

# Variability in the Activity of the Heart

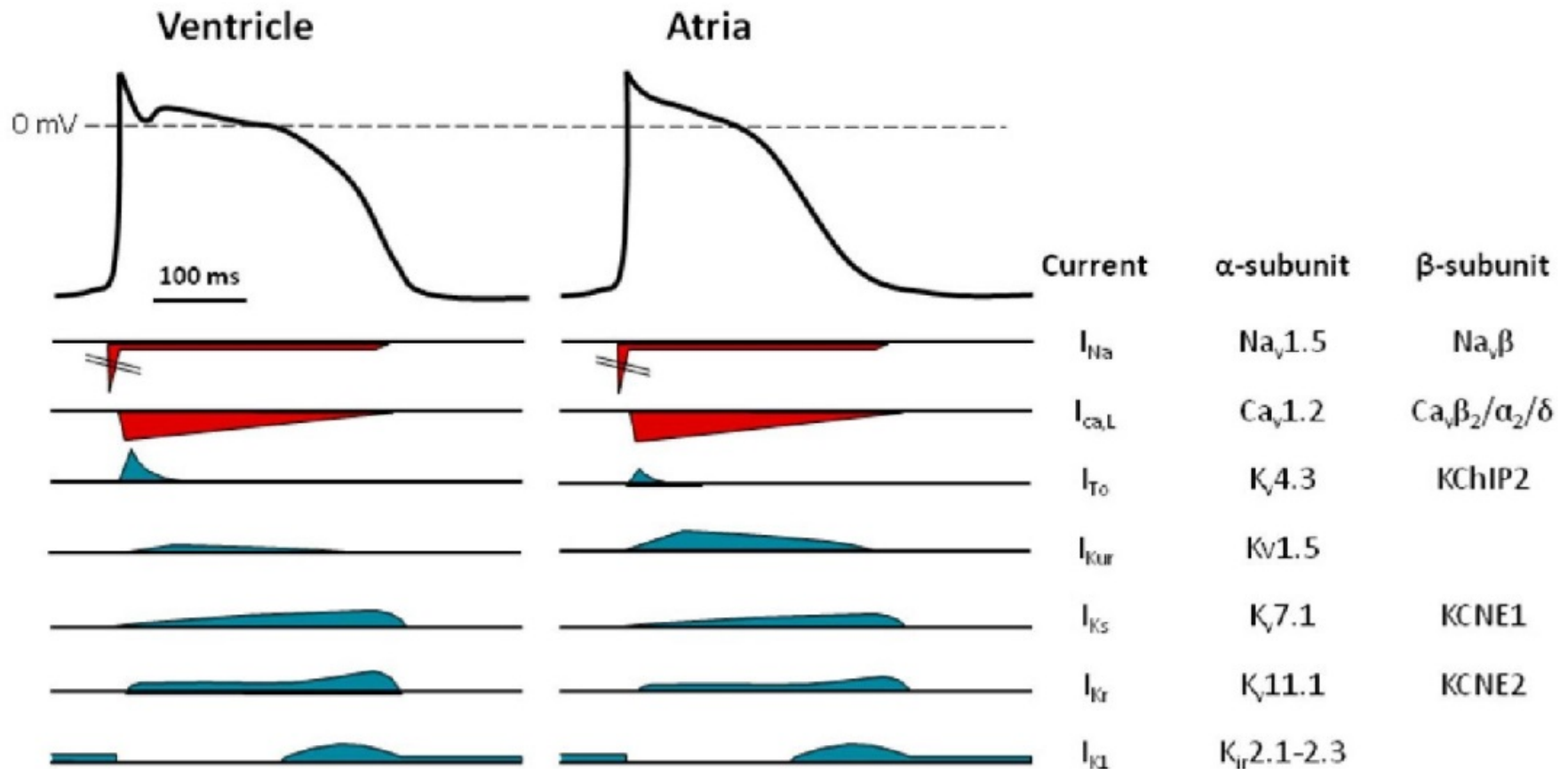
Techniques for the  
calibration of  
populations of models



