

$$f\left(\mathbf{r} + \frac{\mathbf{p}}{m} \Delta t, \mathbf{p} + \mathbf{F} \Delta t, t + \Delta t\right) d^3 \mathbf{r} d^3 \mathbf{p} = f(\mathbf{r}, \mathbf{p}, t)$$

$$dN = f(\mathbf{r}, \mathbf{p}, t) d^3 \mathbf{r} d^3 \mathbf{p}$$

$$\frac{\partial f_i}{\partial t} + \frac{\mathbf{p}_i}{m_i} \cdot \nabla f_i + \mathbf{F} \cdot \frac{\partial f_i}{\partial \mathbf{p}_i} = \left( \frac{\partial f_i}{\partial t} + \frac{\mathbf{p}_i}{m_i} \cdot \nabla f_i + \mathbf{F} \cdot \frac{\partial f_i}{\partial \mathbf{p}_i} \right)$$

$$\int A F_j \frac{\partial f}{\partial p_j} d^3 \mathbf{p} =$$

$$\hat{\mathbf{L}}_{NR} = \frac{\partial}{\partial t} + \frac{\mathbf{p}}{m} \cdot \nabla + \mathbf{F} \cdot \frac{\partial}{\partial \mathbf{p}}$$

# BDT Project: Image-based FE modelling

Peter Hunter

Auckland Bioengineering Institute

University of Auckland, NZ

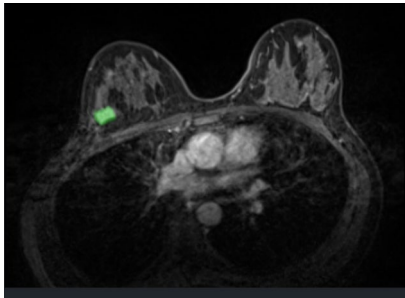
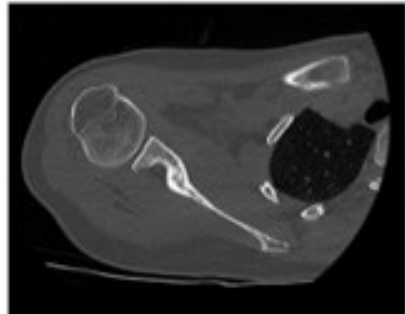
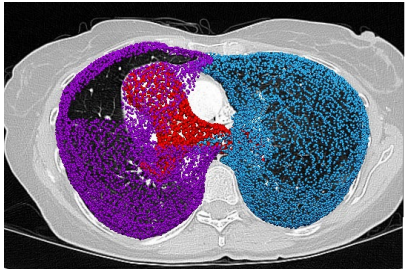
September 30, 2024

