



Advances Towards the Virtual Human Twin at BSC

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HPC a National and European Infrastructure

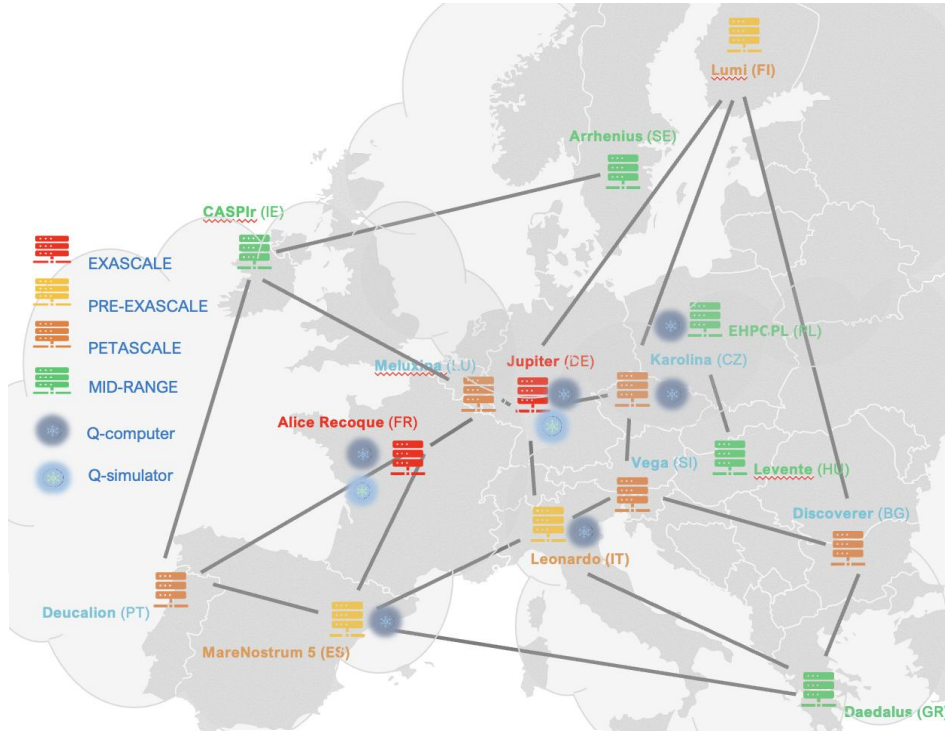
#EuroHPC Joint Undertaking

The European High Performance Computing Joint Undertaking (EuroHPC JU) will pool European resources to develop top-of-the range exascale supercomputers for processing big data, based on competitive European technology.

Member countries are Austria, Belgium, Bulgaria, Croatia, Cyprus, Czechia, Denmark, Estonia, Finland, France, Germany, Greece, Hungary, Iceland, Ireland, Italy, Latvia, Lithuania, Luxembourg, Malta, Montenegro, Netherlands, North Macedonia, Norway, Poland, Portugal, Romania, Serbia, Slovakia, Slovenia, Spain, Sweden, Türkiye and United Kingdom.



EuroHPC
Joint Undertaking



MareNostrum5

ACC – Accelerated
Intel Sapphire Rapids
NVIDIA Hopper
Peak perform: 260 Pflops
Sustained HPL: **163 Pflops**

GPP – General Purpose
Intel Sapphire Rapids
Peak perform: 45,4 Pflops
Sustained HPL: **35,4 Pflops**

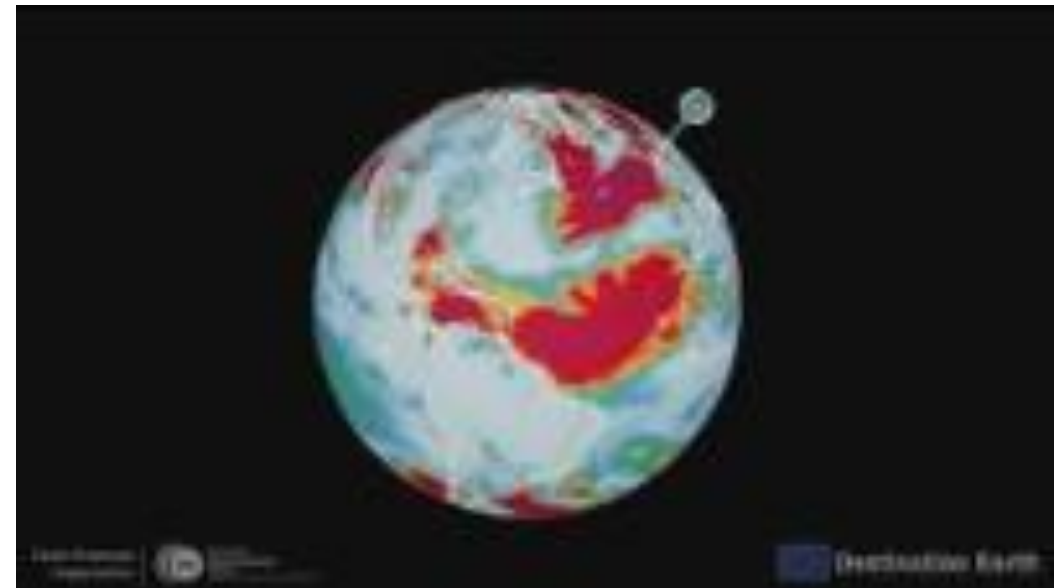
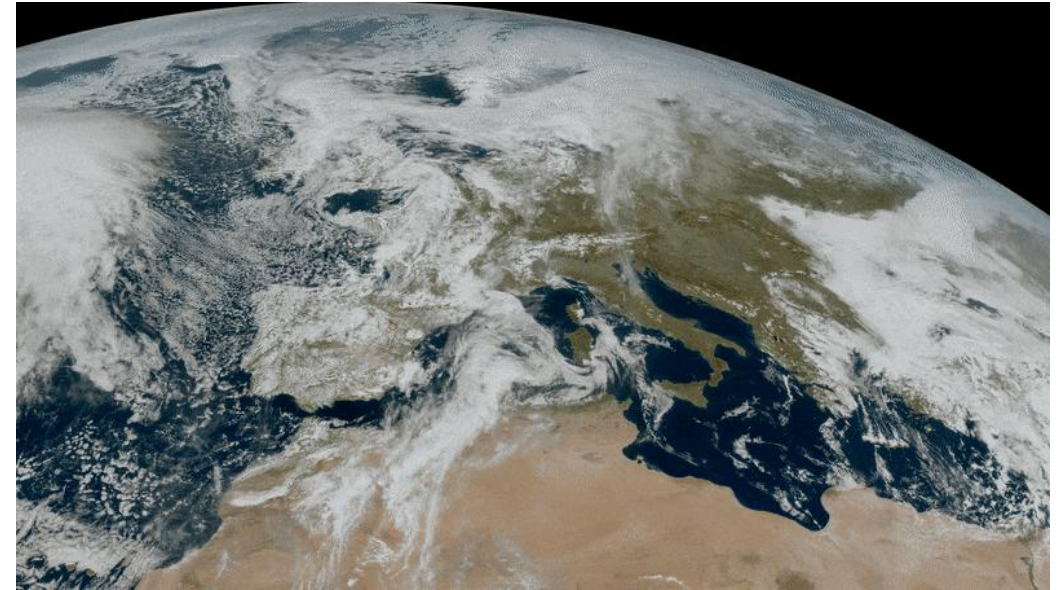
NGT ACC – Next Generation
Peak perform: 6 Pflops
Sustained HPL: **4,24 Pflops**