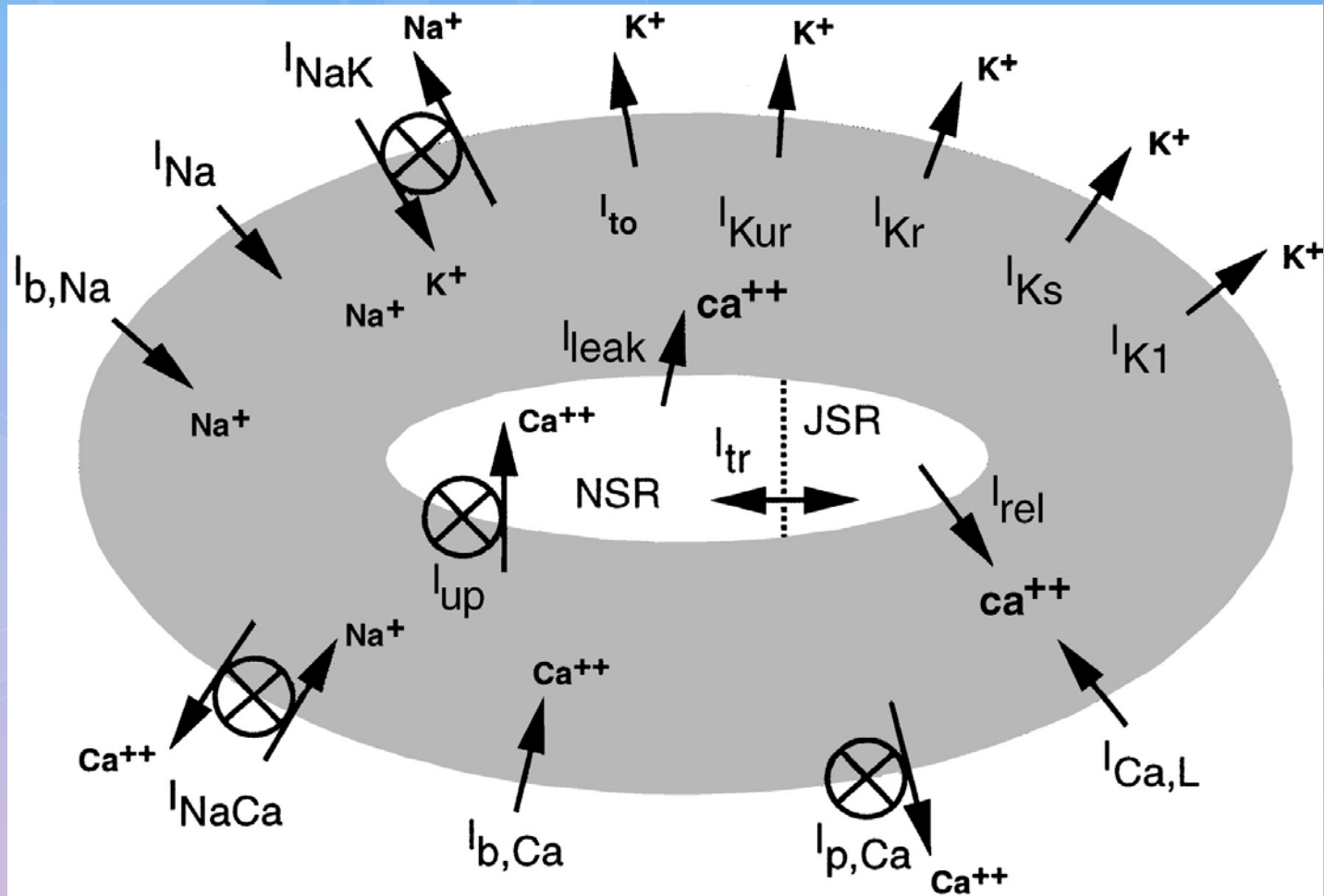
AUSTRALIAN RESEARCH COUNCIL CENTRE OF EXCELLENCE FOR  
MATHEMATICAL AND STATISTICAL FRONTIERS

- ◉ **Within-subject Variability:** Individual heart cells show considerable variance in electrophysiological properties, even cells of the same type
- ◉ **Between-subject Variability:** Further variance occurs between different members of the population. Factors include: genetics, gender, age, pathologies

- I work with single cell data for the **action potential (AP)**:

