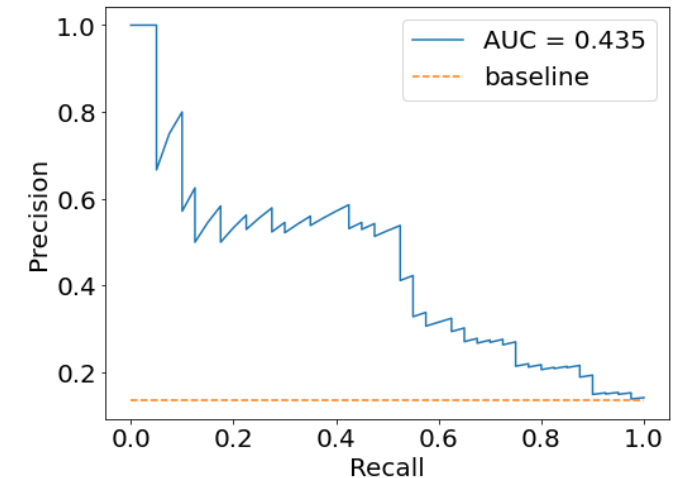
- Logistic Regression with Regularisation (statsmodels.api)

Impute	Oversample	Scale	Model	Decision Boundary
<ul style="list-style-type: none"><li>• KNNImputer</li><li>• Neighbouring samples = 5</li></ul>	<ul style="list-style-type: none"><li>• SMOTE</li><li>• Sampling strategy = 0.85 (smallest overfit)</li></ul>	<ul style="list-style-type: none"><li>• StandardScaler</li></ul>	<ul style="list-style-type: none"><li>• Regularisation strength</li><li>• L1 Alpha = 50 (simpler model but not too simple)</li></ul>	<ul style="list-style-type: none"><li>• Probability &gt; 0.35 <math>\Rightarrow</math> Category = 1 (Die) (balance overfit, recall, precision)</li></ul>



Recall (overfit)  
CV Train = 0.920  
CV Validate = 0.875 (-4.85%)

Recall (overfit)  
Train = 0.915  
Test = 0.800 (-12.7%)



Precision (overfit)  
Train = 0.226  
Test = 0.212 (-6.09%)

# Data Science Process

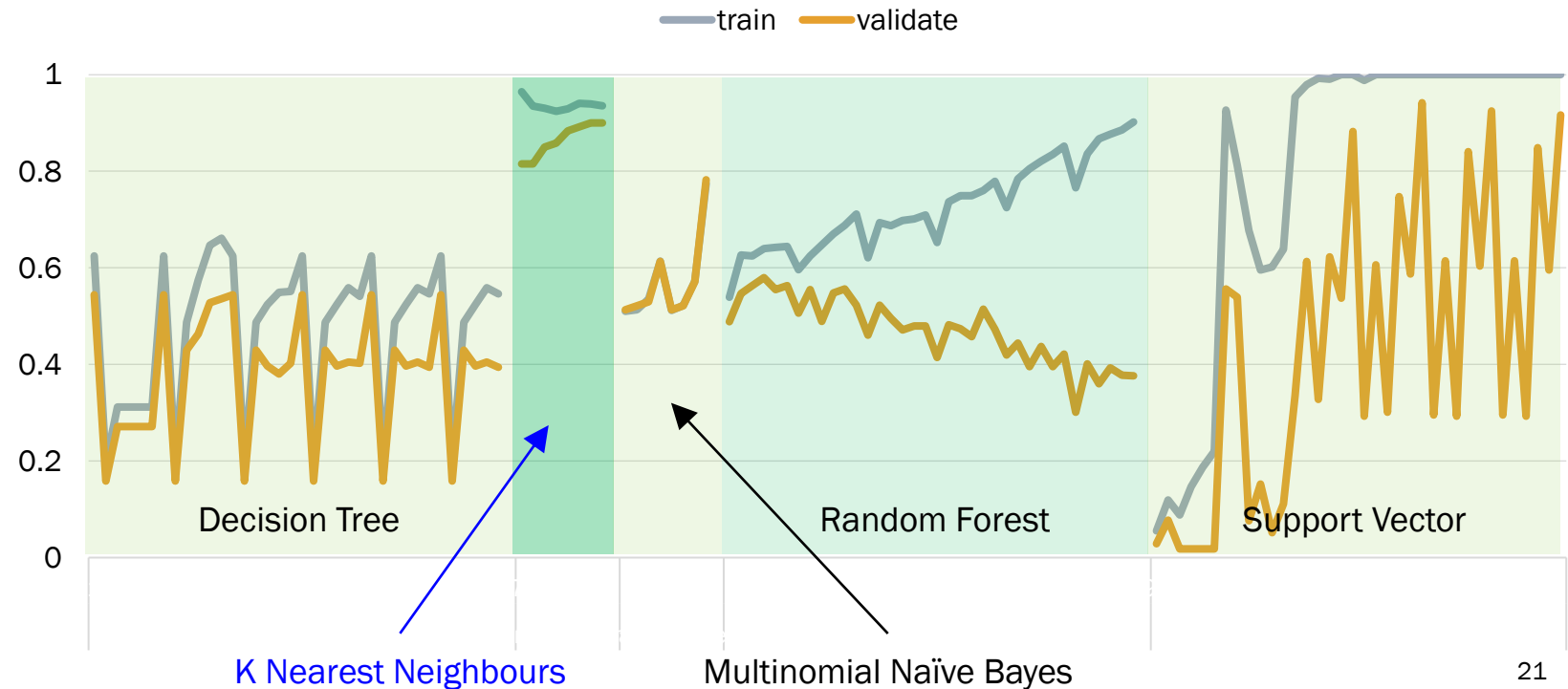
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- Pipeline (train=fit\_transform, validate=transform)
  - KNNImpute
  - SMOTE
  - StandardScaler
  - **Model**
    - Decision Tree
    - Random Forest
    - Multinomial Naïve Bayes
    - K Nearest Neighbours
    - Support Vector Machine
- Hyperparameter search settings
- GridSearchCV (return train score, scoring = recall, 15-fold)
- Fit
- CV results
  - Train recall
  - Validate recall