Consortium



Destination Earth: towards a DT for climate change adaptation

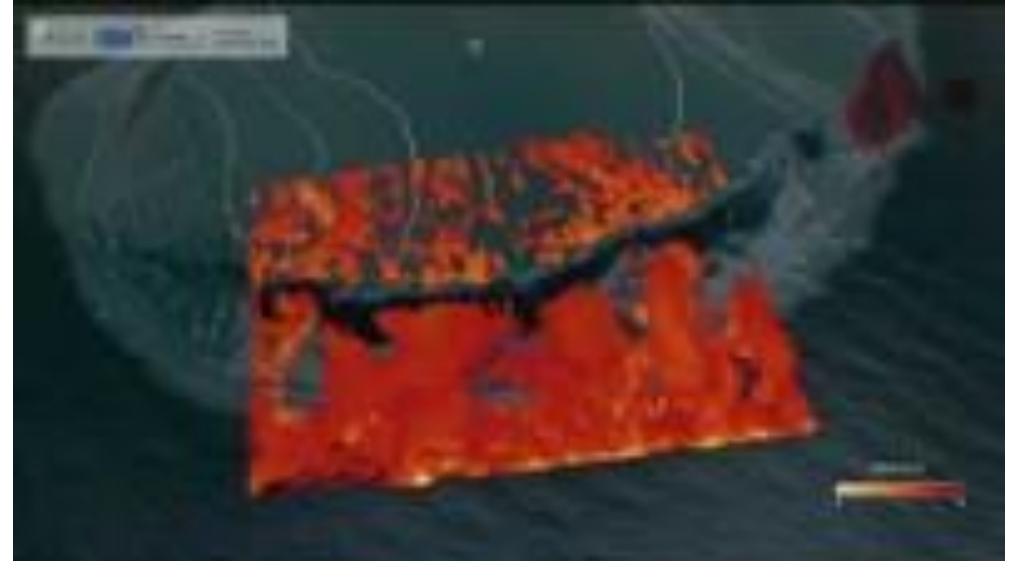
- The **Digital Twin for Climate Change Adaptation**, in which BSC plays a crucial role, is implemented by a partnership procured by ECMWF and involves 12 EU institutions.
- The **high-resolution simulations** using pre-exascale supercomputers in Europe, such as **MareNostrum 5** or LUMI.
- This Climate Digital Twin is a pioneering effort to
 - the **production of global climate projections** for the upcoming decades and
 - provide **globally consistent Earth system** and impact sector information from global to local scales.
 - <https://platform.destine.eu/>



Alya is a DT-enabling tool to simulate complex physics

- The Alya System is a Computational Mechanics code with two main features:
 - Specially designed for **running efficiently in HPC**
 - Capable of **solving different physics**, each one with its own model characteristics, in a **coupled way**.
 - All complex coupled problems solved by Alya retain the efficiency.
- Among the problems it solves are In/Compressible Flows, Non-linear Solid Mechanics, Thermal Flows, N-body collisions, etc.