IMAG/MSM Teaming4BDT

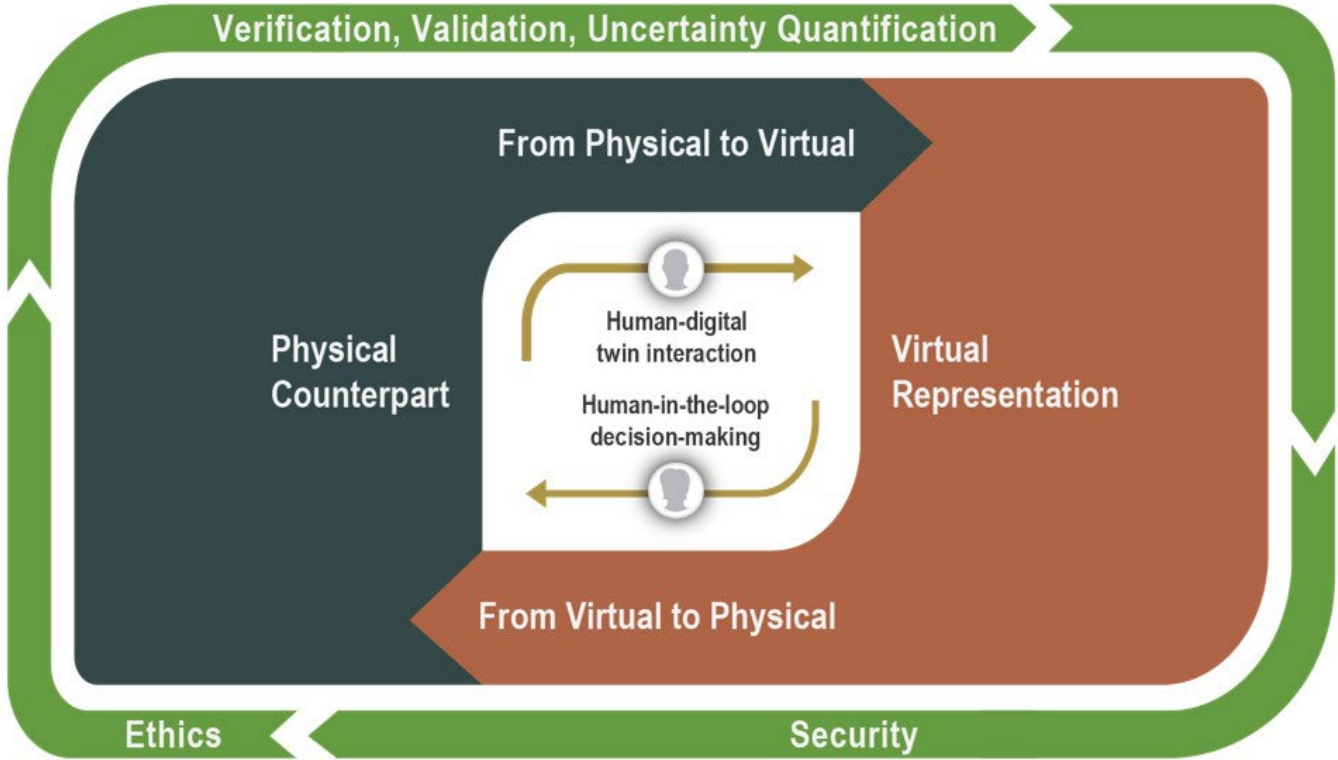
Meeting

# NASEM Loop: Image-based FE modelling

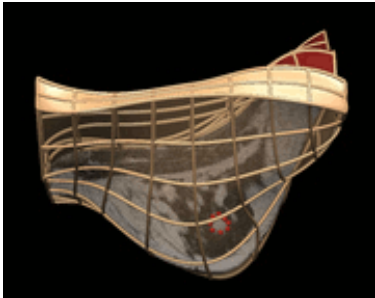
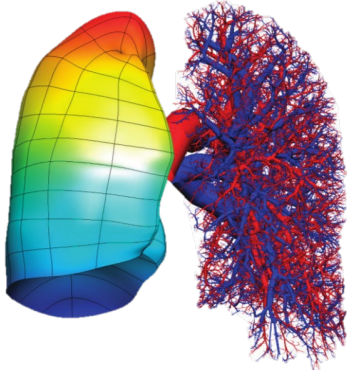
## Clinical images



The FE models are personalised and, with appropriate boundary conditions, are used to solve the eqns of continuum mechanics