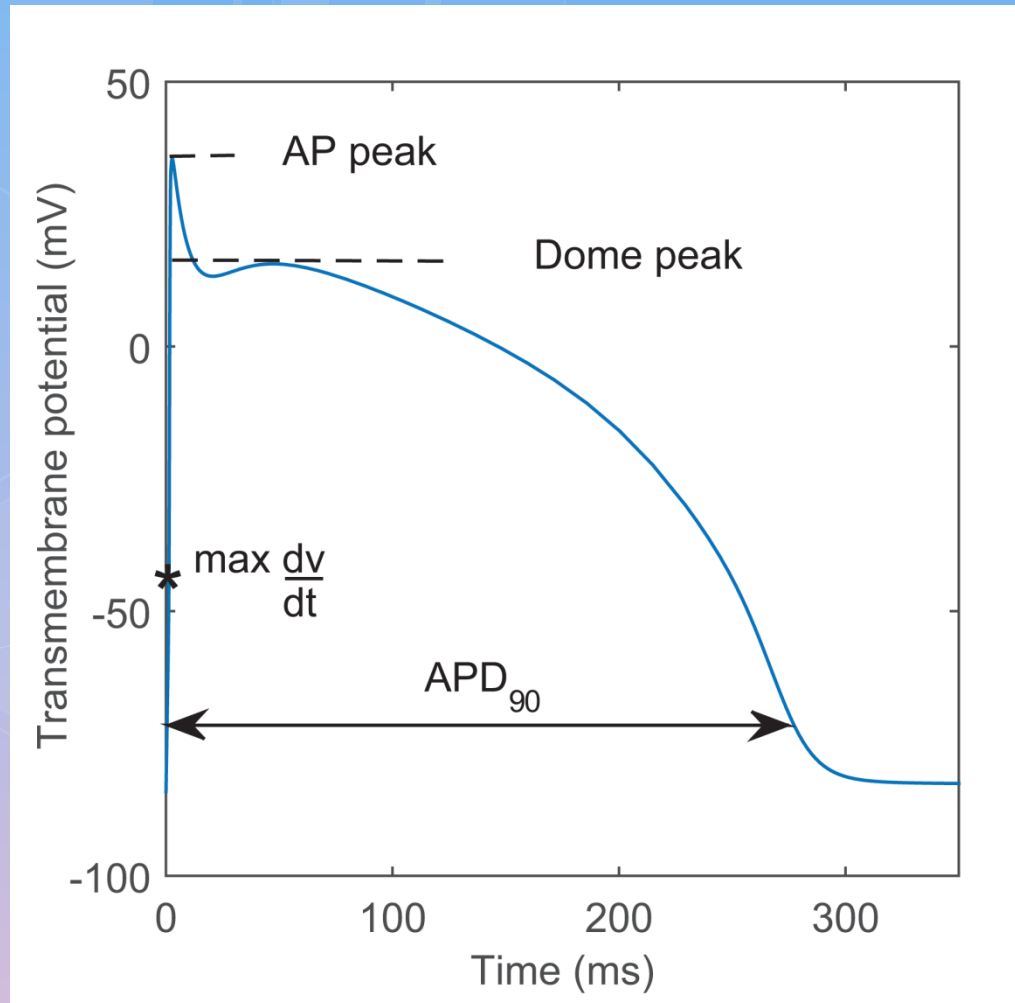


## Ion Channels



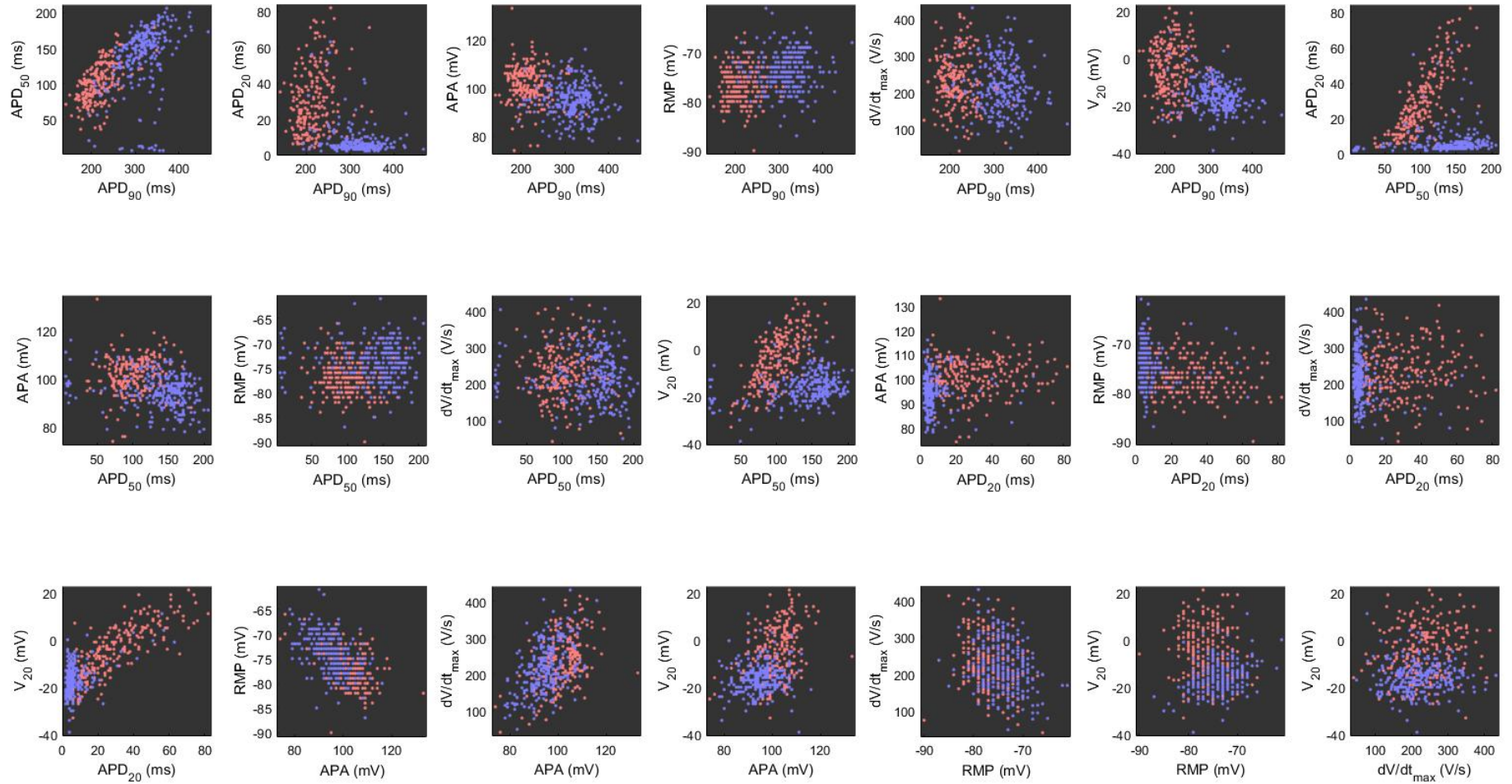
- Cell polarisation is achieved by flow of ions, predominantly  $Na^+$ ,  $K^+$  and  $Ca^{2+}$ .
- These currents activate and deactivate in a complex fashion that produces the action potential.

- In a single cell, represent the changing values in time of:
  - Voltage
  - Gating variables
  - Ion concentrations
- Systems of 40+ ODEs, highly nonlinear
- Jolt them with stimulus current repeatedly until steady state