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- Model
  - Decision Tree
  - Random Forest
  - Multinomial Naïve Bayes
  - **K Nearest Neighbours (best balanced performance in recall and overfit)**
  - Support Vector Machine



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## • K Nearest Neighbours

Impute	Oversample	Scale	Model	Decision Boundary
<ul style="list-style-type: none"> <li>• KNNImputer</li> <li>• Neighbouring samples = 5</li> </ul>	<ul style="list-style-type: none"> <li>• SMOTE</li> <li>• Sampling strategy = 0.70, 0.85, 1.00</li> </ul>	<ul style="list-style-type: none"> <li>• StandardScaler</li> </ul>	<ul style="list-style-type: none"> <li>• Nearest neighbours = 100, 150, 200</li> <li>• Weight = Uniform, Distance</li> <li>• Algorithm = Auto, Brute</li> <li>• P = 1, 2</li> </ul>	<ul style="list-style-type: none"> <li>• Default in K Nearest Neighbours</li> </ul>

Over	Train	Valid	Over	NN	Train	Valid	Over
0.70	0.863	0.698	-19.14%	100	0.908	0.774	-14.76%
0.85	0.915	0.792	-13.42%	150	0.907	0.780	-13.99%
1.00	0.941	0.848	-9.98%	200	0.905	0.784	-13.37%

p	Train	Valid	Over	Weight	Train	Valid	Over	Algo	Train	Valid	Over
p=1	0.865	0.688	-20.49%	Dist	1.000	0.781	-21.87%	auto	0.907	0.779	-14.04%
p=2	0.948	0.871	-8.17%	Unif	0.813	0.777	-4.42%	brute	0.907	0.779	-14.04%