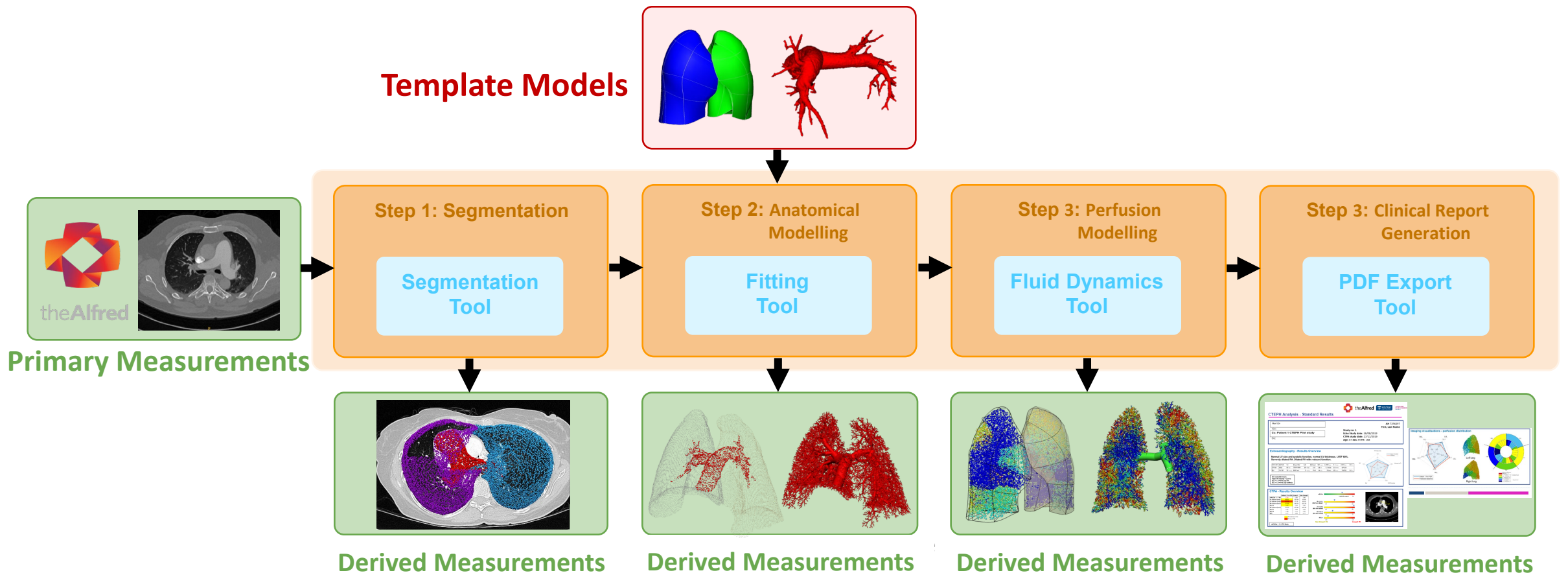
## FE models



The output of the modelling are clinical reports

# Physical and Virtual Assets and Their Interaction

The physical asset is the clinical image (3D and time varying). The virtual asset is the FE model. The model is fitted to the images for each individual patient and used to predict outcomes of treatment strategies.



# Problem and BDT Solution

The FE models are personalised and are currently used to help clinicians diagnose pathologies and development treatment strategies.

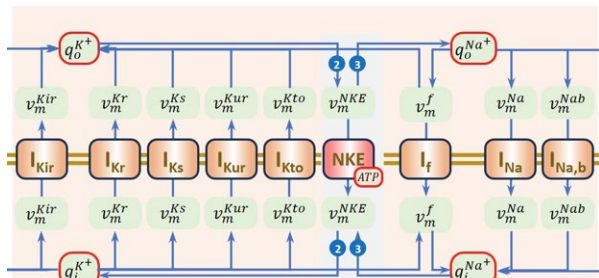
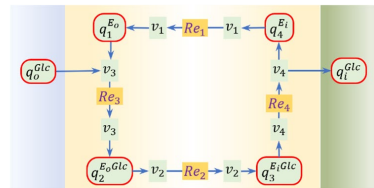
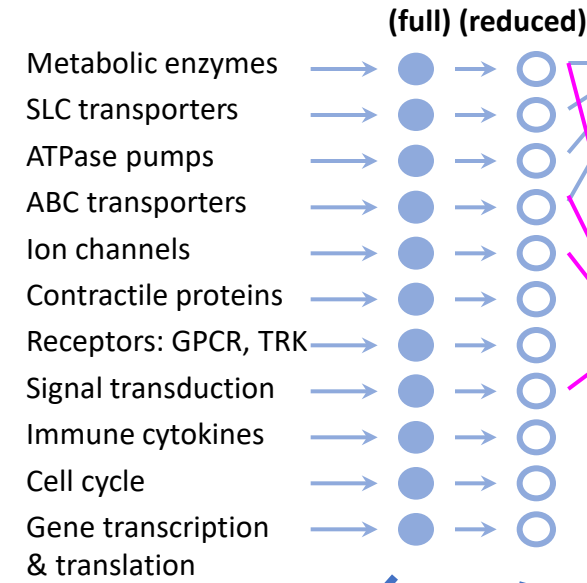
The problem is that they are not yet multiscale – we would like to include information on tissue biopsies, blood biomarkers and genetic testing to improve the clinical utility of the models when these additional data are available.

