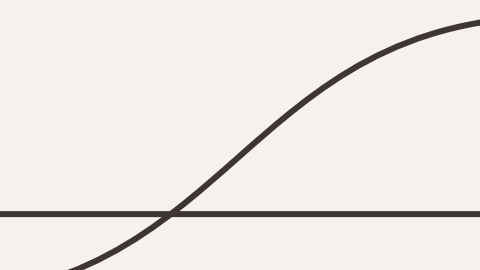


# Context-Aware Weighted KNN with Locally Adaptive $k$

Anieesh Saravanan

03/28/25



You know that feeling when you  
meet someone and your heart skips  
a beat?

Yeah, that's arrhythmia. You can die  
from that.

<https://rusafu.com/quotes/tag/arrhythmia/>

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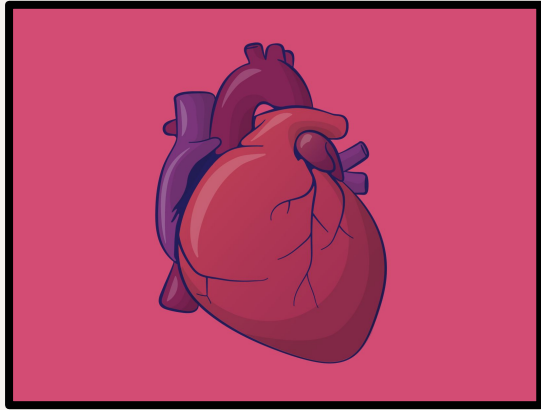
Configuring Parameters  
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Results and Future Work

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# 00

## Introduction

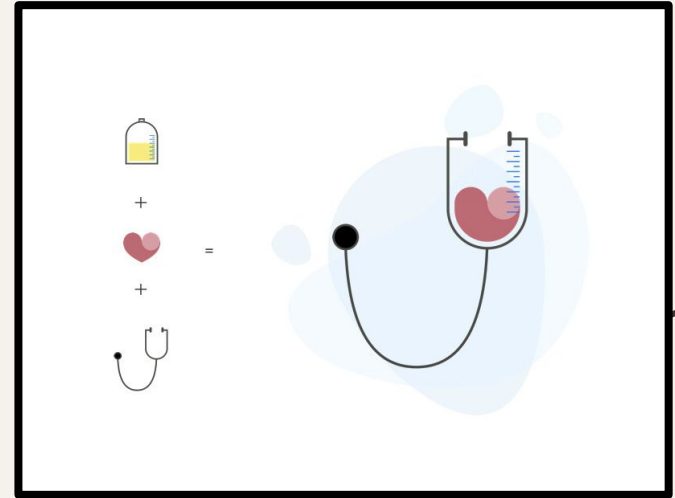
# Project Goal

- Develop a binary classification model for arrhythmia
- Diagnose patients with arrhythmia based on patient biometrics and ECG readings
- Streamline diagnosis process with supplementary outlook of data



# Arrhythmia Overview

- Arrhythmia refers to irregular heartbeats
- Antiarrhythmic medications/therapies exist
- No simple guidelines for diagnosis
- Widely varying clinical manifestations and minimal warning signs
- Even experienced cardiologists may misinterpret ECG patterns



# Common Myths

**Myth:** Arrhythmia always presents with severe, dramatic symptoms

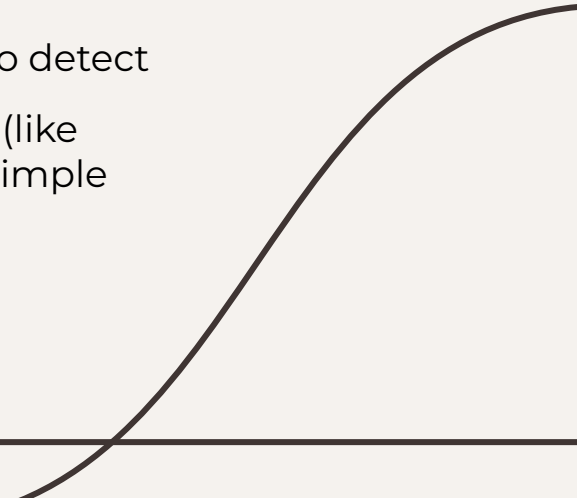
**Fact:** Many arrhythmias can be **subtle or even asymptomatic**