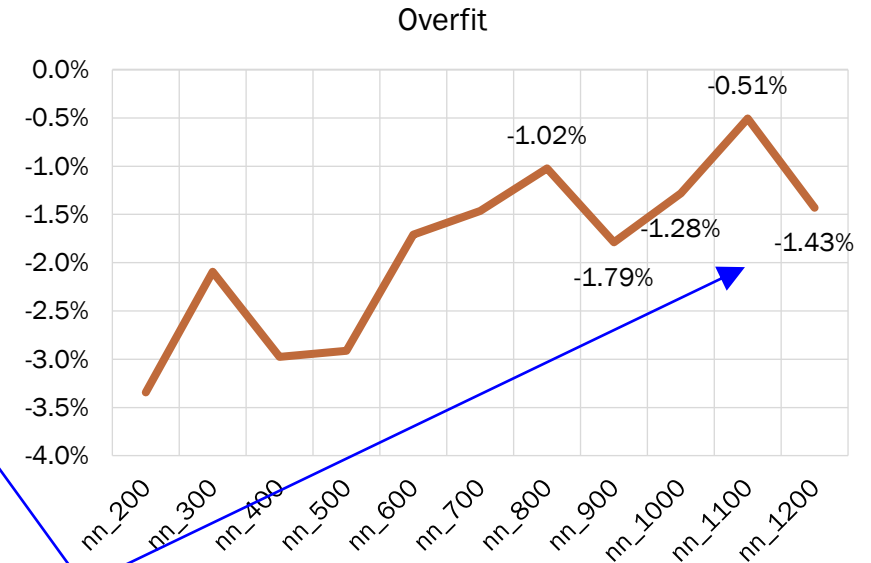
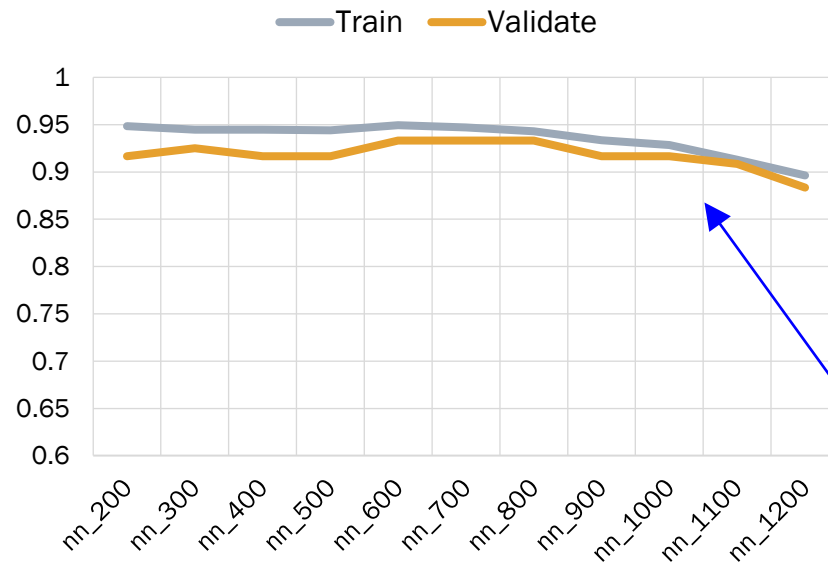
Parameter and hyperparameters with smallest overfit indicated in blue

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## • K Nearest Neighbours

Impute	Oversample	Scale	Model	Decision Boundary
<ul style="list-style-type: none"><li>• KNNImputer</li><li>• Neighbouring samples = 5</li></ul>	<ul style="list-style-type: none"><li>• SMOTE</li><li>• Sampling strategy = 1.00 (smallest overfit)</li></ul>	<ul style="list-style-type: none"><li>• StandardScaler</li></ul>	<ul style="list-style-type: none"><li>• Nearest neighbours = 200 to 1200 (to refine further)</li><li>• Weight = Uniform (smallest overfit)</li><li>• Algorithm = Brute</li><li>• P = 2 (smallest overfit)</li></ul>	<ul style="list-style-type: none"><li>• Default in K Nearest Neighbours</li></ul>



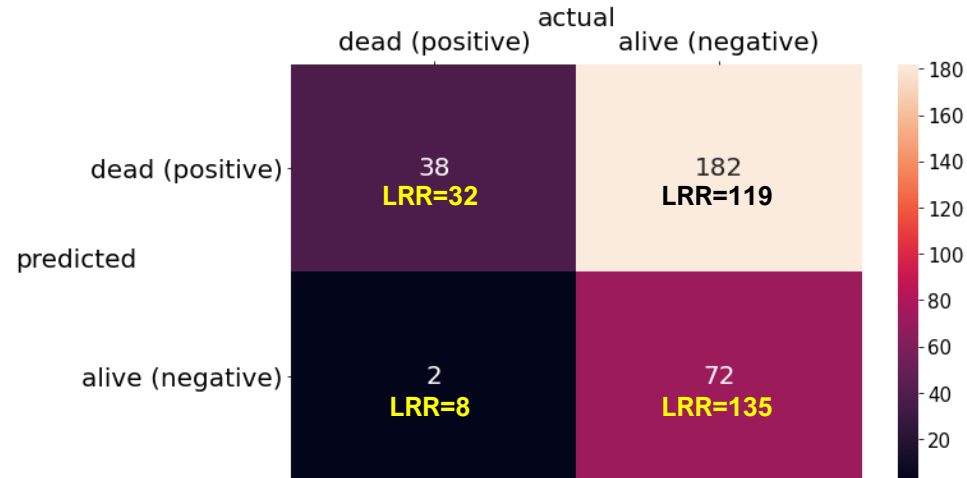
More nearest neighbours, lower performance and smaller overfit