- Can be solved by e.g. MATLAB's *ode15*



- Data is provided in terms of **biomarkers**, measurements of the key properties of the curve

# Cardiac Data

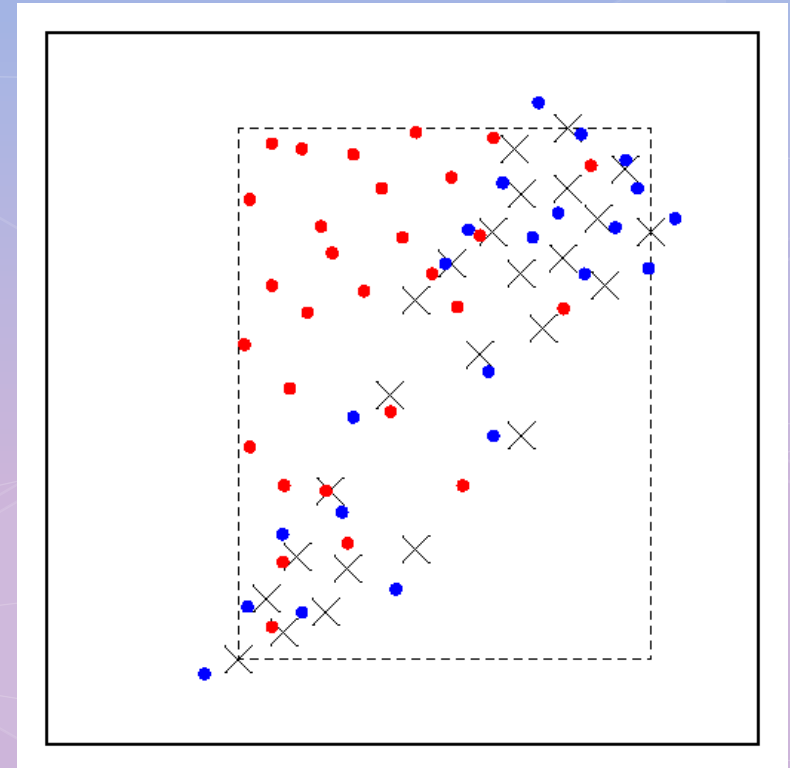


Blue – healthy sinus rhythm, Red – chronic atrial fibrillation

- Highly variable data – how do we handle that with a single deterministic model?
- Perhaps **multiple** models, then?
- Representing a population of people with a *population of models* (POM) – the same model, but with different parameters to represent each individual's properties



