

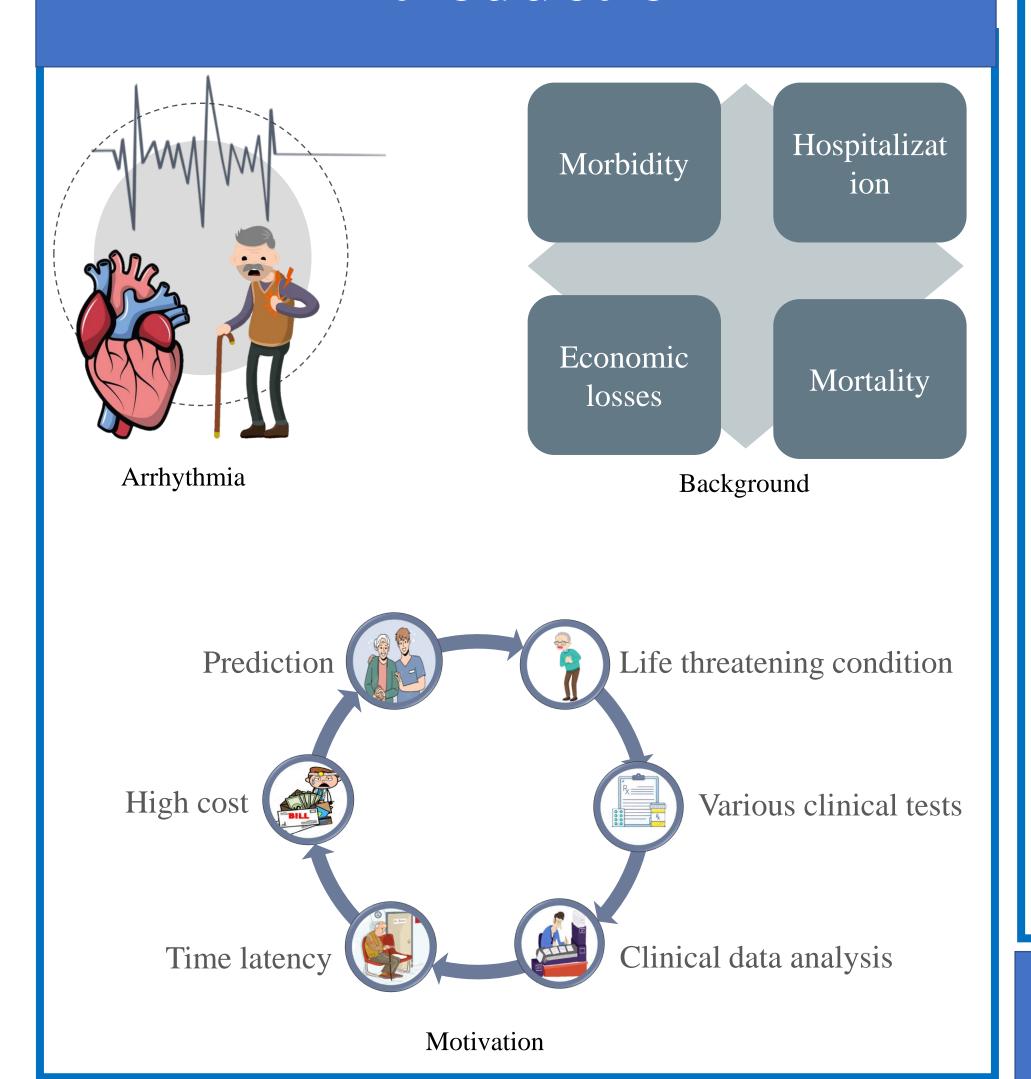
Prediction of Arrhythmia after Acute Myocardial Infarction using Machine Learning and Statistical Techniques

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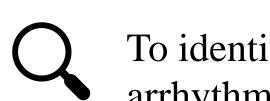
Abstract

Arrhythmia is a life-threatening condition and every year millions of people die because of it. The diagnosis process of conventional health-care system is time consuming and it worsens patients' condition which may lead to death of patients. With the help of conventional statistical and machine learning analysis, we have extracted important predictors so that arrhythmia can be predicted faster and more accurately. The statistical and machine learning techniques have returned the features RA (up and down), BBB, P-R, RA (up and down), heart beats and DD-P of the AMI dataset. With the help of an optimized random forest classifier we have predicted arrhythmia which has shown an accuracy 69.71% with all features and 68.89% with the extracted features. We have also analyzed the time complexity of the model to analyze how quickly our model can predict arrhythmia.

