

A modeling framework for Biomedical Digital Twins

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Community resources

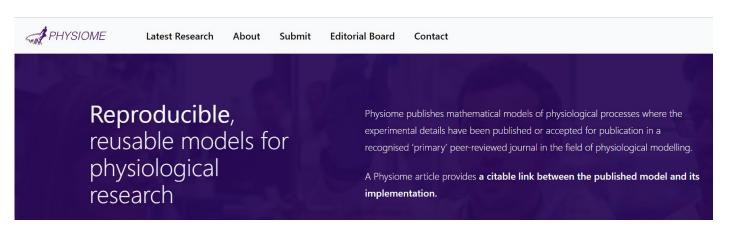


CellML homepage

https://www.cellml.org/



OpenCOR software https://opencor.ws/ (Download, documentation, tutorial)

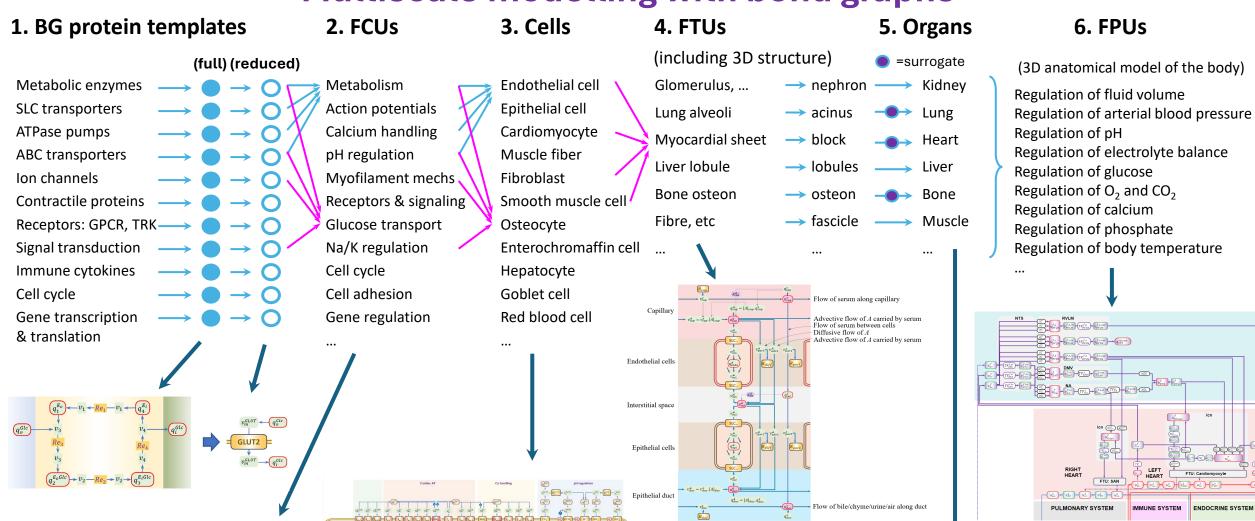


https://journal.physiomeproject.org/

Mapping tools

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

Multiscale modelling with bond graphs



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

Example: SLC transporters

458 genes



66 gene families



Examples:

