



Group 12: HCC Survival

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Challenge

Predict whether a Patient will survive the diagnosed liver cancer.

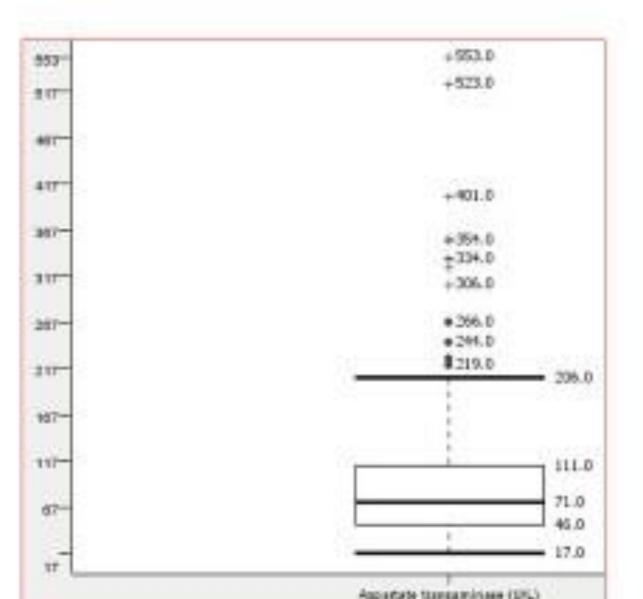
Class 0: patient Dies Class 1: patient Survives

Dataset

- Contains data of 165 real patients diagnosed with HCC(liver Cancer)
- · 49 features + 1 Class Attribute
- Nominal features: 23
 Ordinal features: 3
 Continuous features: 23

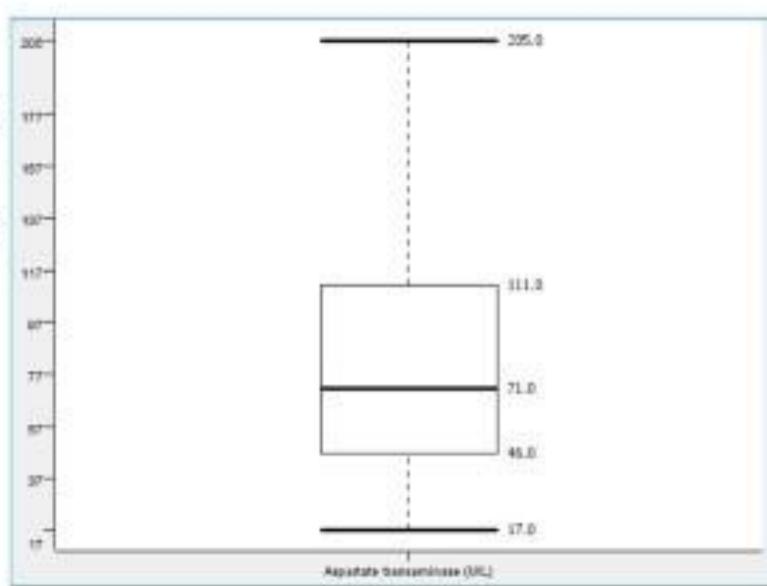
Data Understanding

- Converting Nominal and Ordinal Features to Strings
- Out of 165, only 8 patients have complete information.
- Observed Class Imbalance-Data Skewed towards Class 1
- Positive Correlation between
 Total and Direct Bilirubin(mg/dL)
- Outliers detected using Box plots



Data Preparation

- Missing Values were replaced.
- Correlations were filtered using a threshold of 0.8
- Treated Numerical Outliers with Closest Permitted Values.



Modeling

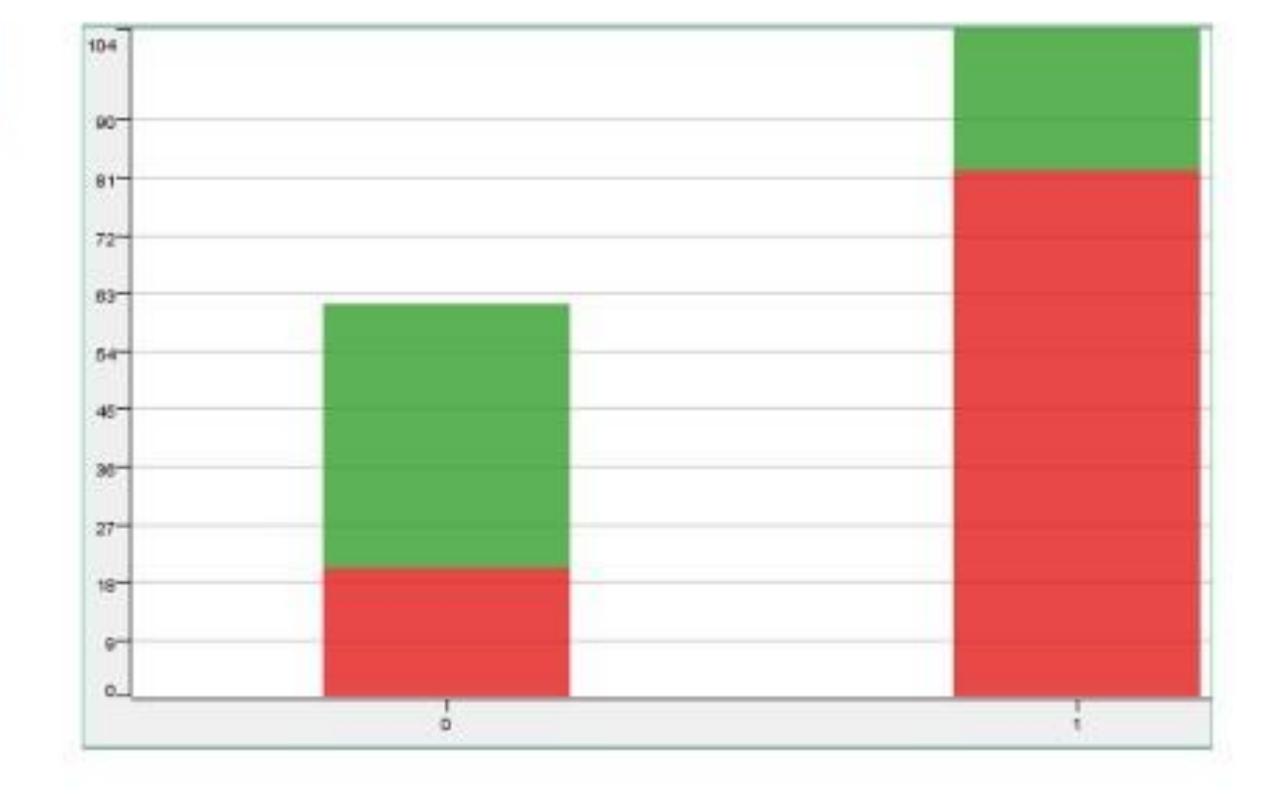
- Class Imbalance was fixed using SMOTE node (minority-Class 0 oversampled)
- X- Partitioner with Leave one- out was used to generate Training and Test data
- Classification Models Tested:-
 - Random Forest
 - Decision Tree
 - K- Nearest Neighbour

Evaluation

- Comparison of Models based on Minimization of False Positives(Diseased patient predicted as "Survived") and Better Accuracy.
- Random Forests excels in both Accuracy and Number of False Positive Predictions.

Results

- Accuracy of Random Forest Model is around 75%
- False Positive Rate is approx. 35%.
- Use of SMOTE reduces the skewed training of the model and increases accuracy of prediction.



Knime-Workflow