Introduction



Objectives



To identify important features for predicting arrhythmia



To reduce time complexity



To investigate the acute myocardial infraction (AMI) dataset

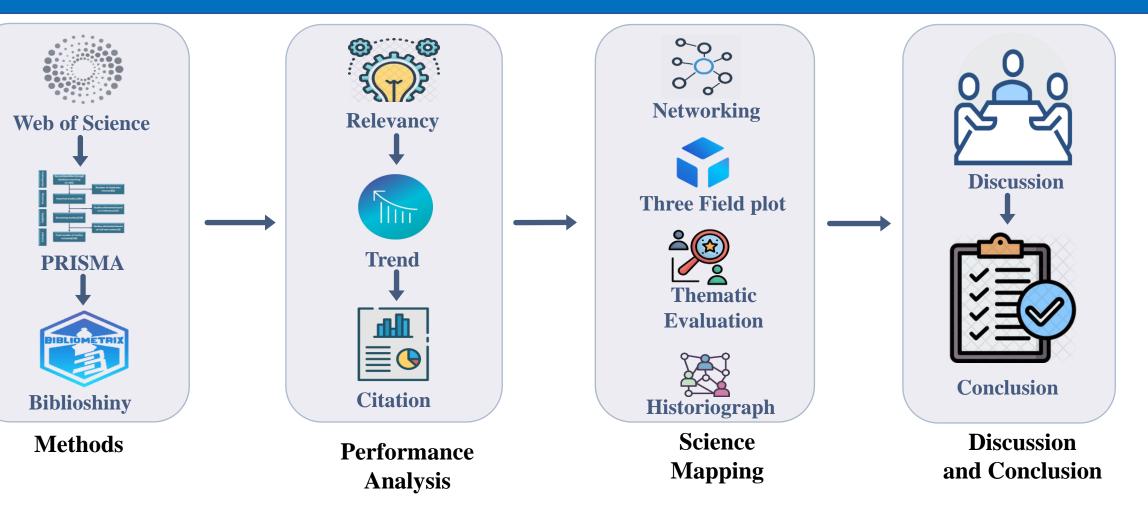


To analyze the model performance

Conclusion & Future Work

- Similar types of features returned from statistical and machine learning analysis
- Improved accuracy than the previous work [2]
- Model will be trained with more datasets
- Neural network will be applied to improve accuracy
- Real-time server-based patient monitoring system will be developed where this model will predict arrhythmia

Literature Review



Methodology of Bibliometric Analysis



 Imported studies (285)

Studies eliminated based on irrelevancy (47)

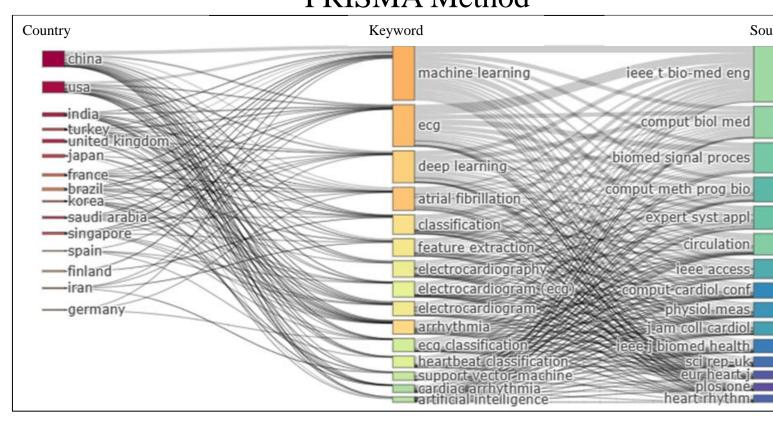
Remaining studies (238)