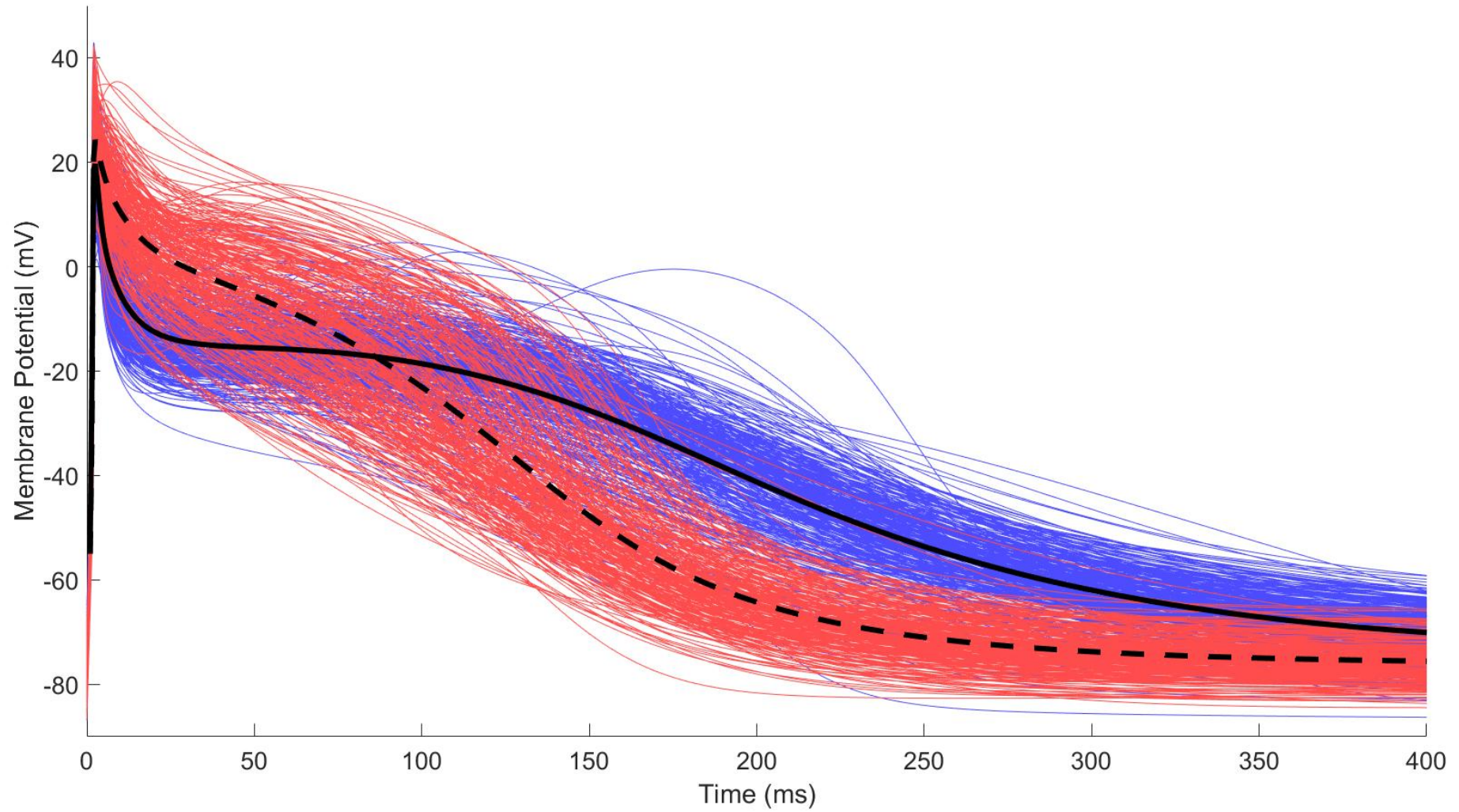
- We can select models from some parameter space, but how do we choose?
- Calibration is typically performed by ensuring all model outputs fall within the ranges of the data (and are hence physiologically realistic)