ELEM
THE VIRTUAL
HUMANS FACTORY

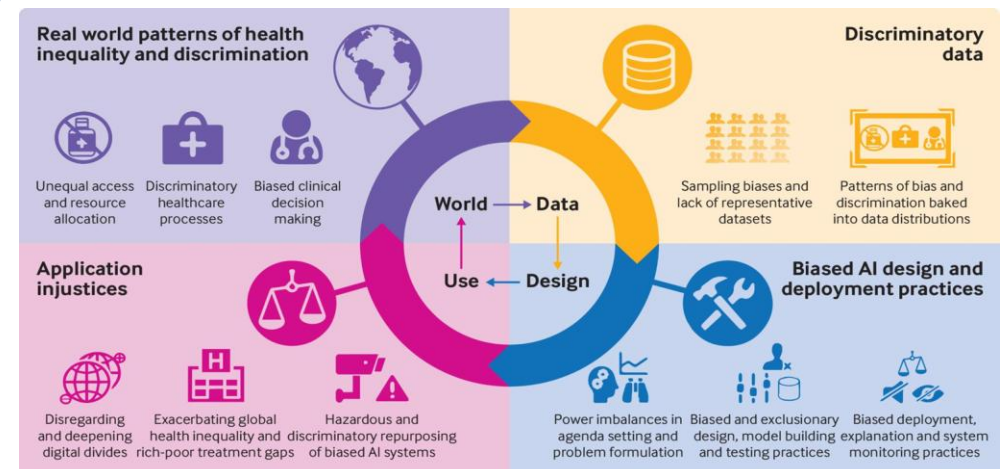


AI Factories will be used to complement digital twins

- Call for Expression of Interest to improve EuroHPC supercomputers with **AI capabilities**
 - AI-optimised supercomputer
 - Associated “AI Factories” activities and services
- AI Factories will provide the **computing power, data, support and talent**, to offer a wide and exhaustive **range of services to AI startups and researchers** needed for the development of **European generative AI models and applications**.
- Global budget: 130 – 150 M€
 - I+D services: 30 – 50 M€
 - AI Compute: 90 – 100 M€
- Partnerships: local and global



EU AHEAD – coordination evaluation of ML/AI models



AI agents are poised to revolutionise DTs

- AI agents are **autonomous software systems** that use AI to achieve specific goals and perform tasks on behalf of users, often with minimal human intervention.
 - They can **perceive their environment**,
 - make decisions**, and
 - take actions** to achieve their objectives.
- These agents **leverage technologies like large language models (LLMs)** and can be integrated into complex workflows to automate processes and improve efficiency

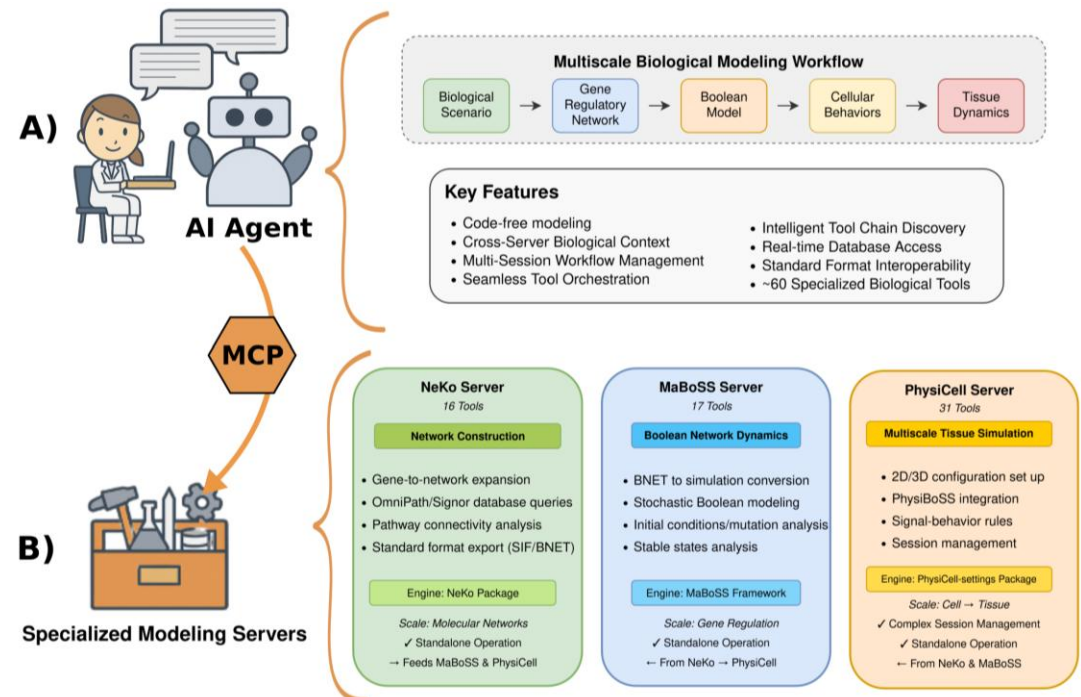
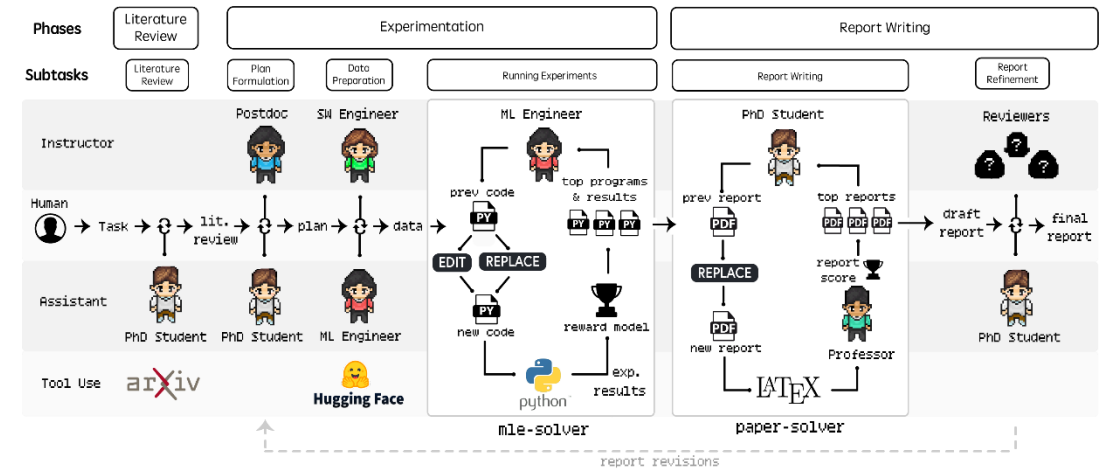
Intelligent Tool Orchestration for Rapid Mechanistic Model Prototyping: MCP Servers as AI-Biology Interfaces