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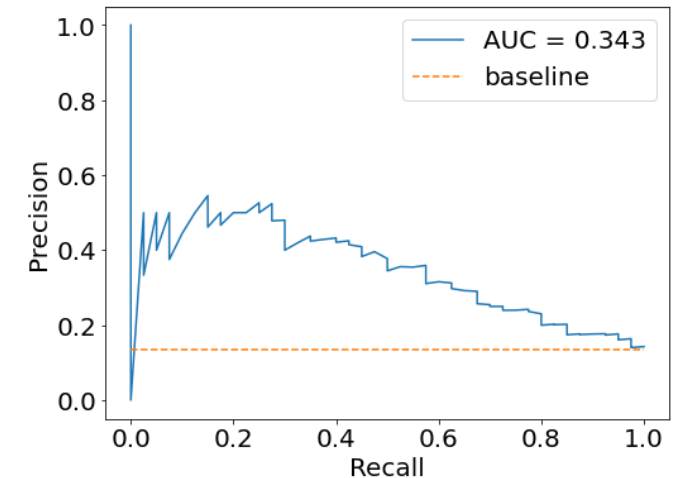
## • K Nearest Neighbours

Impute	Oversample	Scale	Model	Decision Boundary
<ul style="list-style-type: none"><li>• KNNImputer</li><li>• Neighbouring samples = 5</li></ul>	<ul style="list-style-type: none"><li>• SMOTE</li><li>• Sampling strategy = 1.00 (smallest overfit)</li></ul>	<ul style="list-style-type: none"><li>• StandardScaler</li></ul>	<ul style="list-style-type: none"><li>• Nearest neighbours = 1000 (smallest overfit)</li><li>• Weight = Uniform (smallest overfit)</li><li>• Algorithm = Brute</li><li>• P = 2 (smallest overfit)</li></ul>	<ul style="list-style-type: none"><li>• Default in K Nearest Neighbours</li></ul>



Recall (overfit)  
CV Train = 0.929  
CV Validate = 0.917 (-1.29%)

Recall (overfit)  
Train = 0.933  
Test = 0.950 (1.85%)



Precision (overfit)  
Train = 0.171  
Test = 0.173 (1.15%)