**Myth:** Only unhealthy/elderly individuals develop arrhythmia

**Fact:** Arrhythmia can occur in healthy individuals (without heart conditions)

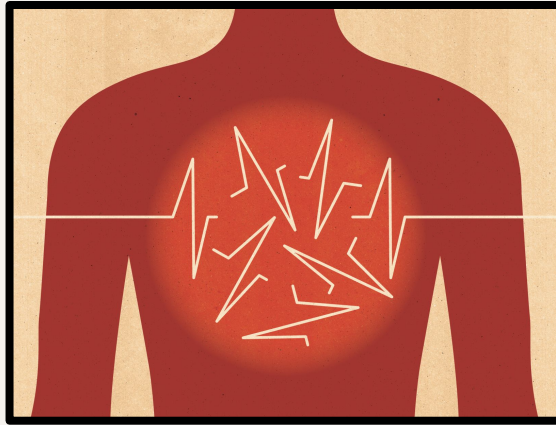
**Myth:** Abnormal heart rhythms are always obvious and easy to detect

**Fact:** Many arrhythmias require **specialized diagnostic tools** (like extended ECG monitoring) that may not be apparent with a simple pulse check



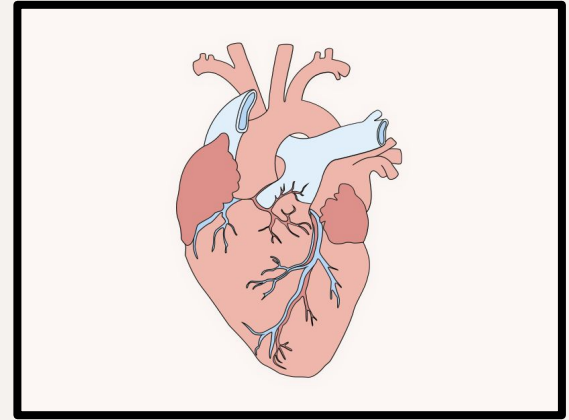
# Impact

In 2021, arrhythmia was mentioned on at least **232,030** death certificates



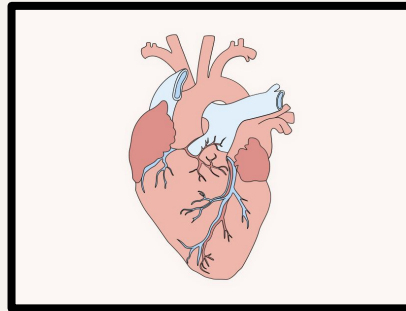


# 01 Project Overview



# Dataset Overview

- **Quantitative/Categorical** dataset with **452 instances** and **279 attributes**
- Class attribute is the diagnosis of the patient (nonarrhythmia or type of arrhythmia)
- Sourced from UCI and originally compiled by H. Altay Guvenir
- Missing values: instances regarding vector angles (ECG electrode similarity)
- Class distribution: 54.20% nonarrhythmia, 45.80% arrhythmia



# Attributes

1. **class: Class Code 01-16**
  - a. Code 01: Normal
  - b. Code 02: Ischemic changes (Coronary Artery Disease)
  - c. Code 03: Old Anterior Myocardial Infarction
  - d. Code 04: Old Inferior Myocardial Infarction
  - ...
  - p. Code 16: Others
2. age: Age in years, linear
3. sex: Sex (0 = male; 1 = female), nominal
4. height: Height in centimeters, linear
5. weight: Weight in kilograms, linear
6. QRSduration: Average of QRS duration in milliseconds, linear
7. PRinterval: Average duration between onset of P and Q waves in milliseconds, linear
8. Q-Tinterval: Average duration between onset of Q and offset of T waves in msec., linear
9. Tinterval: Average duration of T wave in milliseconds, linear
10. Pinterval: Average duration of P wave in milliseconds, linear
- ...
279. chV6\_QRSTA: Of channel V6