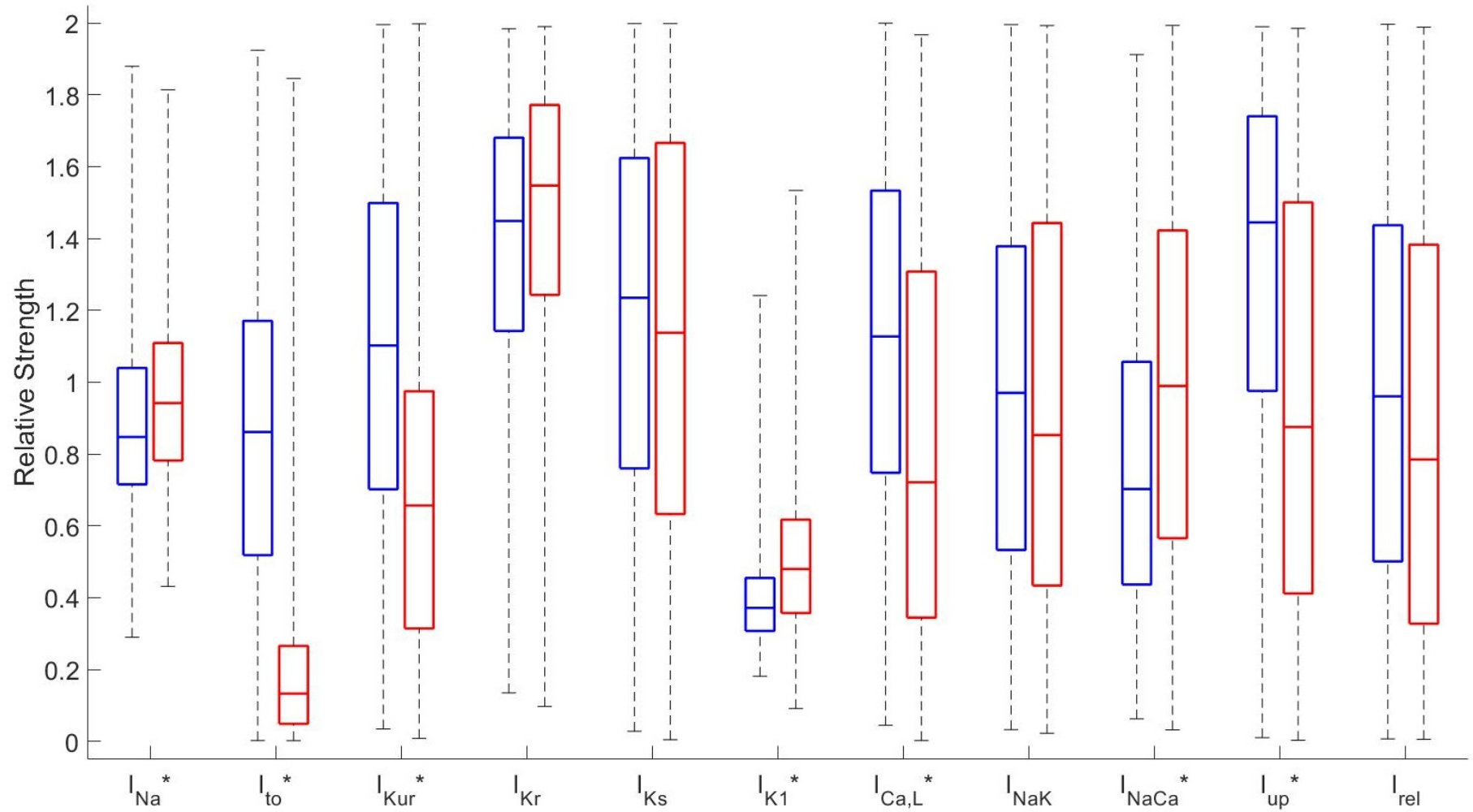
- Not good enough!

- We want to calibrate to the **distribution** of the data directly:
  1. Estimate distribution (MVKDE)
  2. Try to sample according to this distribution (SMC)
  3. Clean the results using an optimally-selected subpopulation (simulated annealing)

## Results 1



## Results 2



- Can we use this technique on other stratified data? What about outside of cardiac electrophysiology?
- Could we solve this instead as a fully Bayesian inverse problem?

$$\mathbf{b} = \mathcal{M}(\boldsymbol{\theta}) + \boldsymbol{\varepsilon}$$

- What new research questions can we tackle once adding in the spatial dimension? (Travelling waves of excitation in multiple cells)