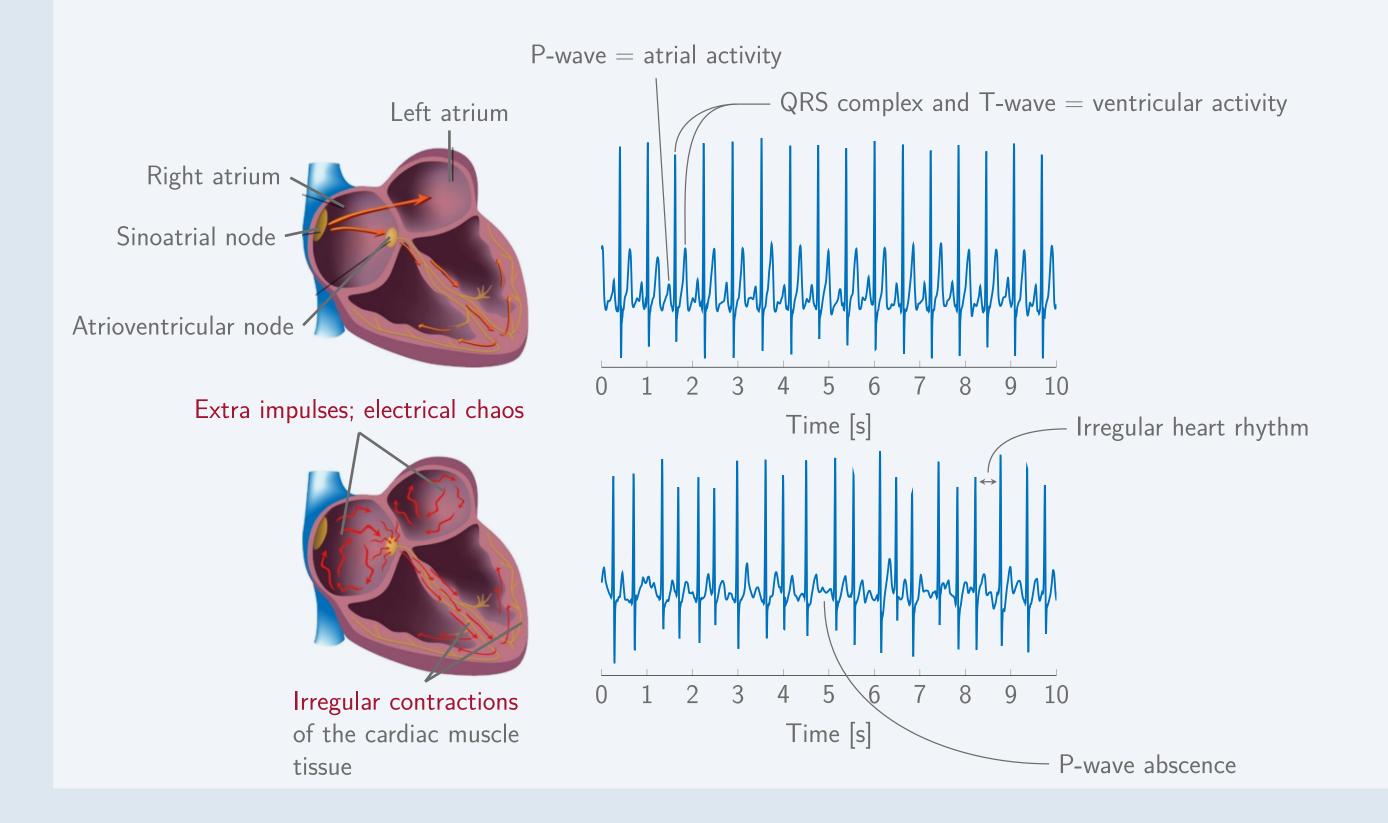
Tensor-based ECG Signal Processing Applied to Atrial Fibrillation Detection Simon Geirnaert

Joint work with G. Goovaerts, S. Padhy, M. Boussé, L. De Lathauwer, S. Van Huffel