# Related Work

- **Signal Pre-Processing and Fixed-Feature Extraction**

  - Classification of Arrhythmia using KNN-Classifer

  - Heart Arrhythmia Classification Using Machine Learning Algorithms

- **Classifier Comparisons with Feature Selection**

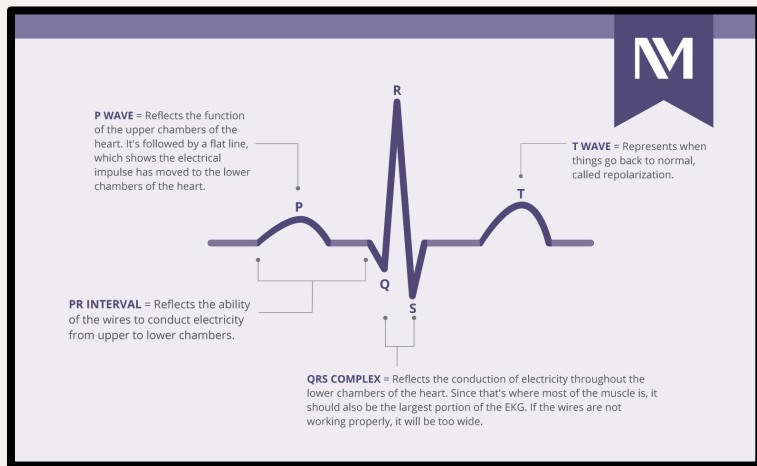
  - Classification of Arrhythmia

- **Algorithmic Approaches with Feature Projection**

  - A Supervised Machine Learning Algorithm for Arrhythmia Analysis

  - Identifying Best Feature Subset for Cardiac Arrhythmia Classification





## 02 Preprocessing

# Missing Value Handling

Imputed using the mean for numerical features and mode for categorical features.

```
numeric_pipeline = Pipeline(  
    [("imputer", SimpleImputer(strategy = "mean")),  
      ("scaler", StandardScaler())]  
)  
  
categorical_pipeline = Pipeline(  
    [("imputer", SimpleImputer(strategy = "most_frequent")),  
      ("onehot", OneHotEncoder(handle_unknown = "ignore", drop = "if_binary"))]  
)
```

scikit-learn pipelines were used to streamline preprocessing

# Feature Encoding

Used one-hot encoding on all categorical features  
(necessary for distance computations)

```
categorical_pipeline = Pipeline(  
    [("imputer", SimpleImputer(strategy = "most_frequent")),  
     ("onehot", OneHotEncoder(handle_unknown = "ignore", drop = "if_binary"))]  
)
```

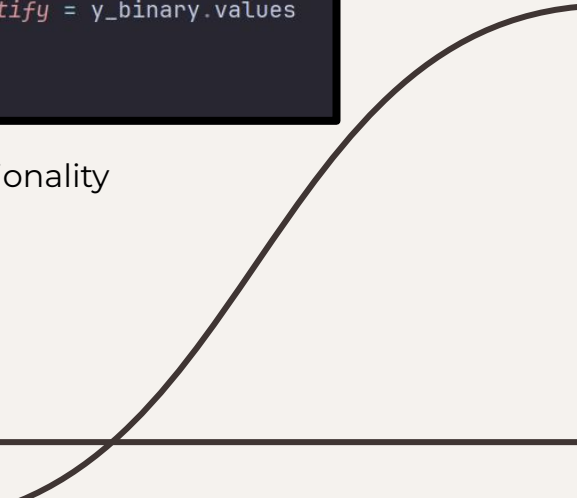
Encodings using scikit-learn functionality

# Dataset Splitting

- Training Set: 80% of data for model training
- Testing Set: 20% for unbiased evaluation of model performance

```
x_train, x_test, y_train, y_test = train_test_split(  
    *arrays: x_scaled, y_binary.values, test_size = 0.2, random_state = 42, stratify = y_binary.values  
)
```

Split with stratified distribution using scikit-learn functionality





# Feature Selection

```
pca = PCA(n_components = 0.95, random_state = 42)  
x_pca = pca.fit_transform(x_processed)
```

Preserves 95% of the variance while avoiding the curse of dimensionality

Eliminates all redundancy from correlated features.

# Converting Dataset to OvR

```
y_binary = y.apply(lambda x: 0 if str(x).strip() == "1" else 1)
```

Class Code 1 is separated from Class Codes 2 - 16

**class: Class Codes 01-16**

- a. Code 01: Normal
- b. Code 02: Ischemic changes (Coronary Artery Disease)
- c. Code 03: Old Anterior Myocardial Infarction
- ...
- p. Code 16: Others

# 03 Analysis



# Feature Weighting

- **Mutual Information (MI):**
  - Captures nonlinear dependencies between features and target
- **Absolute Pearson Correlation:**
  - Quantifies linear relationships

$$w = \alpha \times \frac{MI}{\Sigma MI + \epsilon} + (1 - \alpha) \times \frac{|Corr|}{\Sigma |Corr| + \epsilon}$$

# Neighborhood Size

$$k = k_{\min} + \frac{\text{local density} - \text{minimum density}}{\text{maximum density} - \text{minimum density} + \epsilon} \times (k_{\max} - k_{\min})$$

$k_{\min}$  defaults to 3 and  $k_{\max}$  defaults to 15

```
self.training_densities = np.median(distances[:, 1:], axis = 1)
self.min_density = np.min(self.training_densities)
self.max_density = np.max(self.training_densities)
```