Marco Ruscone, Miguel Vazquez, Alfonso Valencia

doi: <https://doi.org/10.1101/2025.09.10.675105>

Agent Laboratory: Using LLM Agents as Research Assistants

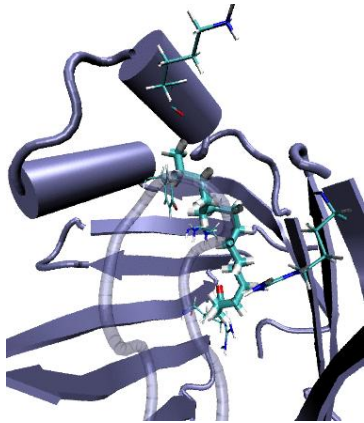
Samuel Schmidgall^{1, 2}, Yusheng Su¹, Ze Wang¹, Ximeng Sun¹, Jialian Wu¹, Xiaodong Yu¹, Jiang Liu¹, Zicheng Liu¹ and Emad Barsoum¹



BSC pioneers DT in Life Sciences

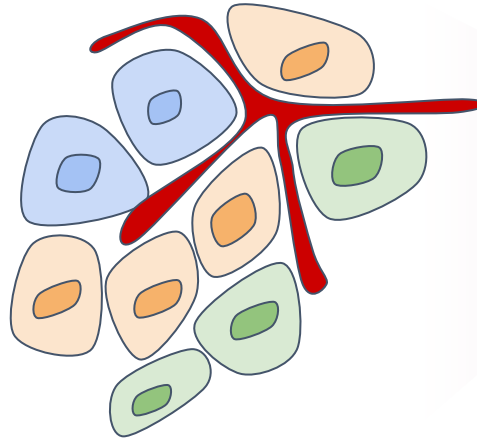
From cells to tissues and organs

PELE tool



Borrelly et al. (2005) J. Chem.
Theo. Comput

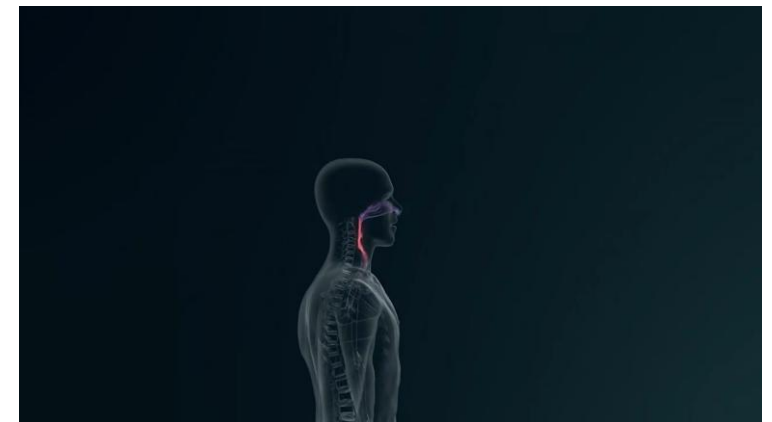
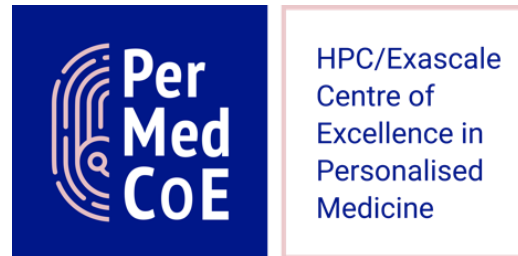
PhysiBoSS tool



ALYA tool



<https://www.bsc.es/news/bsc-news/alya-red-computational-heart>



<https://www.youtube.com/watch?v=yyBLPaU057A>

Virtual Human Twins initiative for health and care

European Virtual Human Twins

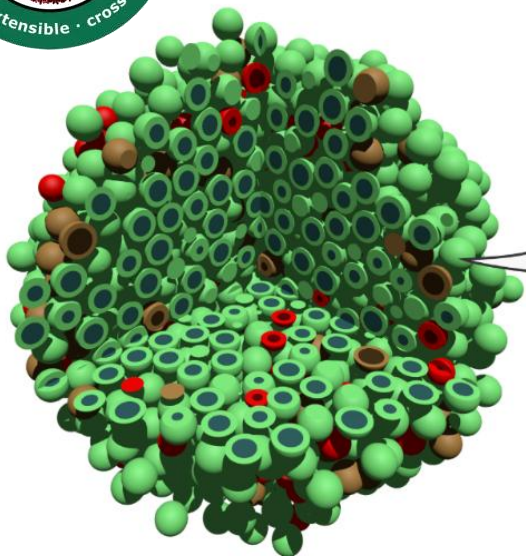
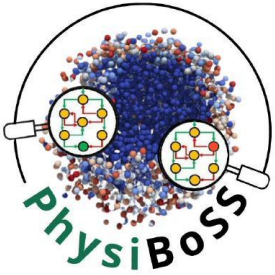


