

## Introduction

Atrial fibrillation (AF) is a condition in which the atria, the upper chambers of the heart, contract in an uncoordinated fashion. AF reduces pumping efficiency and causes an irregular heart rate that cannot adapt anymore to the body's demands. AF is caused by a malfunction of the electrical activation mechanism that orchestrates the contraction of the heart. In a normal heart this activation is generated at regular intervals in a single location, called the sinus node, and from there spreads over the atria. In AF, the activation can run in circles, resulting in a chaotic and irregular contraction. Moreover, AF maintains and aggravates itself by causing structural changes. Treatment options depend on the disease stage. Our purpose is to develop better diagnostic methods to identify the stage of AF in individual patients. We conducted this project to develop a computer model of the heart that could help us to better understand the relation between the stage of AF and the characteristics of the electrocardiogram (ECG) measured on the skin of the patient.

The structural remodeling process that maintains AF causes heterogeneous changes in the atrial tissue's capacity to propagate the electrical activation. Therefore, the complexity of the propagating activation wavefront could be a measure of the disease stage. It would be desirable to assess this complexity from the ECG measured at the body surface. We hypothesized that more complex conduction patterns in the atria generate more complex potential patterns on the body surface. The purpose of our project was to find and validate measures for the complexity of ECG signals that correlate well with the complexity of wavefront propagation in the atria. To achieve this, we developed a large-scale computer model of the human atria with which we could accurately simulate the atrial activation pattern and the electrical signals (ECG and local electrograms) that result from it.

## Results of this project

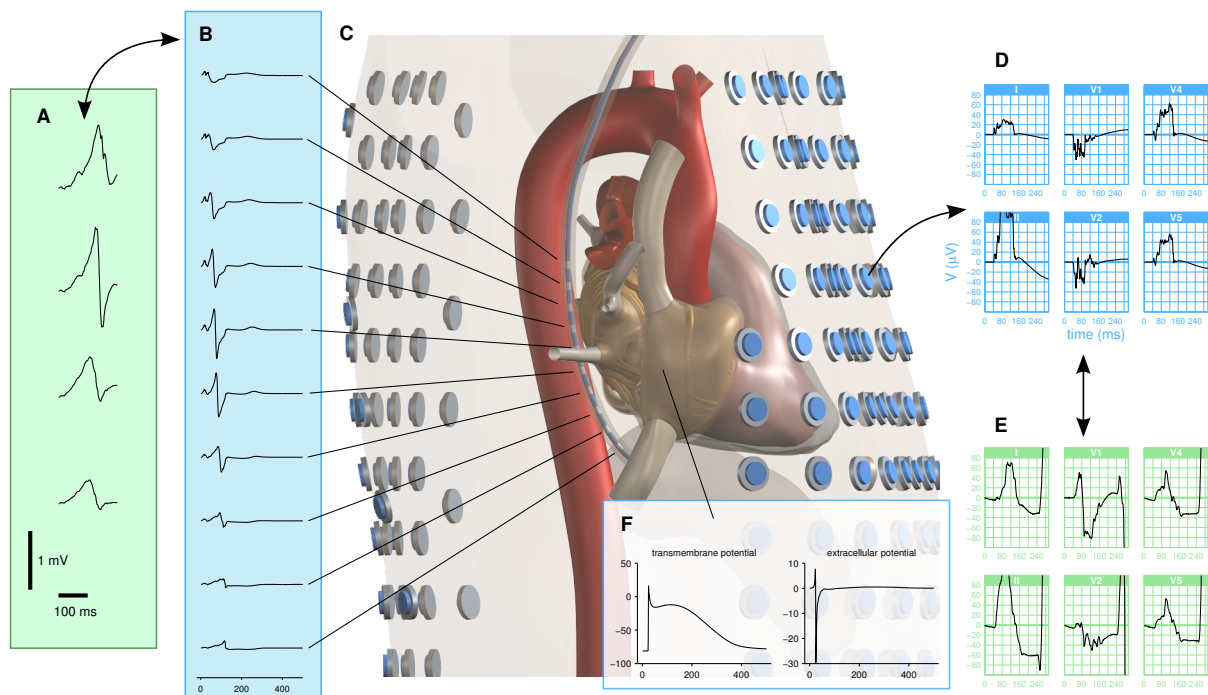
We have built an anatomical model of the atria that accurately represents the structural inhomogeneities that are present even in the healthy atria, and that allows pathological changes in the structure to be represented. Notably, the model includes a clear representation of the thin atrial wall and the thicker muscle bundles that are invariably present. With this model we simulated the electrical activation mechanism of the atria based on the transmembrane ionic currents in the atrial muscle cells. From the results we computed the ECG together with electrograms in the atria (which can be measured with endocardial catheters in real patients) as well as electrograms in the esophagus (which can also be measured in patients using a special probe).

We discovered that the model could not reproduce the atrial ECG as it is commonly described in cardiology textbooks and observed in clinical ECGs. Even after a long period of tuning, the simulated ECG remained considerably more complex than what is seen on the clinical ECG. However, we were aware that clinical ECGs are electronically filtered to remove noise and interference – unwanted signals that are almost as large as the atrial ECG itself. We suspected that these filters hide much of the complexity that is present in the real physiological signals. We therefore decided to measure ECGs ourselves, using techniques that avoid filtering. We used specialised high-quality ECG equipment to record ECGs with extremely low noise and interference levels. Moreover, we made long recordings (5 to 10 minutes) to obtain hundreds of heartbeats. These recordings were cut into single-beat segments, each aligned on the P wave (the part of the ECG generated by the atria). In each of these segments, the physiological signal is supposed to be the same, while the noise and interference are different each time. Therefore, when the average of hundreds of segments is computed, the real signals add up while the noise and interference cancel out. Thus a very clean signal is obtained without

any filtering. We performed this analysis on normal volunteers of age varying from 23 to 75 years.

The P waves that we obtained in this manner were very different from the filtered signals that are seen in the clinic. Despite the absence of heart disease, the signals were very complex, with 2 to 10 peaks appearing in a single P wave, even in the youngest subjects.

This unexpected discovery and the period of model tuning that convinced us to do the research that led to it has delayed the project significantly. However, it has important consequences for our thinking about the ECG changes to be expected in AF patients, because the complexity that we expected to see in their ECGs is already present in normal ECGs, and takes specialised equipment to measure. Moreover, since the true shape of the atrial ECG is so poorly known, our work will lead to an important publication, which is currently being edited.



*Model results compared with measured data. Measured trans-esophageal ECGs (panel A) are compared with their simulated equivalents (B) extracted from a virtual probe in the torso model (C). Surface ECGs are extracted from hundreds of surface locations (D) and compared to measured ECGs obtained with the same electrode set in human volunteers (E). The model also computes several local signals such as transmembrane potentials and local electrograms (F) which, in our study, have no measured equivalents, but can be compared to data from other studies.*

## Followup

Although the project is finished, the work is continuing with funding from other sources. Finally convinced that our model is accurate, we are performing simulations of reentrant activity in the atria. This work is continued with various degrees of structural remodeling in the atria, while we are quantifying the complexity of both the activation pattern and the simulated ECG.

By combining model results with measured data we have been able to show that the atrial ECG, when adequately recorded, reflects detailed anatomical features of the atria. This finding may have important clinical implications on the long term, which we are currently investigating.