Atrial fibrillation (AF) is the most common arrhythmia



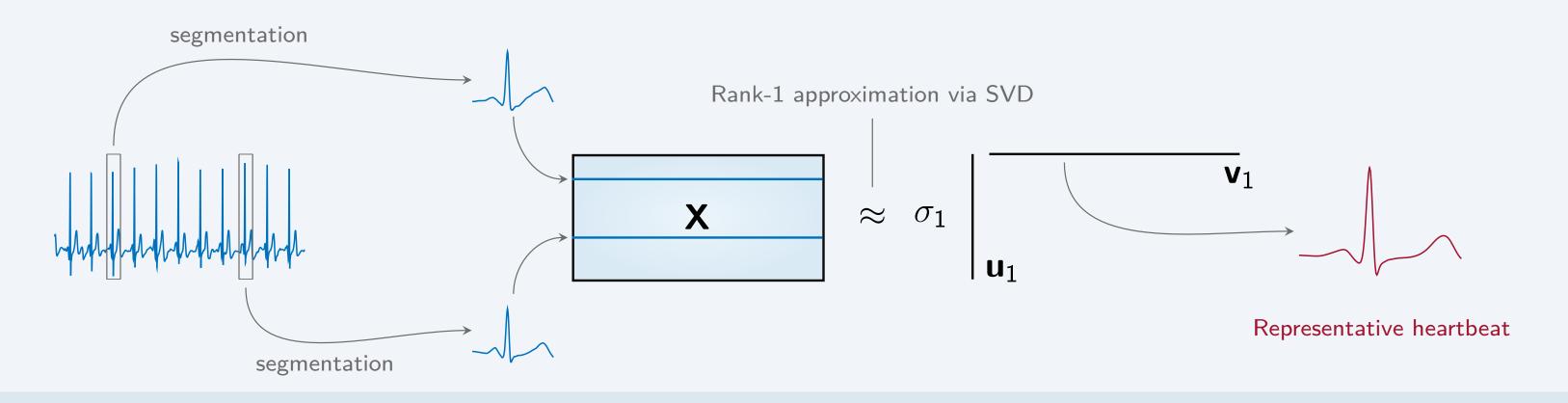
Some facts:

- ► 1 out of 4 will develop AF
- \blacktriangleright 1% of general population
- \blacktriangleright Latent risks: cloths of blood \rightarrow pulmonary embolism, stroke, ... Some trends:
- 1 Need for accurate and early detection
- 2 More and more data become available (*mHealth*) Motivate this research:

Goal

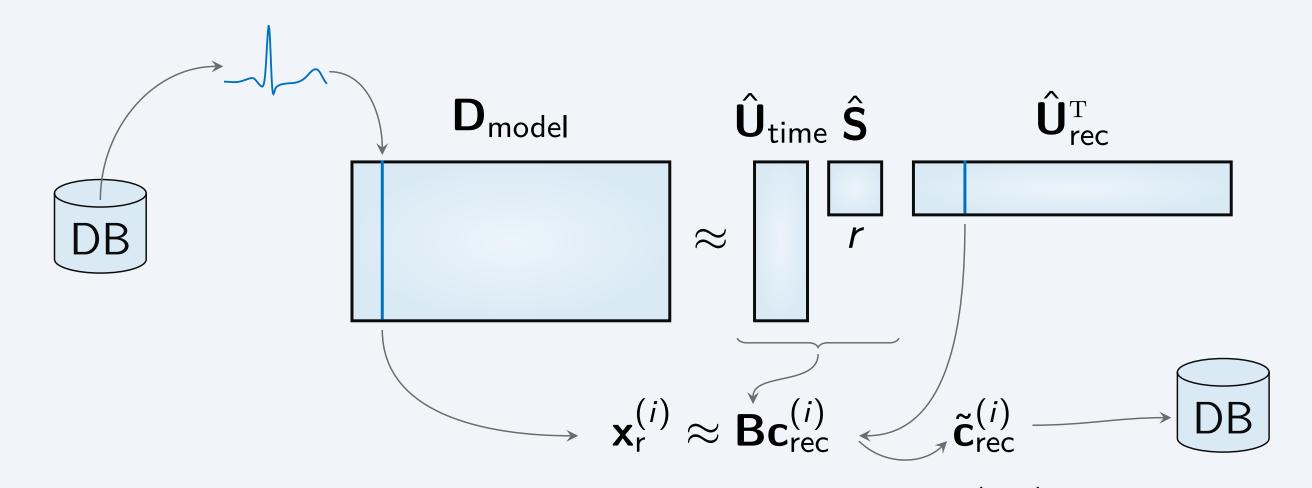
The *automatic* detection of AF in ECG signals

Represent ECG recording by compressing it in single representative beat: results in fixed-length representation and extra denoising



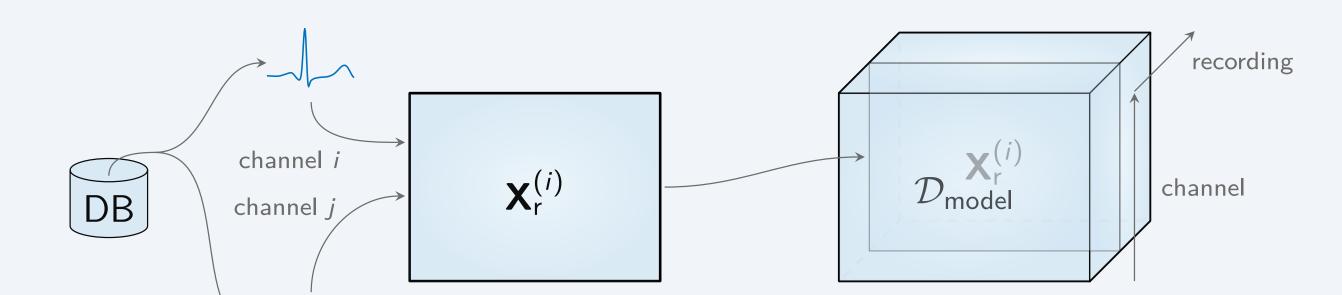
SVD-based detection for single-channel ECG