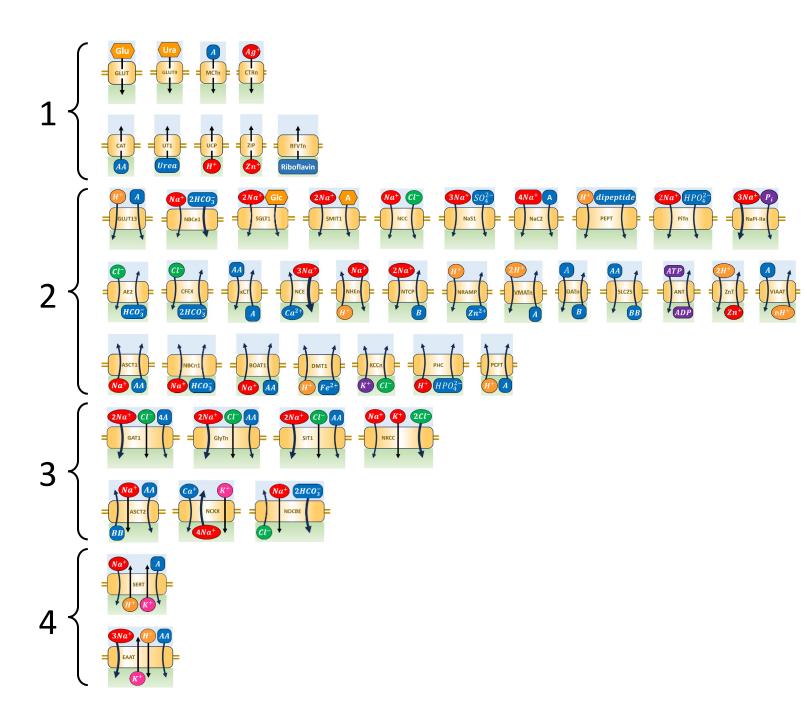
SLC1: (EAAT) Glutamate transport

SLC2: (GLUT) Facilitated glucose transport

SLC4: (AE, NBC) Bicarbonate cotransporters

SLC5: (SGLT) Na-glucose cotransport

SLC8: (NCE) Na-Ca exchange SLC9: (NHE) Na-H exchange

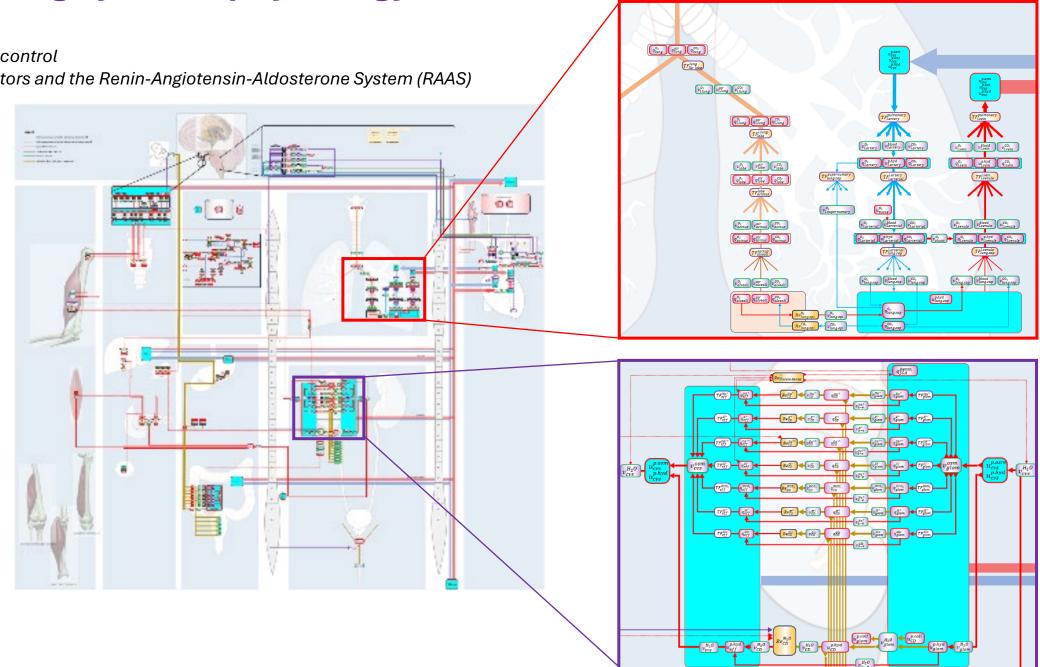


Linking systems physiology with molecular mechanism

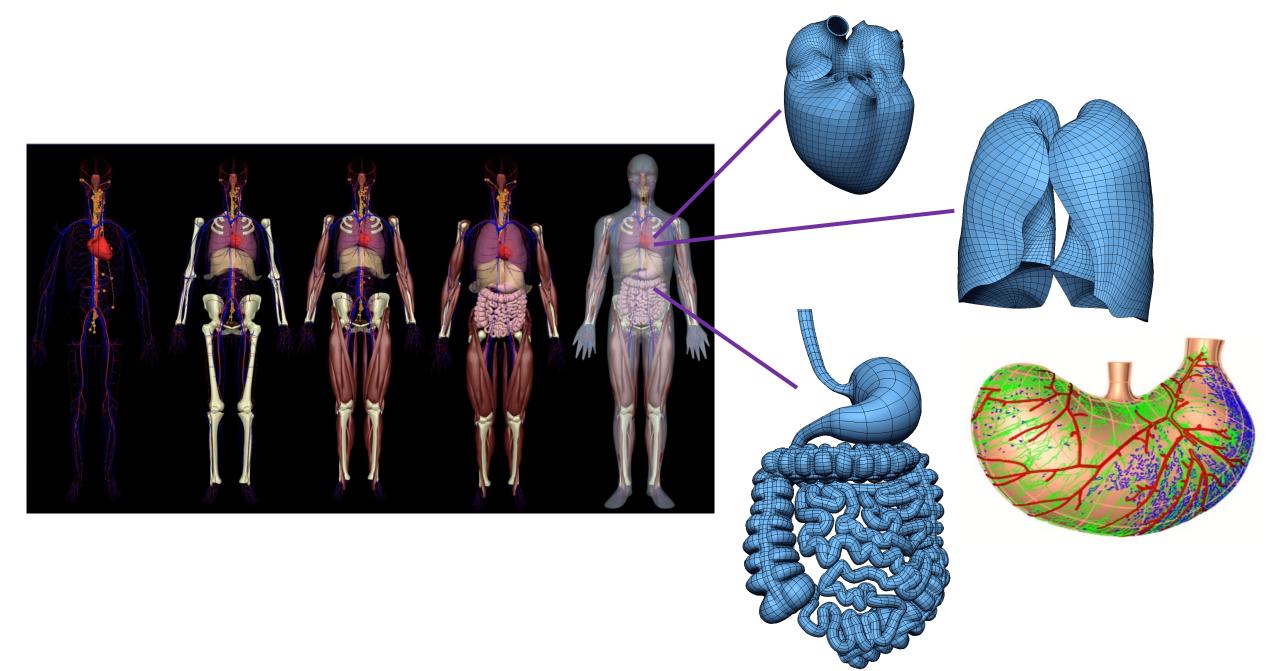