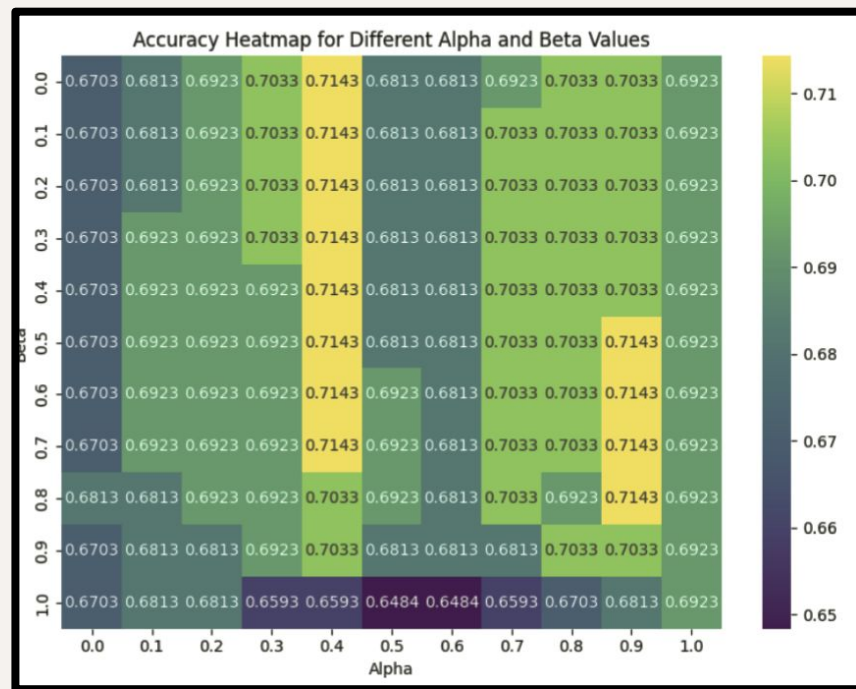
“distances” variable contains distances from  $r$  nearest neighbors

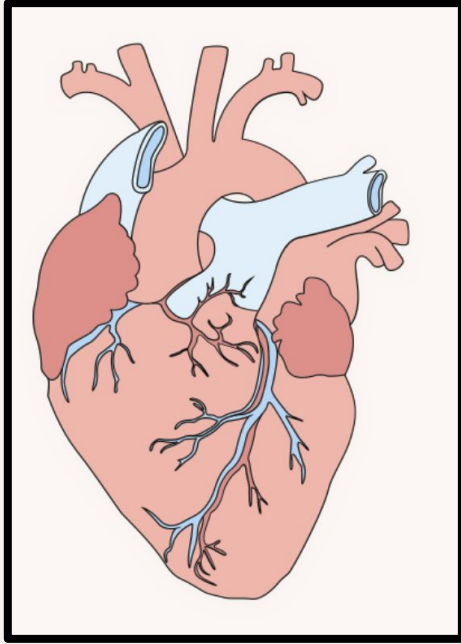
# Distance Metrics

- **Weighted Euclidean Distance:**
  - Incorporates feature weights directly
- **Local Mahalanobis Distance:**
  - Uses a locally computed covariance matrix to account for interdependencies in features.

$$d = \beta \times d_{euclidean} + (1 - \beta) \times d_{mahalanobis}$$

# Grid Search





# 04 Conclusion

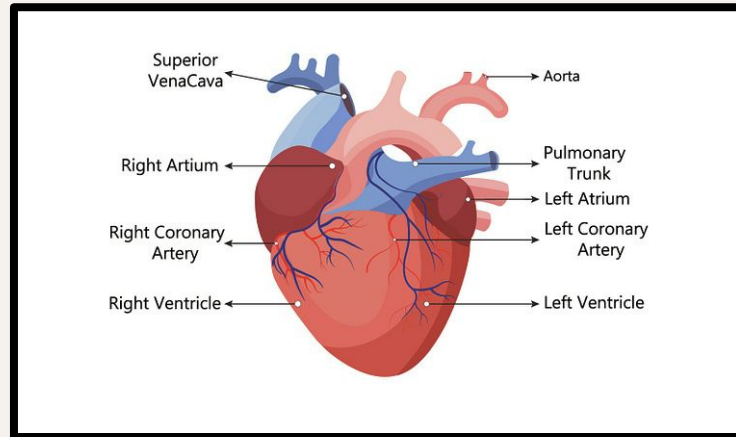


# Results

	Accuracy (%)	Precision (%)	Recall (%)	F1 Score (%)
Default Parameters	68.13	84.21	38.10	52.46
Optimized Parameters	71.43	86.00	40.50	55.07

# Future Work

- Hyperparameter tuning using cross-validation
- Application to other high-dimensional medical datasets
- Hybrid models that integrate adaptive KNN



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