Modeling phase: truncated SVD



MLSVD-based detection for multi-channel ECG

Tensorization



Classification phase: classifying a new ECG recording $\mathbf{x}_{r}^{(new)}$

Morphological features: Solve:

$$\mathbf{x}_{r}^{(new)} = \mathbf{B}\mathbf{c}_{rec}^{(new)}
ightarrow \mathbf{\tilde{c}}_{rec}^{(new)}$$

Compute for each class c, $f_c^{(new)}$, based on:

$$f_c^{(\text{new})} = \sum_{i \in c} w_i s_i, \text{ with } s_i = \tilde{\mathbf{c}}_{\text{rec}}^{(\text{new})^{\mathrm{T}}} \tilde{\mathbf{c}}_{\text{rec}}^{(i)}, \forall i : 1 \leq i \leq M$$



A clear separation between the AF class and normal class (NSR) shown by the morphological features on the PhysioNet/Cinc challenge 2017 dataset.

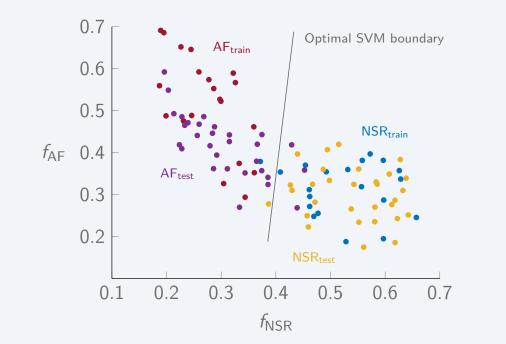
Modeling phase: truncated MLSVD

$$\mathcal{D}_{\text{model}} \approx \hat{S} \cdot_{1} \hat{\mathbf{U}}_{\text{channel}} \cdot_{2} \hat{\mathbf{U}}_{\text{time}} \cdot_{3} \hat{\mathbf{U}}_{\text{rec}}.$$
For one slice $\mathbf{X}_{r}^{(i)}$:
 $\mathbf{X}_{r}^{(i)} \approx \hat{S} \cdot_{1} \hat{\mathbf{U}}_{\text{channel}} \cdot_{2} \hat{\mathbf{U}}_{\text{time}} \cdot_{3} \mathbf{c}_{\text{rec}}^{(i)^{\text{T}}} \Leftrightarrow \text{vec} \left(\mathbf{X}_{r}^{(i)}\right) \approx \mathbf{B}$

 Classification phase: classifying a new multi-channel ECG recording X^(new)
 Morphological features: Solve:

$$\operatorname{vec}\left(\mathbf{X}_{\mathsf{r}}^{(\mathsf{new})}
ight)=\mathbf{B}_{(3)}^{\scriptscriptstyle\mathrm{T}}\mathbf{c}_{\mathsf{rec}}^{(\mathsf{new})}
ightarrow \mathbf{ ilde{c}}_{\mathsf{rec}}^{(\mathsf{new})},$$

compute **f**^(new) in a similar way



An almost perfect separation is possible based on the morphological features (MIT-BIH Arrhtyhmia + AFTDB dataset).

Feature set	P(%)	F_{1n}	F_{1a}	F_{1o}	F_1
HRV	77.6	0.85	0.76	0.59	0.73
SVD	70.0	0.81	0.57	0.40	0.59
SVD + HRV	80.2	0.87	0.80	0.65	0.77

Combining the new morphological features with classical HRV features with an SVM classifier leads to higher performances. This is shown on the PhysioNet/Cinc challenge 2017 dataset.

Morphological + classical heart rate variability (HRV) features

Feature set	AUC	F_1
HRV	1.000	0.983
MLSVD	0.990	0.933
MLSVD + HRV	1.000	1.000

The new morphological features result in a very high performance, using an SVM classifier (MIT-BIH Arrhtyhmia + AFTDB dataset).

Conclusion

The designed morphological features contain a lot of information by themselves and complement the classical HRV features when other rhythms, besides NSR and AF, are present (clinically relevant)









 $(i)_{3} c_{rec}^{(i)}$.