The next slides illustrate our approach to physics-based multiscale modelling.



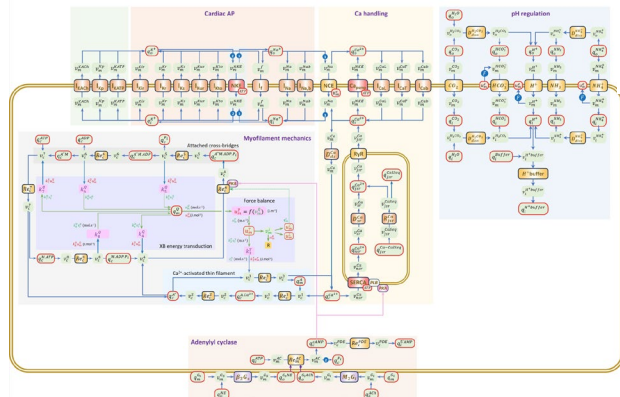
# Multiscale modelling with bond graphs

## 1. BG protein templates



## 2. FCUs

- Metabolism
- Action potentials
- Calcium handling
- pH regulation
- Myofilament mechs
- Receptors & signaling
- Glucose transport
- Na/K regulation
- Cell cycle
- Cell adhesion
- Gene regulation



## 3. Cells

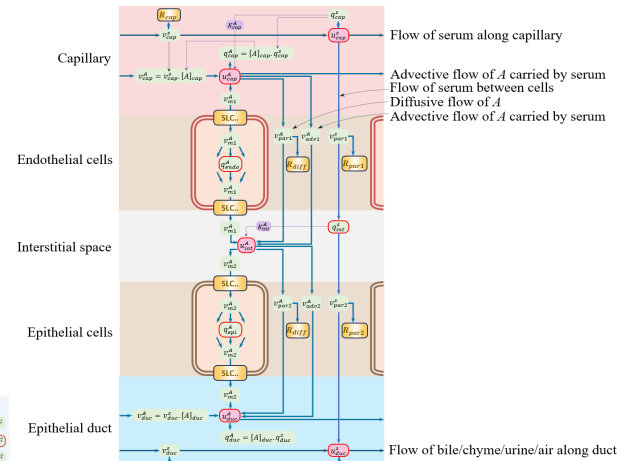
- Endothelial cell
- Epithelial cell
- Cardiomyocyte
- Muscle fiber
- Fibroblast
- Smooth muscle cell
- Osteocyte
- Enterochromaffin cell
- Hepatocyte
- Goblet cell
- Red blood cell



## 4. FTUs

(including 3D structure)

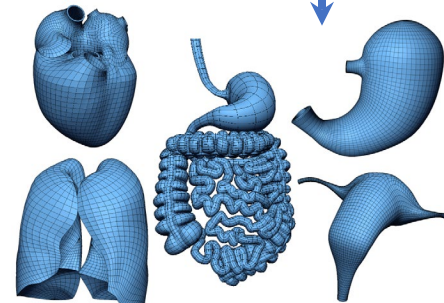
- Glomerulus, ...
- Lung alveoli
- Myocardial sheet
- Liver lobule
- Bone osteon
- Fibre, etc



## 5. Organs

● =surrogate

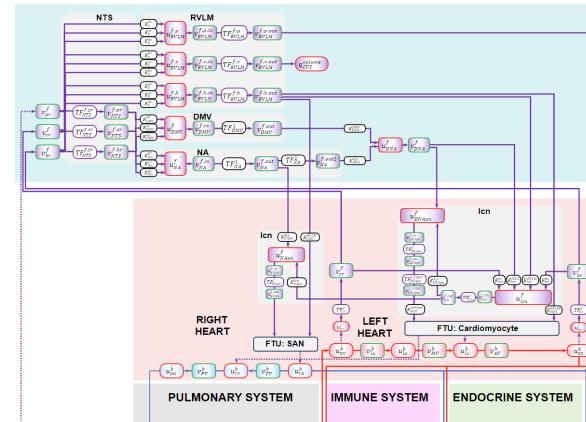
- nephron
- acinus
- block
- lobules
- osteon
- fascicle
- ...