- EU Initiative Goals:
 - Foster **collaboration** among academia, industry, and innovators;
 - integrate advanced **supercomputing and AI**;
 - break silos via a **unified platform**;
 - promote **interoperability and scaling**.
- Major Milestones:
 - Launch of the **VHT Initiative** alongside MareNostrum5.
 - Signature of the Virtual Human Twins Manifesto by 75+ stakeholders.
 - 80 M€ Horizon Europe funding for integrated, multi-scale patient models.
 - 24 M€ platform for model integration and validation started in June 2025.
- Benefits of the Advanced Platform:
 - Secure,
 - open-source,
 - AI-optimised infrastructure

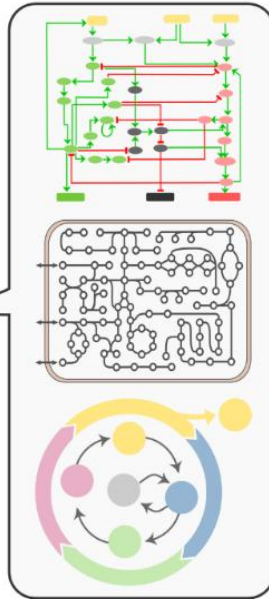




Agent-based for multiscale modelling



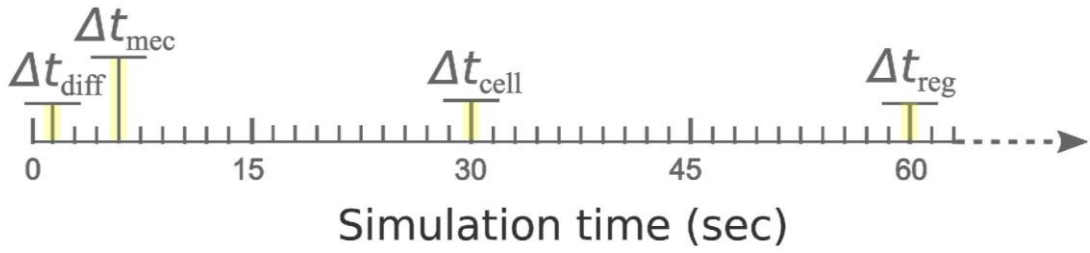
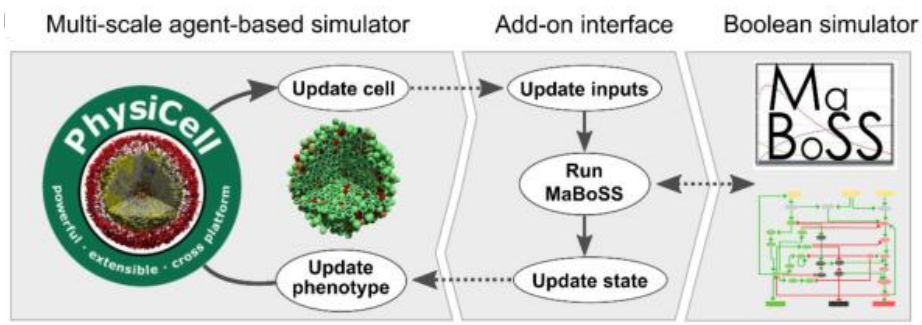
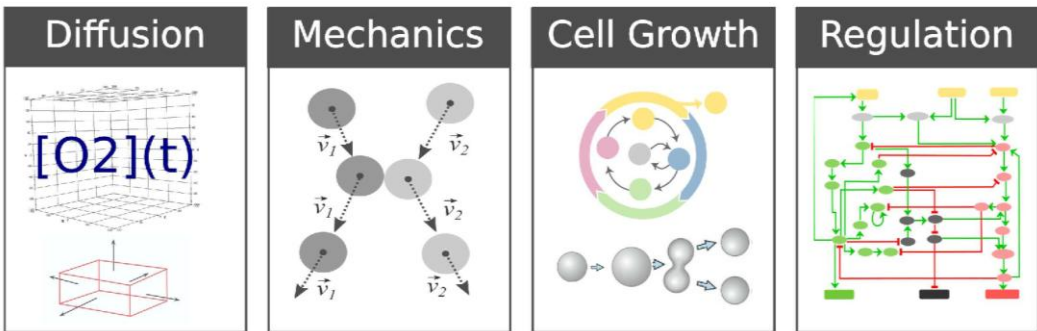
Cell population