Studies eliminated based on full text review (0)

Final number of studies included (238)

PRISMA Method

Record identifies through

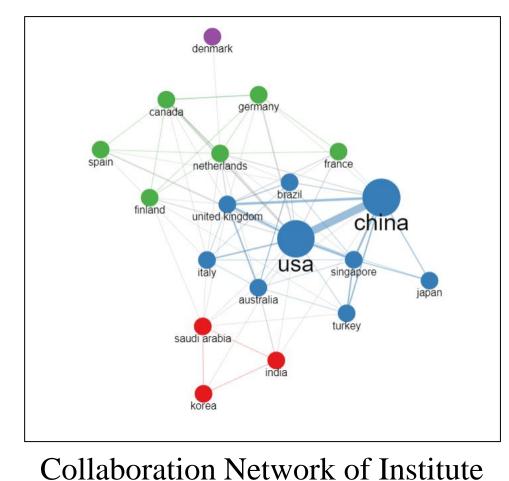


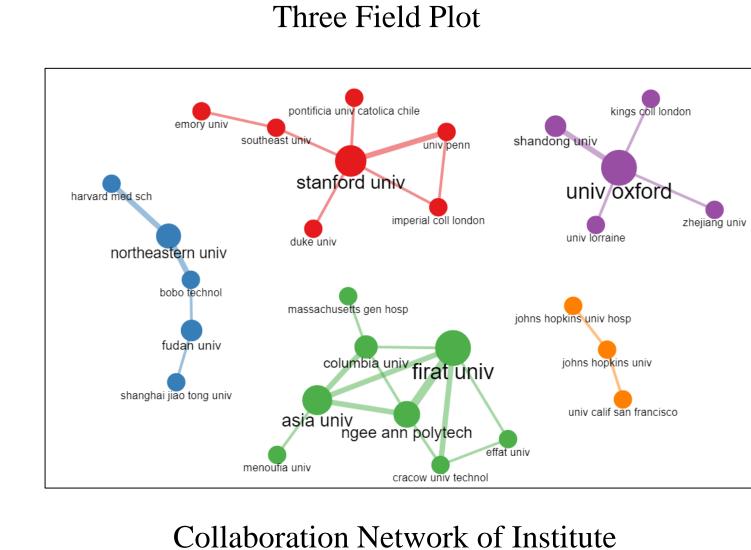
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sensors-basel
inform sciences
expert syst appl
comput biol med
biomed signal proces
biomed signal proces
eaccess
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circulation
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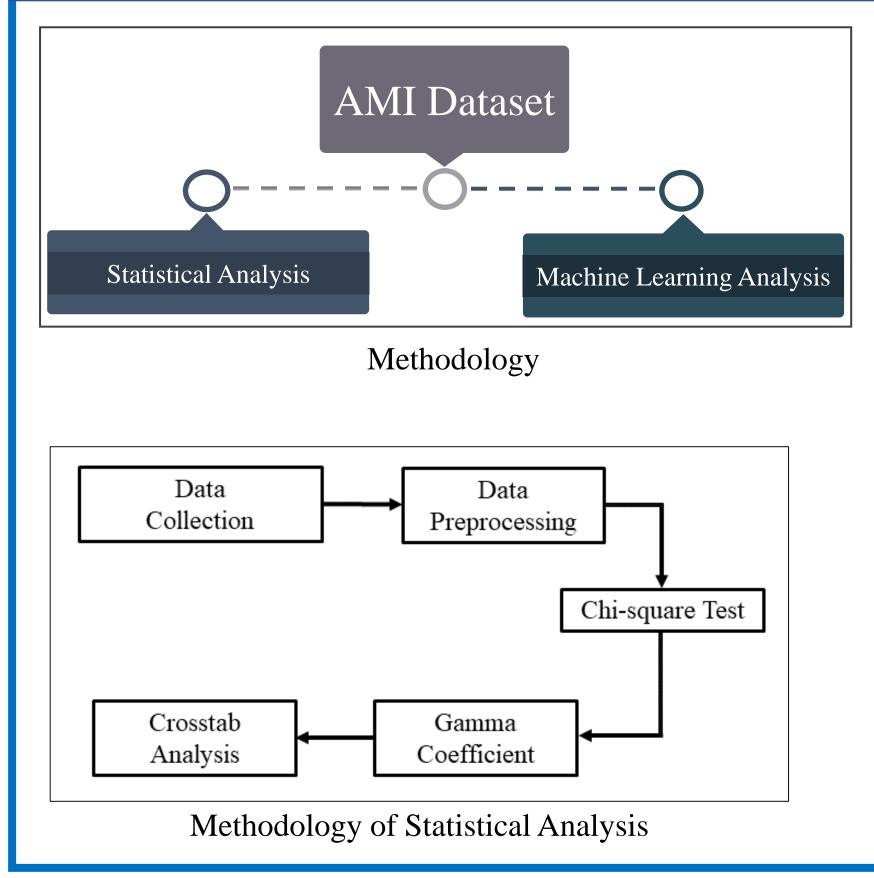
Co-citation Network of Journal