## 6. FPU

(3D anatomical model of the body)

- Regulation of fluid volume
- Regulation of arterial blood pressure
- Regulation of pH
- Regulation of electrolyte balance
- Regulation of glucose
- Regulation of O<sub>2</sub> and CO<sub>2</sub>
- Regulation of calcium
- Regulation of phosphate
- Regulation of body temperature

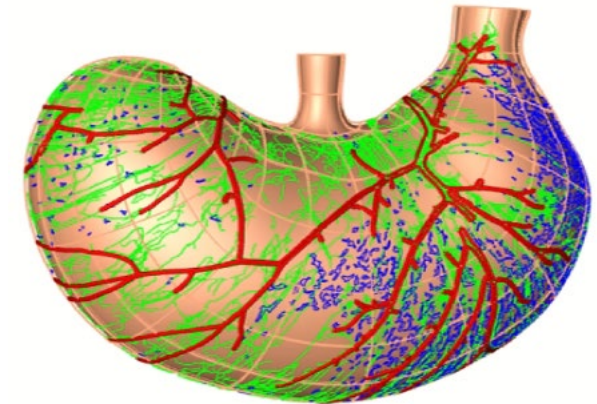
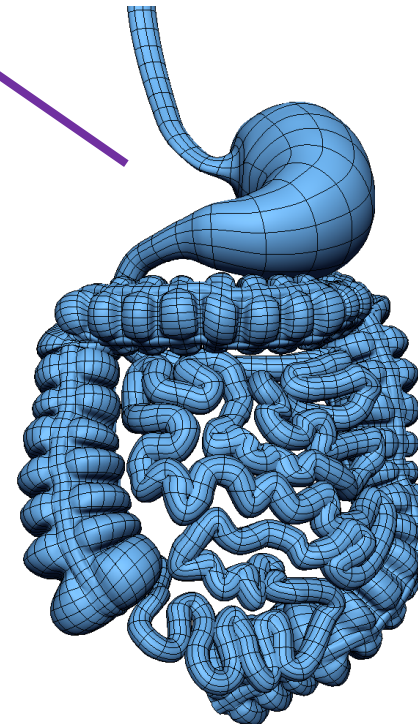
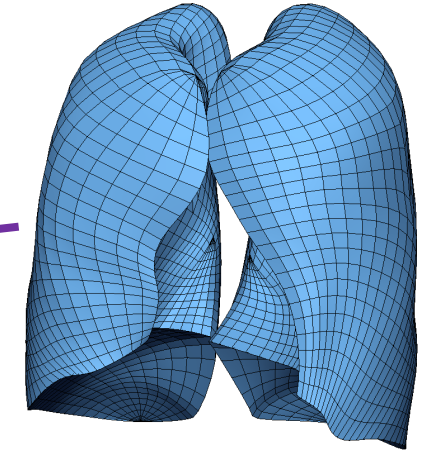
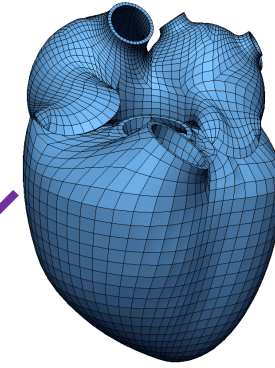


Finite element models that are subsequently reduced (with AI methods) to surrogate models.