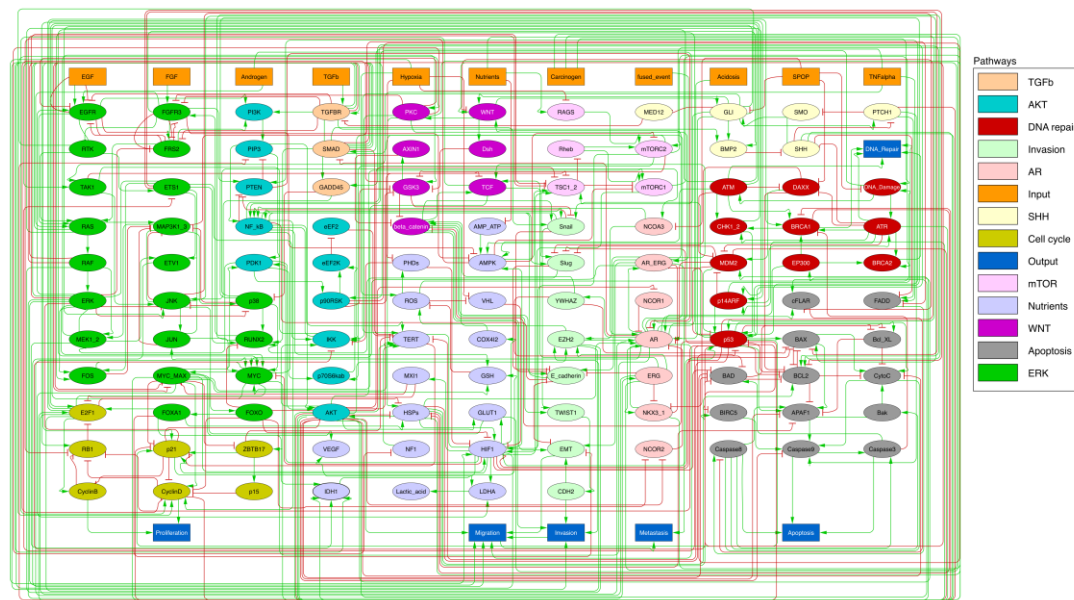
Intracellular models

Time scales

- Δt_{diff} : (diffusion/transport): 0.01 min
- Δt_{mec} : (cell movement): 0.1 min
- Δt_{cell} : (cell processes): 6 min
- Δt_{reg} : (Boolean simulation): 10 min



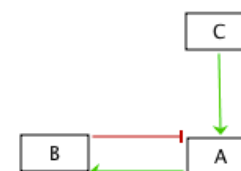
Boolean modelling for signalling pathways



Montagud, et al. (2022). *eLife*, 11:e72626

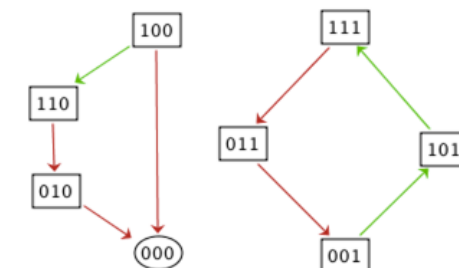
- Set of **discrete variables as abstractions of activity level** linked by logical rules as signed interactions