Food intake and glucose control

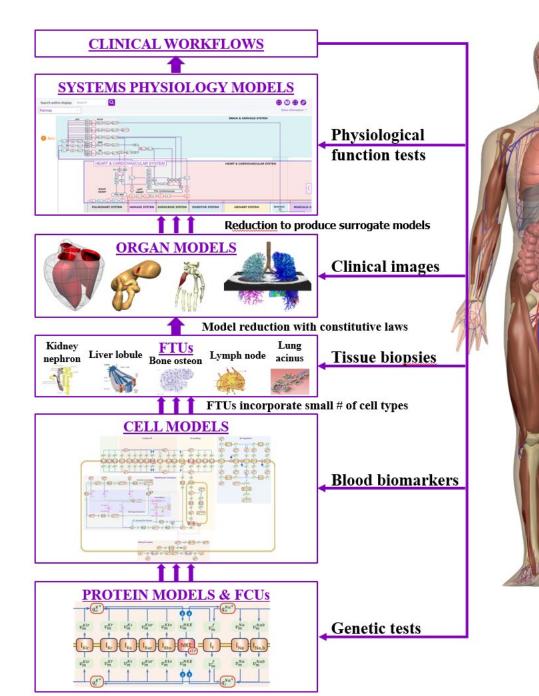
Control of BP: Baroreceptors and the Renin-Angiotensin-Aldosterone System (RAAS)

Control of blood volume Control of metabolism Calcium homeostasis Circadian rhythms Growth regulation



Scaffolds for spatial mapping





Constrained by observed behaviour



Model parameters



Constrained by physics and genetics

Thank you!