

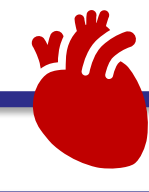
# ECG-Based Unsupervised Clustering in Coronary Artery Disease Associates with Ventricular Arrhythmia

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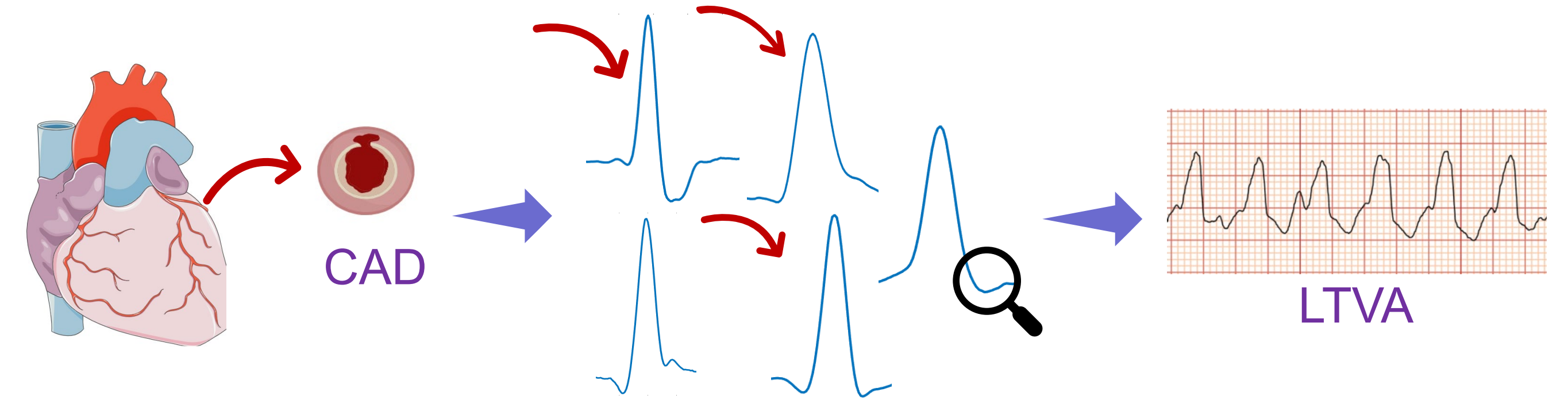
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## Background

- ❖ Coronary Artery Disease (CAD) is a leading cause of life-threatening ventricular arrhythmias (LTVAs).
- ❖ The presence of CAD slows ventricular conduction heterogeneously across individuals, manifesting as different QRS morphologies.



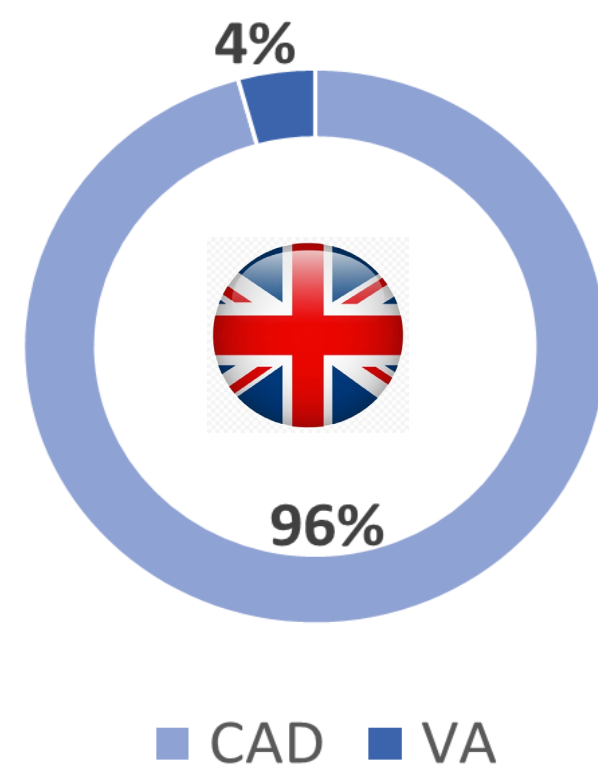
## Objective

- ❖ This study aimed to identify distinct clusters of CAD individuals based on QRS morphology using unsupervised learning.