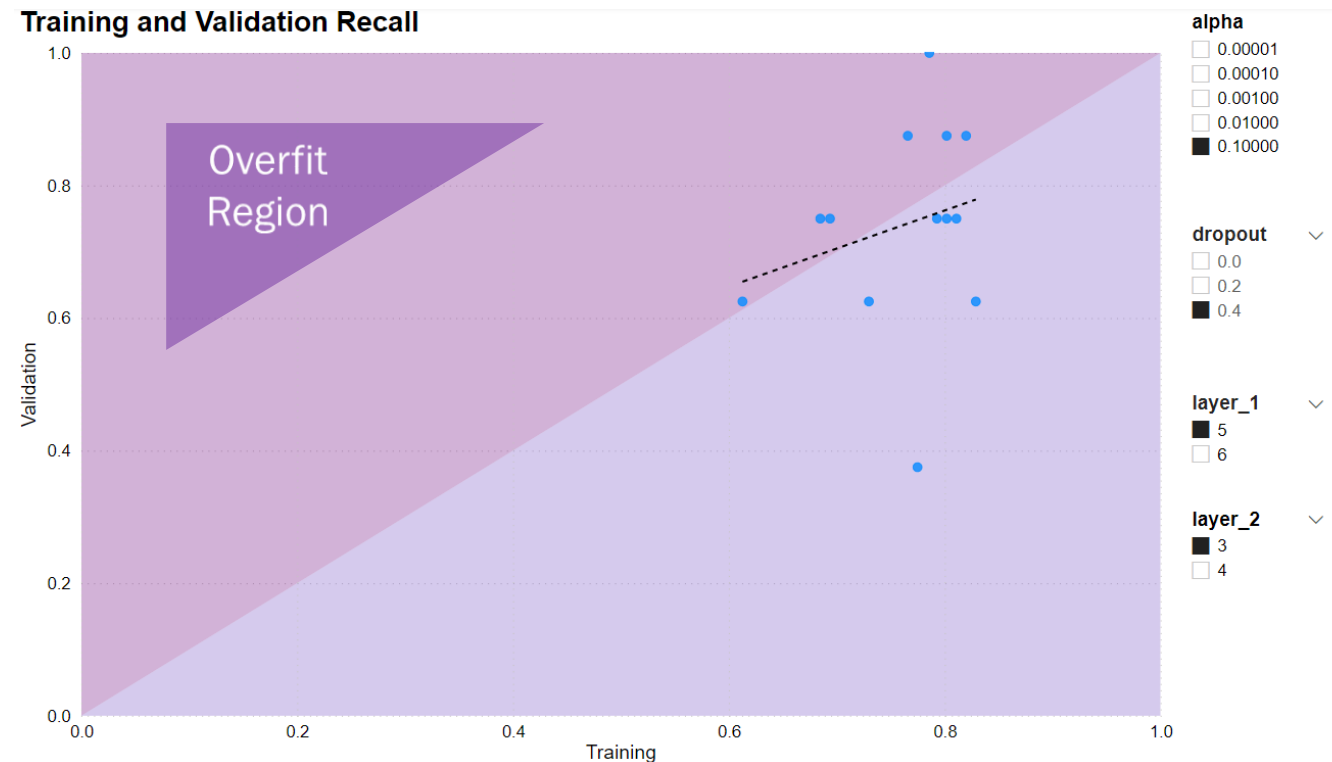


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- Neural Network with Regularisation, Dropout Rate, Early Stopping

Impute	Oversample	Scale	Model	Decision Boundary
<ul style="list-style-type: none"><li>• KNNImputer</li><li>• Neighbouring samples = 5</li></ul>	<ul style="list-style-type: none"><li>• SMOTE</li><li>• Sampling strategy = 1.00</li></ul>	<ul style="list-style-type: none"><li>• StandardScaler</li></ul>	<ul style="list-style-type: none"><li>• Layer 1 = 5, 6</li><li>• Layer 2 = 3, 4</li><li>• Dropout = 0.0, 0.2, 0.4</li><li>• L1 Alpha = 0.1, 0.01, 0.001, 0.0001</li></ul>	<ul style="list-style-type: none"><li>• 0.4</li></ul>



Visualisation with  
Microsoft Power BI

Good balance of  
\* slope = 1  
\* small overfit

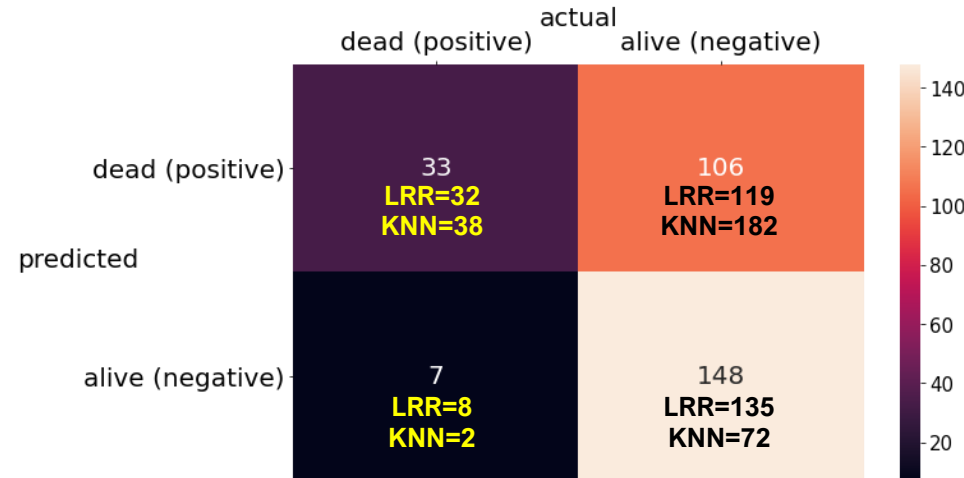
Best hyperparameters  
with  
\* layer1 = 5 neurons  
\* layer2 = 3 neurons  
\* dropout = 0.4  
\* L1 alpha = 0.1