Regulatory graph & logical rules

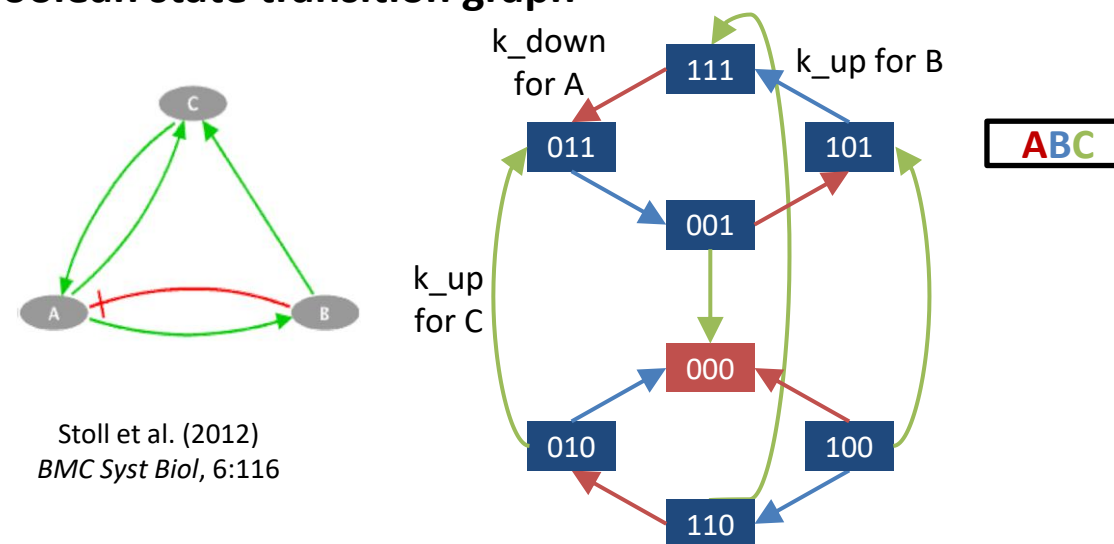


$A = !B \ \& \ C$
 $B = A$
 $C = \text{input}$

Solutions



Boolean state transition graph

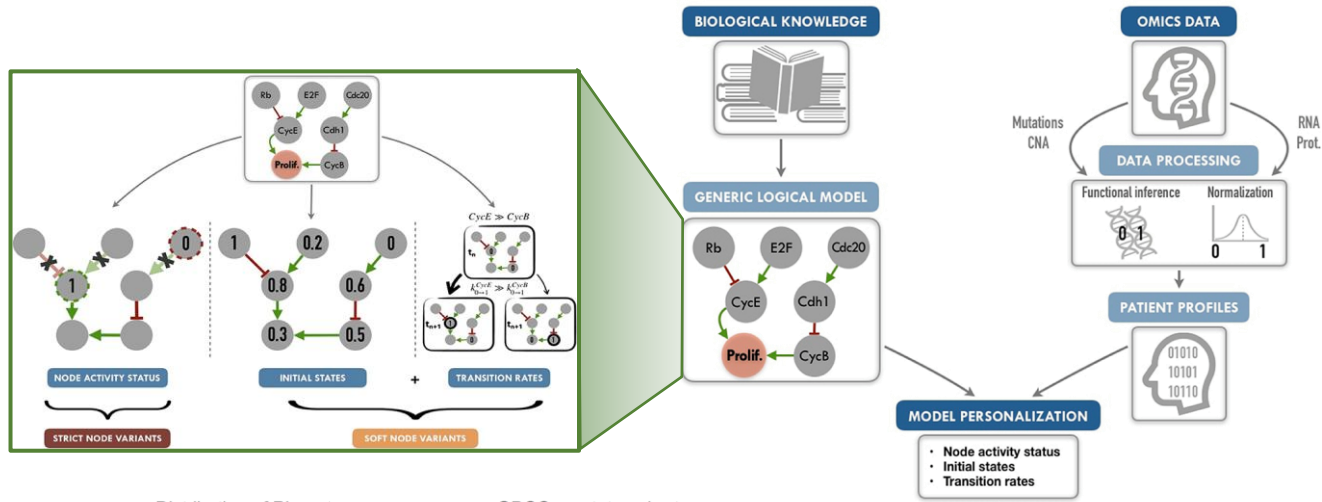


Stoll et al. (2012)
BMC Syst Biol, 6:116

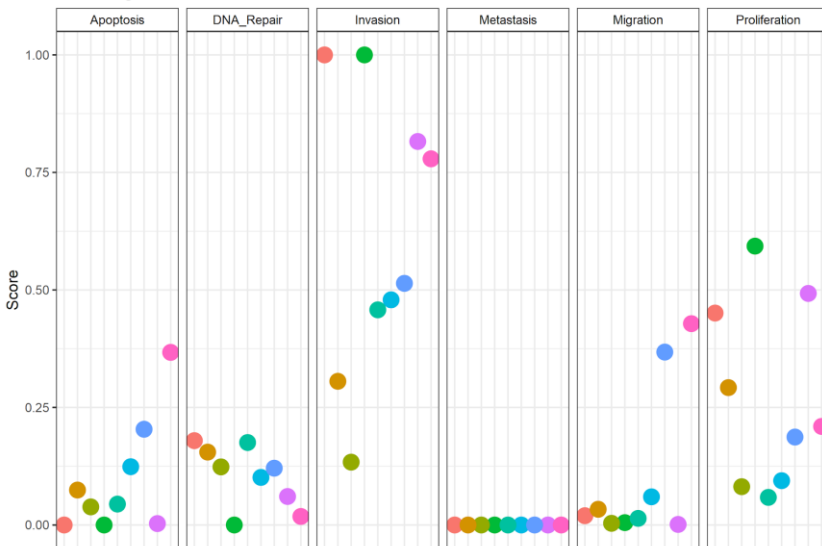
- Continuous time Markov process** on the Boolean transition state space
 - Each **Boolean state** has an associated **probability**
 - Rate of change** associated to each transition
 - rate up and rate down
 - Stochasticity, time, **probabilities**, ...

Simulating patient-specific models

Personalisation of intracellular models

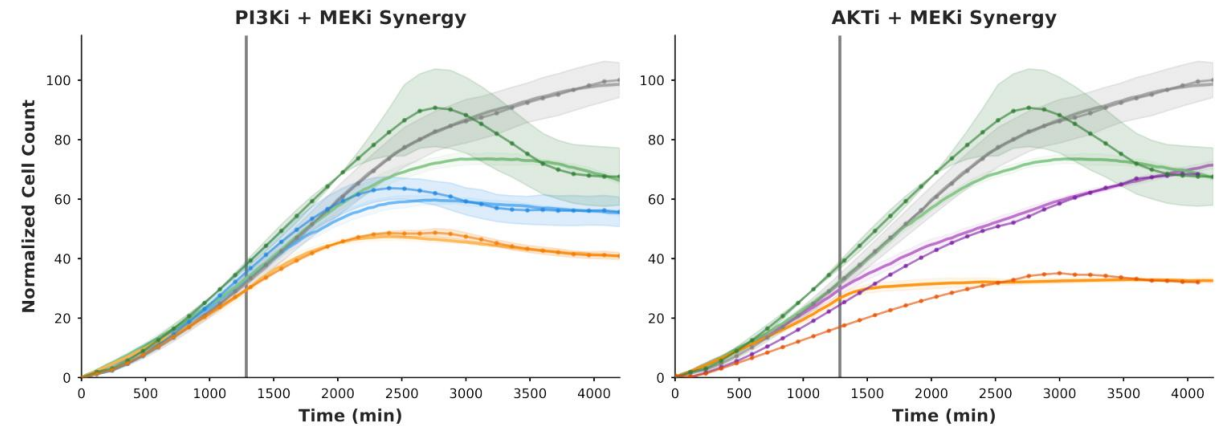
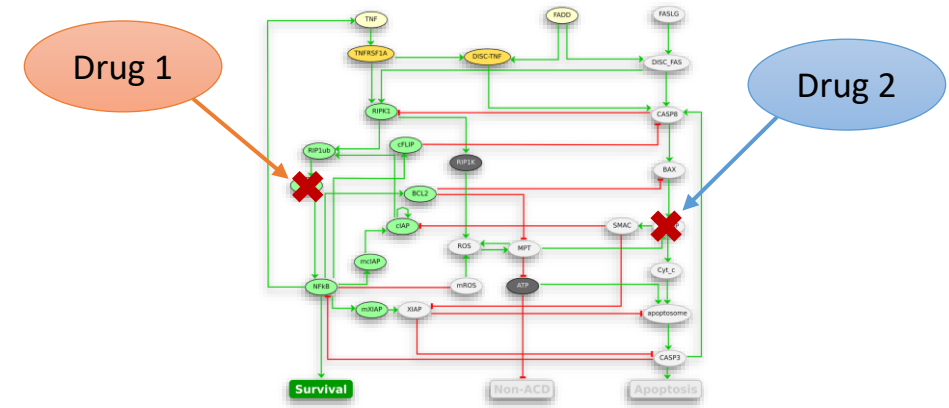


Distribution of Phenotypes scores across GDSC prostate cohort, using mutation as discrete data, RNA as continuous data and random initial conditions