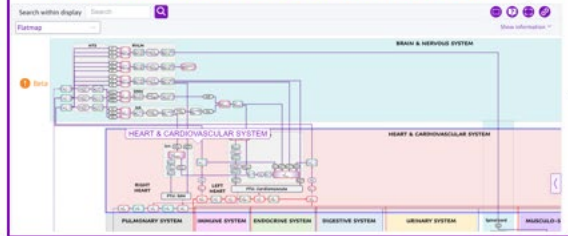


# Scaffolds for spatial mapping



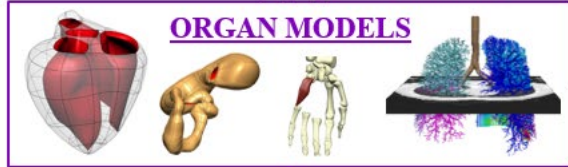
# CLINICAL WORKFLOWS

## SYSTEMS PHYSIOLOGY MODELS



Physiological function tests

Reduction to produce surrogate models



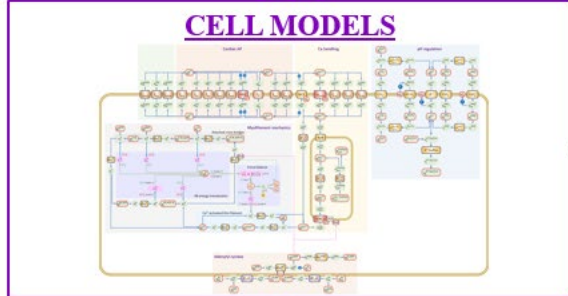
Clinical images

Model reduction with constitutive laws

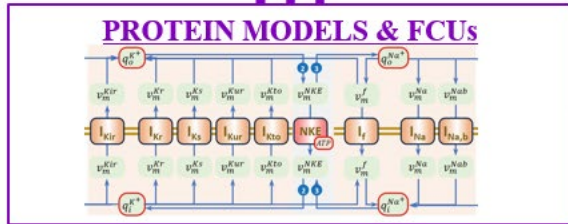


Tissue biopsies

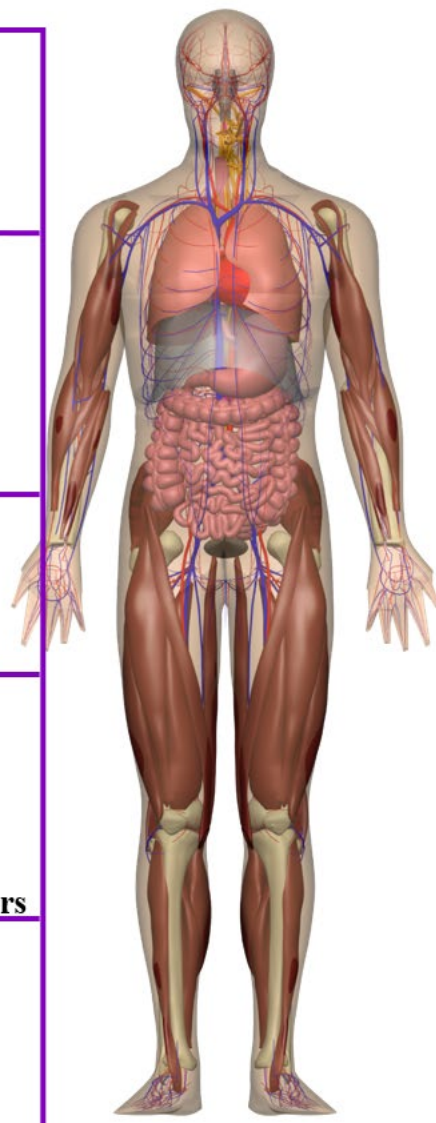
FTUs incorporate small # of cell types



Blood biomarkers



Genetic tests



Constrained by observed behaviour



## Model parameters



Constrained by physics and genetics



# Questions

## Contact me

Email: [p.hunter@auckland.ac.nz](mailto:p.hunter@auckland.ac.nz)

Lab: Auckland Bioengineering Institute: <https://www.auckland.ac.nz/en/abi.html>

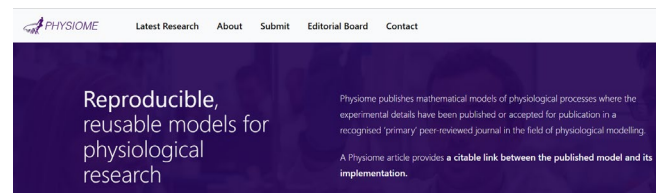
## Resources:



CellML homepage <https://www.cellml.org/>



OpenCOR software <https://opencor.ws/>



<https://journal.physiomeproject.org/>

Mapping tools <https://docs.sparc.science/docs/map-core-scaffold-mapping-tools>