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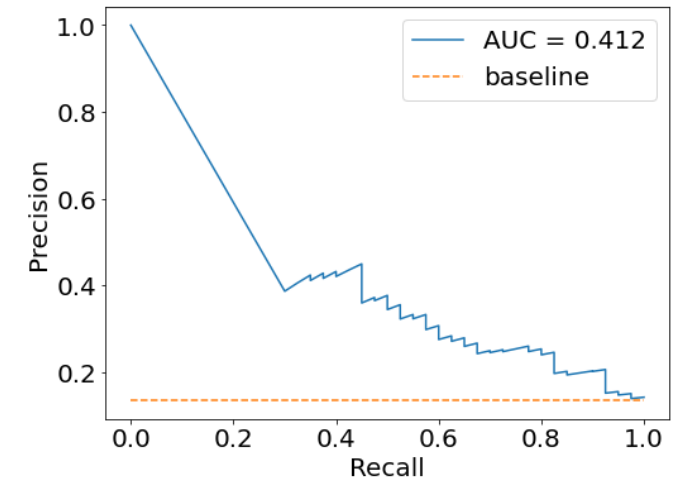
- Neural Network with Regularisation, Dropout Rate, Early Stopping

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Recall (overfit)  
CV Train = 0.769  
CV Validate = 0.742 (-3.54%)

Recall (overfit)  
Train = 0.924  
Test = 0.825 (-10.8%)



Precision (overfit)  
Train = 0.264  
Test = 0.237 (-10.0%)

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- **Logistic Regression using statistical significance**

feature	std_coef	z	p
renal_failure	-0.3382	-4.851	0
bicarbonate	-0.2817	-3.114	0.002
blood_calcium	-0.2525	-3.602	0.000
urine_output	-0.2168	-3.133	0.002
deficiencyanemias	-0.1804	-2.871	0.004
sp_o2	-0.1459	-2.245	0.025
heart_rate	0.1387	2.031	0.042
leucocyte	0.1719	2.298	0.022
atrialfibrillation	0.1725	2.785	0.005
urea_nitrogen	0.3728	4.485	0.000

renal\_failure:

- Categorical, Negative
- Presence will reduce odds of dying compared to absence

bicarbonate:

- Numeric, Negative
- Higher value will reduce odds of dying

heart\_rate:

- Numeric, Positive
- Higher value will increase odds of dying

atrialfibrillation:

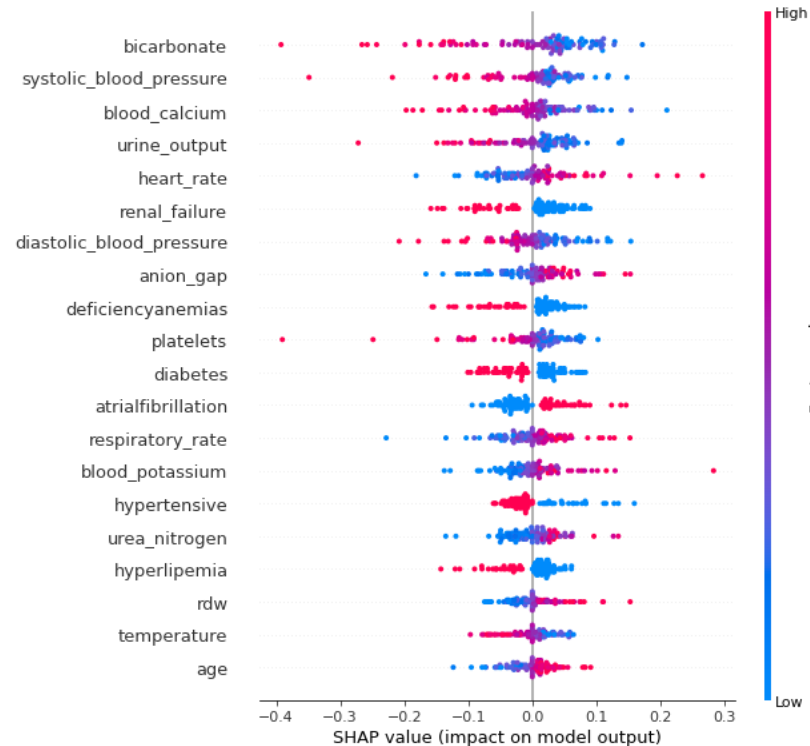
- Categorical, Positive
- Presence will increase odds of dying compared to absence