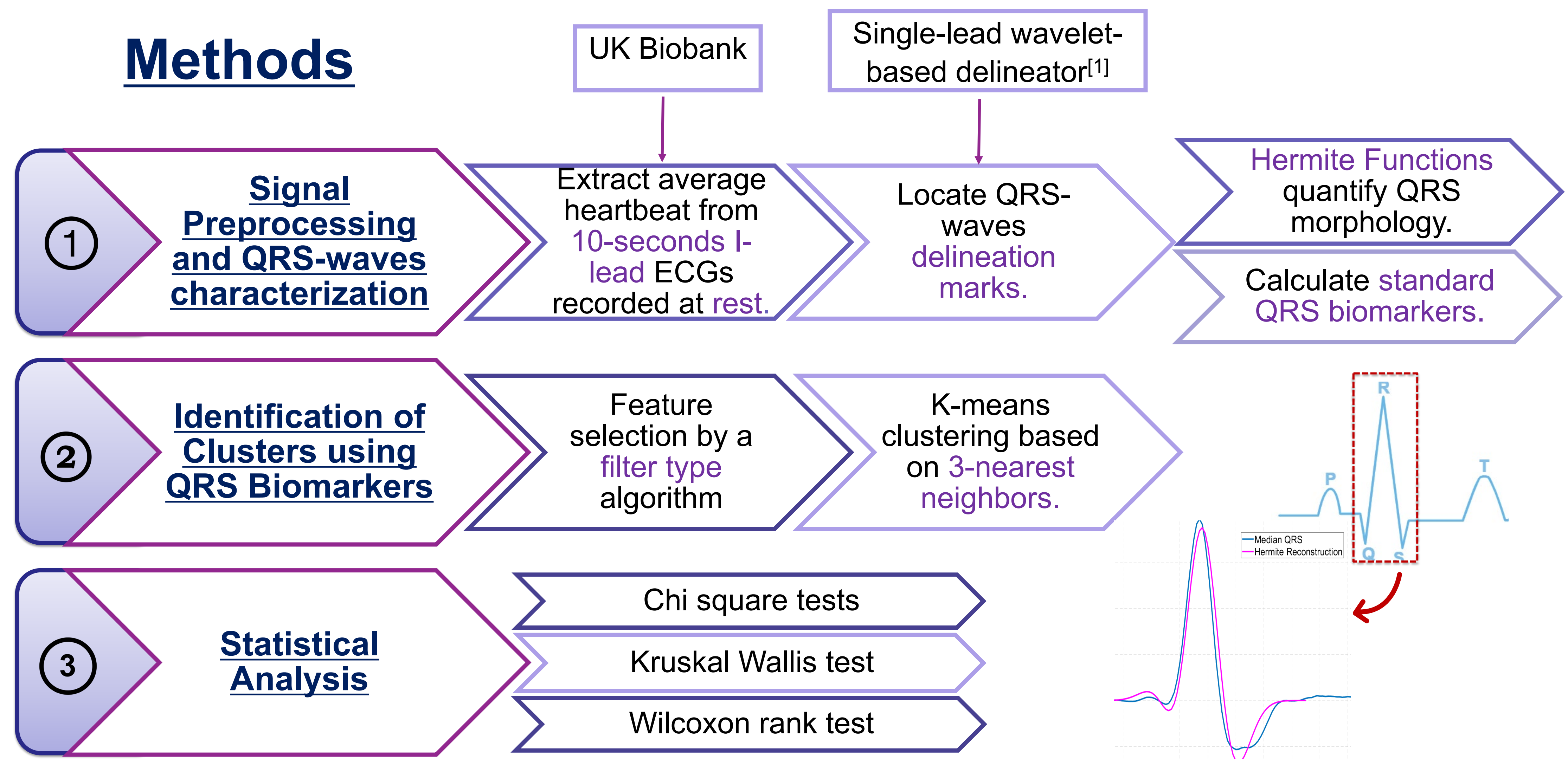
## UK Biobank Cohort

- ❖ The study population consisted of 1,458 individuals diagnosed with CAD in the UK Biobank study. (Application number: 8256)

Population Distribution



## Methods



## Results

- ❖ Cluster 1 was mainly characterized by lower QRS amplitude, flatter down slopes, and a wider QRS than clusters 2 and 3.
- ❖ Significant differences in morphological variations were described by Hermite basis 2, 3 and 5.