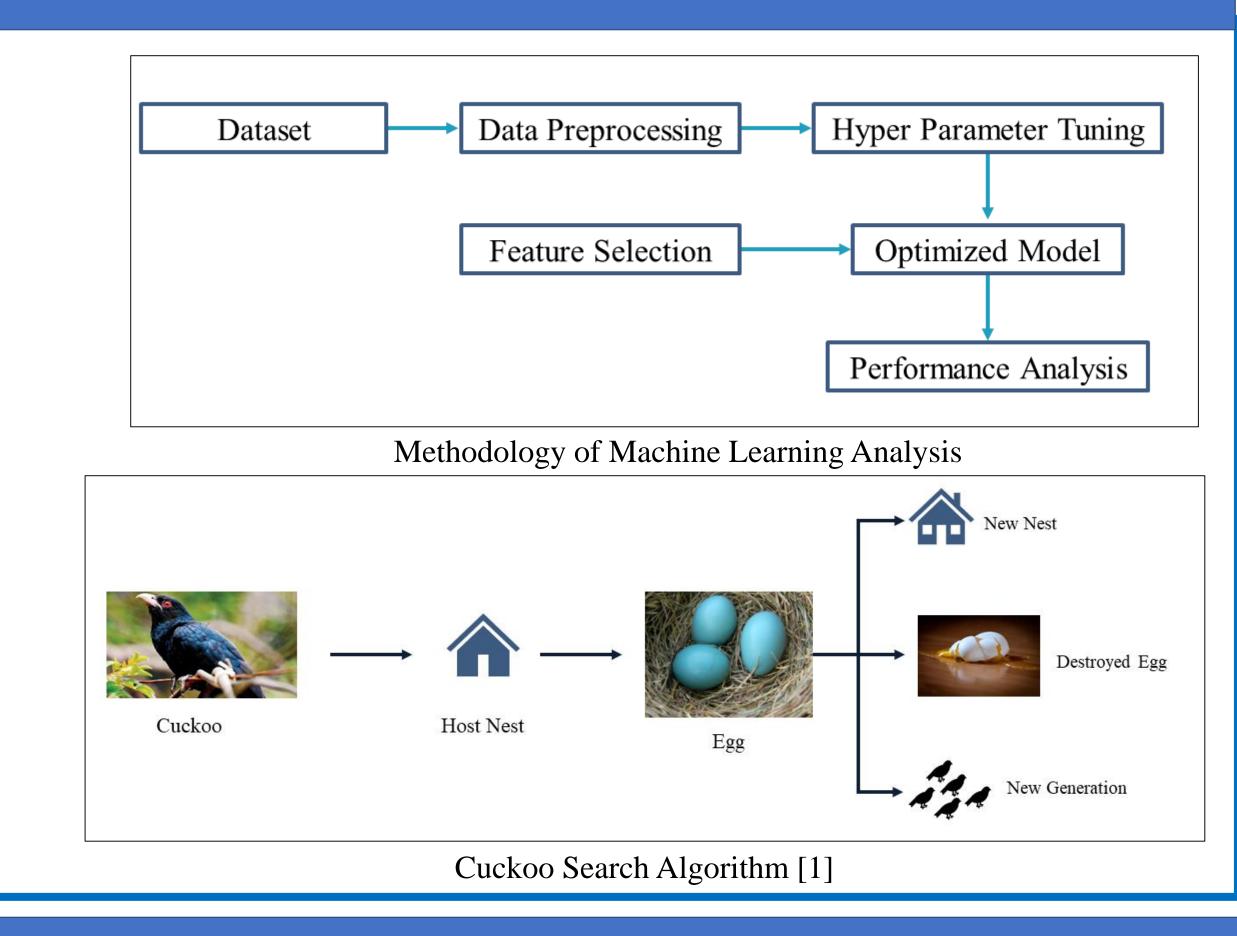
Collabora



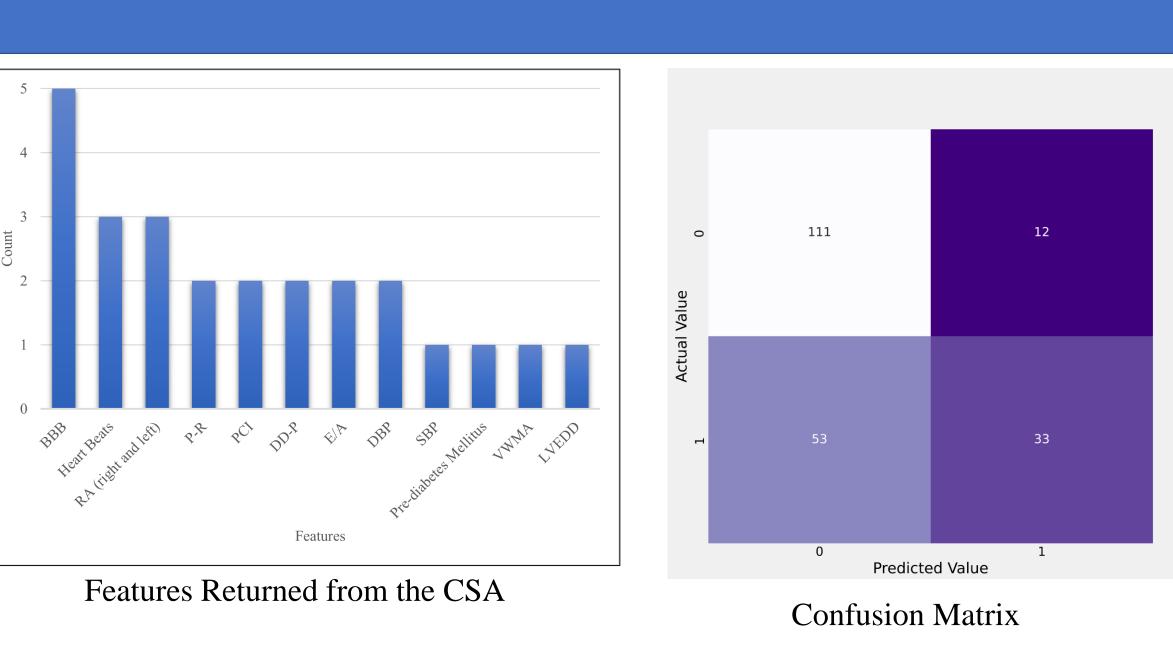


Methodology





Result



1.0 AUC=0.6430799773114011

0.8
0.6
0.0
0.0
0.0
0.2
0.4
0.6
0.8
1.0

ROC Curve

Features Returned from Statistical Analysis: RA (right and left), BBB, P-R and RA (up and down)

References