- Interpretable
  - Causes of death are understood

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- **K Nearest Neighbours using Shapley values from SHAP**



bicarbonate (in agreement to before)

- Numeric
- Higher value (red) will reduce odds of dying (negative SHAP values)

heart\_rate (in agreement to before)

- Numeric
- Higher value (red) will increase odds of dying (positive SHAP values)

renal\_failure (in agreement to before)

- Categorical
- Higher value (presence) will reduce odds of dying (negative SHAP values)

atrialfibrillation (in agreement to before):

- Categorical
- Higher value (presence) will increase odds of dying (positive SHAP values)

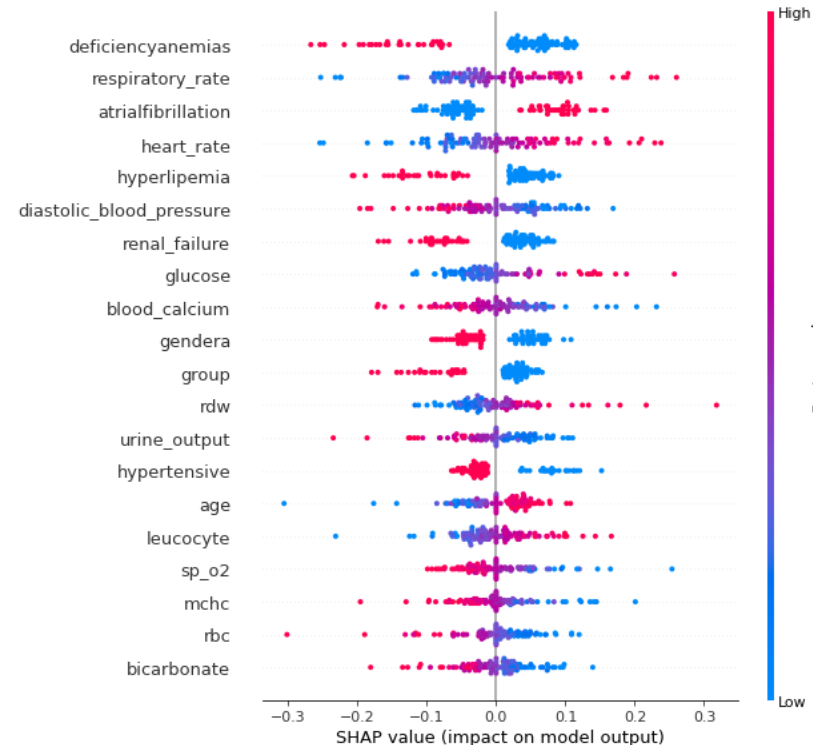
- Interpretable
  - Causes of death are understood

# Data Science Process

- Objective
- Collect
- Explore
- Engineer feature
- Build model
  - Logistic Regression
  - K Nearest Neighbours
  - Neural Network
- **Conclude on objective**
- Deployment workflow

- Develop interpretable machine learning model to predict and influence mortality of patients with heart failure warded in intensive care unit

- **Neural Network using Shapley values from SHAP**



bicarbonate (in agreement to before)

- Numeric
- Higher value (red) will reduce odds of dying (negative SHAP values)

heart\_rate (in agreement to before)

- Numeric
- Higher value (red) will increase odds of dying (positive SHAP values)

renal\_failure (in agreement to before)

- Categorical
- Higher value (presence) will reduce odds of dying (negative SHAP values)

atrialfibrillation (in agreement to before)

- Categorical
- Higher value (presence) will increase odds of dying (positive SHAP values)

- Interpretable
  - Causes of death are understood

## Data Science Process

- Objective
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- Comparison of top important features in 3 machine learning models