Representative Median Beat for each Cluster

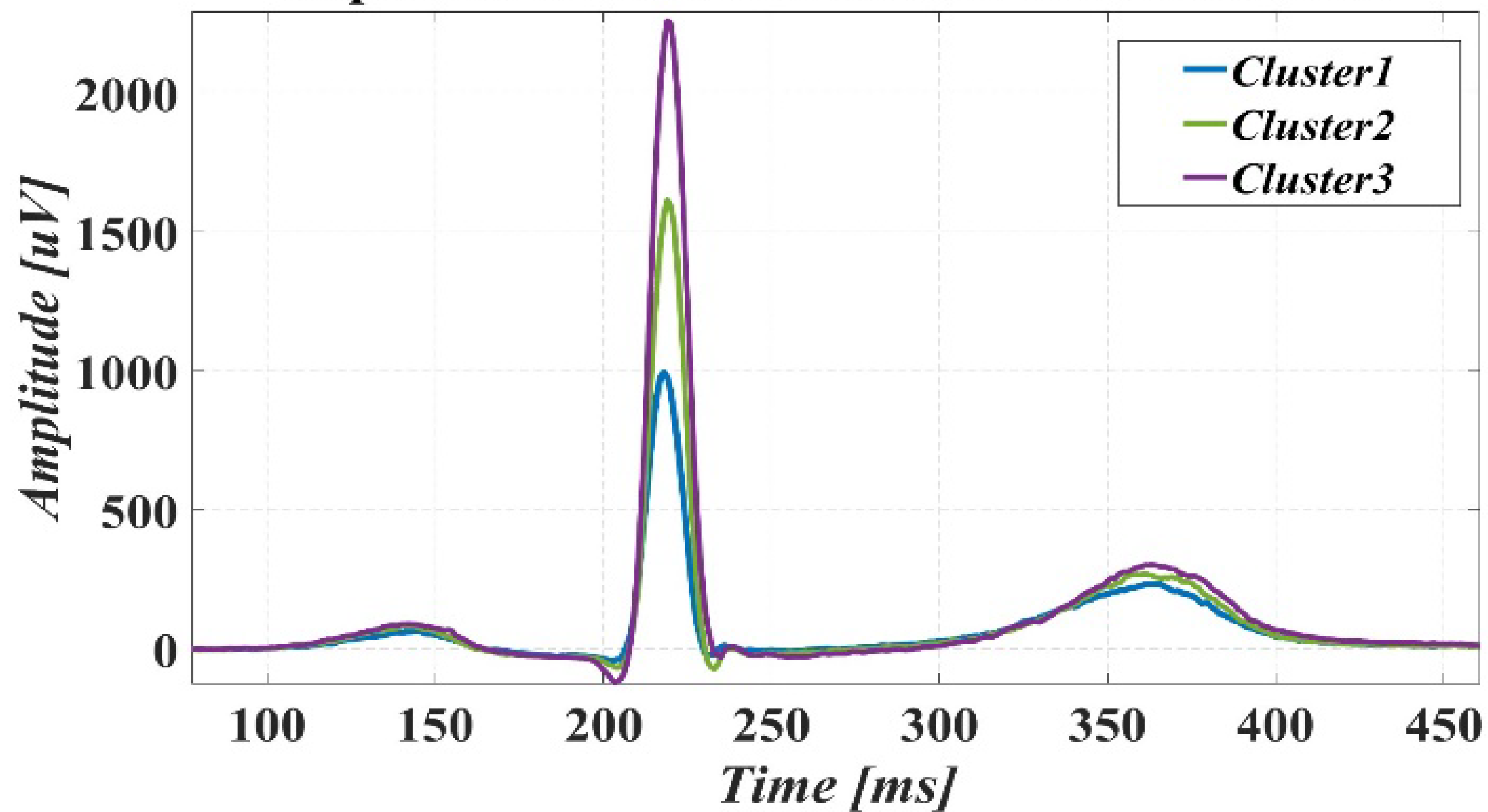
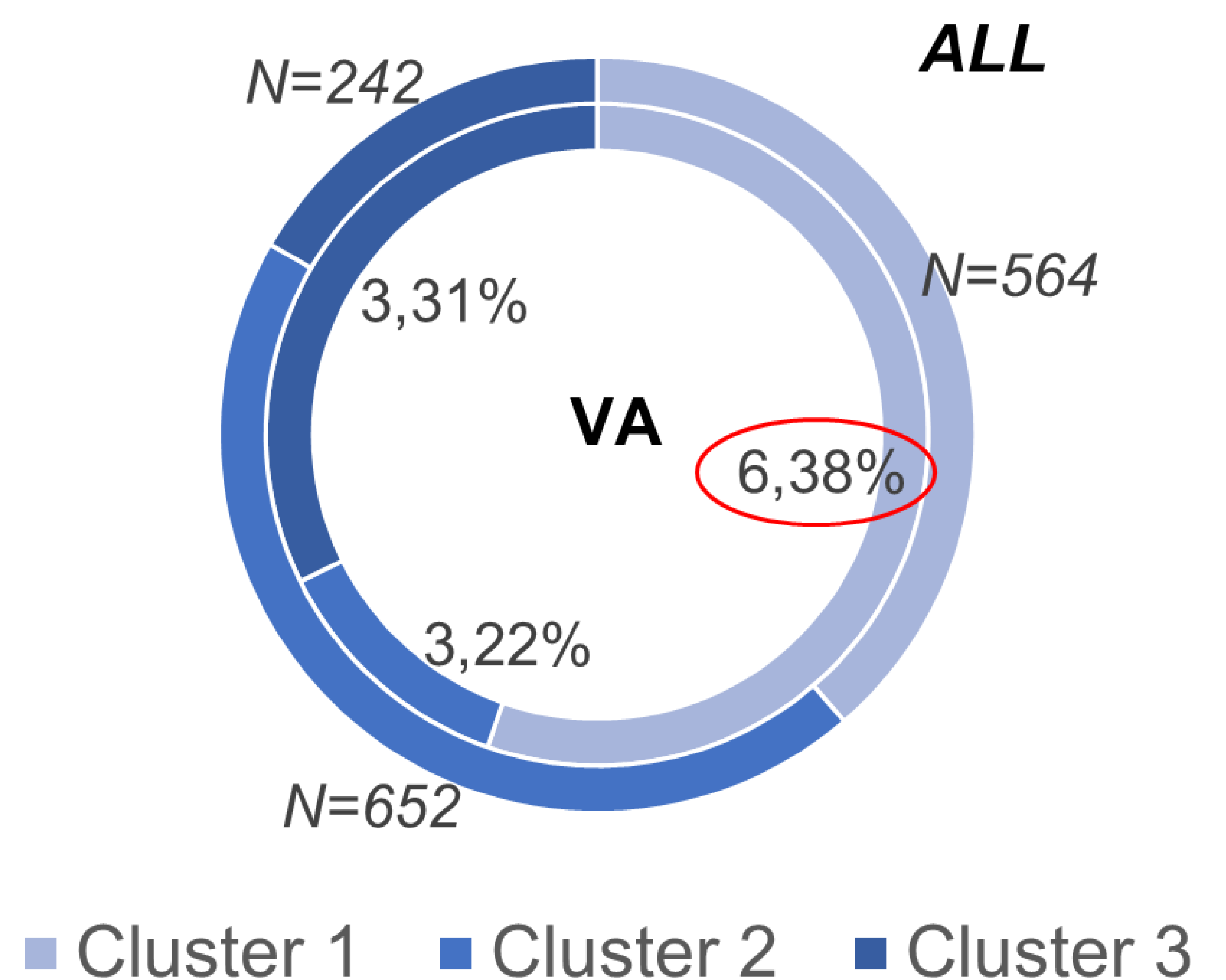


Figure 1. Median beat representative for each cluster obtained by the 3-nearest neighbors clustering algorithm

## Clustering Results



## Conclusions

- ❖ Our analysis has identified in an unsupervised manner three distinct clusters of CAD individuals using the QRS morphology.
- ❖ The cluster with the lowest QRS amplitudes and widest QRS complexes was strongly associated with LTVAs risk.
- ❖ Further studies will investigate the contribution of additional LTVAs risk factors in CAD.