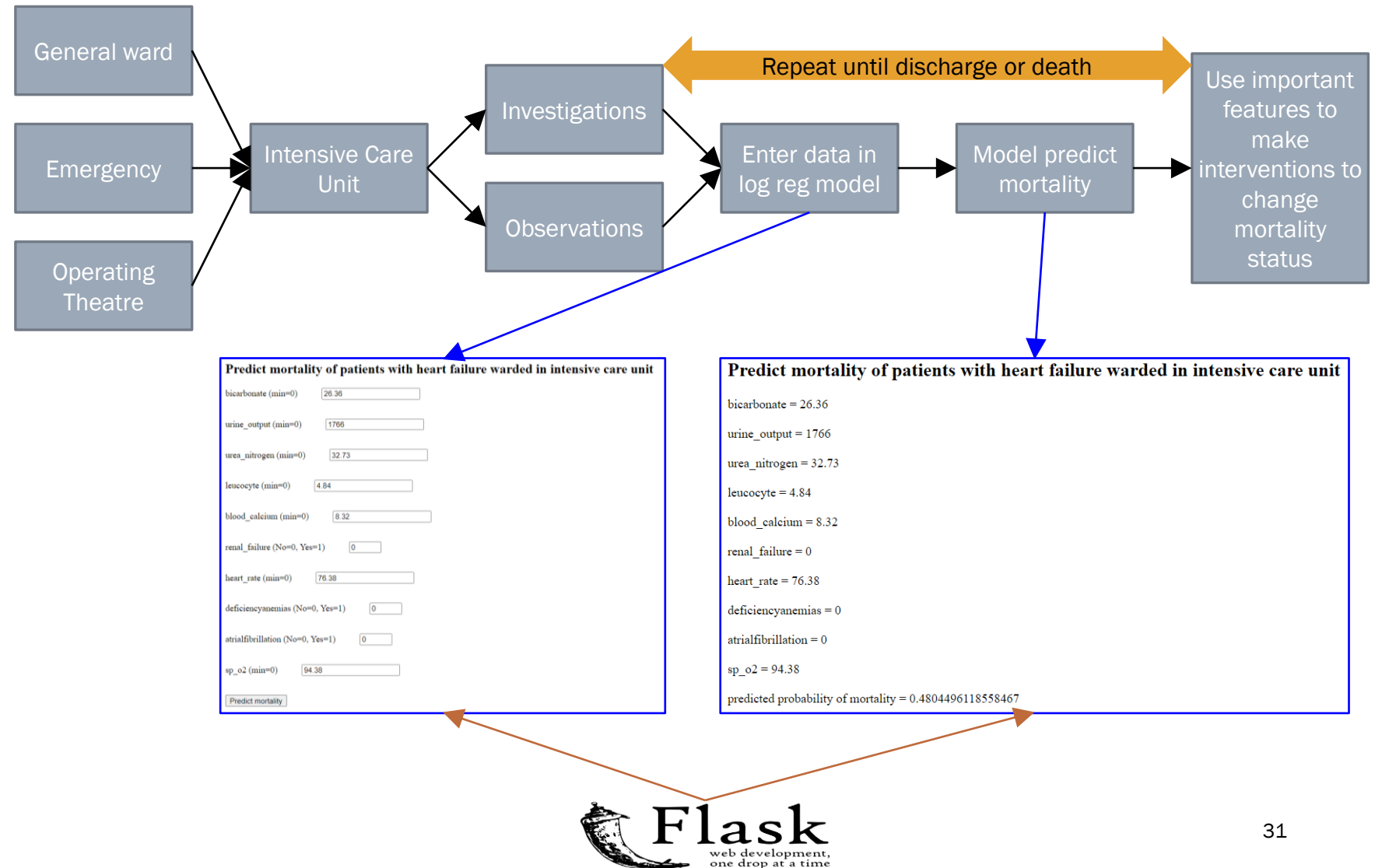
Logistic Regression	K Nearest Neighbours	Neural Network
Renal_failure	Bicarbonate	Deficiencyanemias
Bicarbonate	Systolic_blood_pressure	Respiratory_rate
Blood_calcium	Blood_calcium	Atrialfibrillation
Urine_output	Urine_output	Heart_rate
Deficiencyanemias	Heart_rate	Hyperlipemia
Sp_o2	Renal_failure	Diastolic_blood_pressure
Heart_rate	Diastolic_blood_pressure	Renal_failure
Leucocyte	Anion_gap	Glucose
Atrialfibrillation	Deficiencyanemias	Blood_calcium
Urea_nitrogen	Platelets	Gendera

- BLUE: Presence in 3 models

# Data Science Process

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  - Neural Network
- Conclude on objective
- **Deployment workflow**

- Workflow to execute data science solution:



## Data Science Process

- Objective
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THE END