Béal et al. (2019) *Frontiers in Physiology*, 9:1965

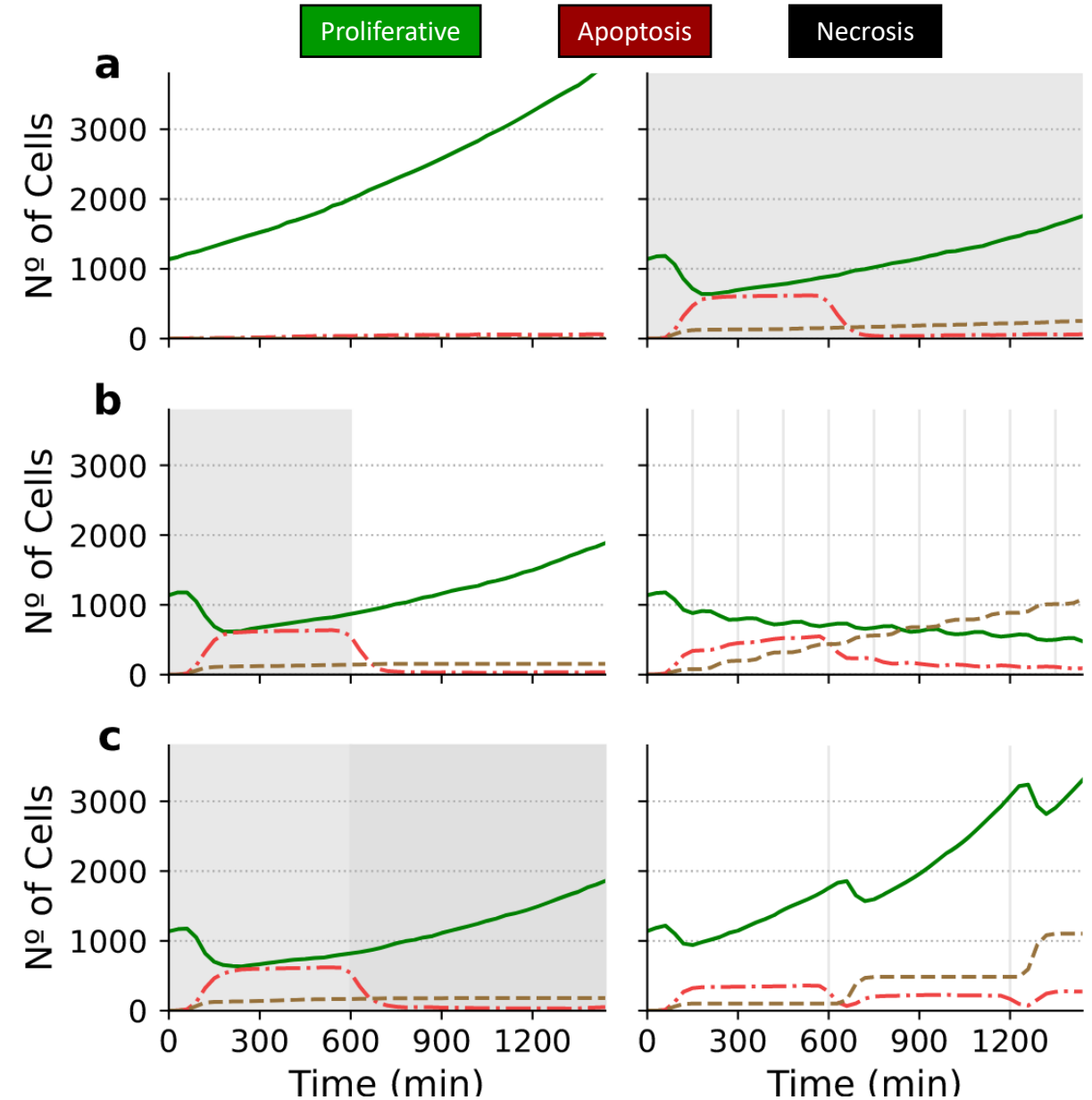
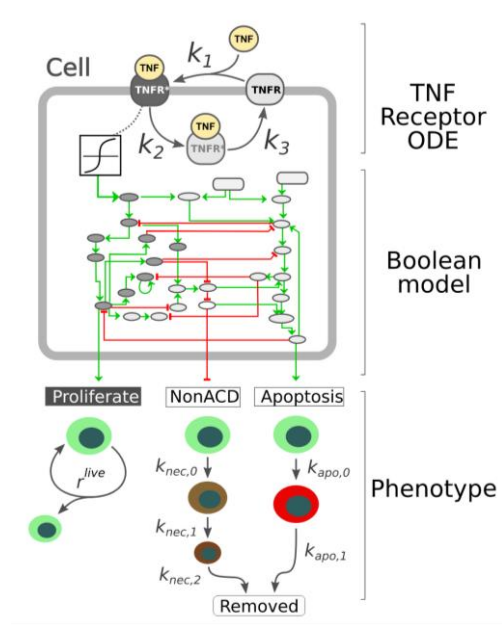
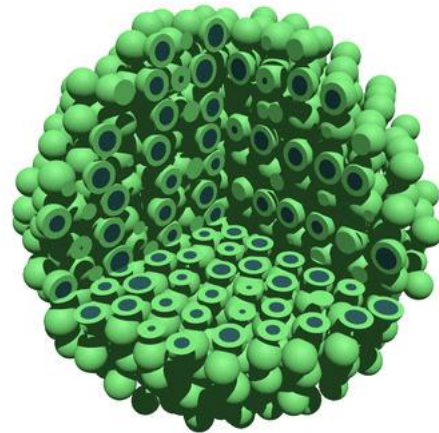
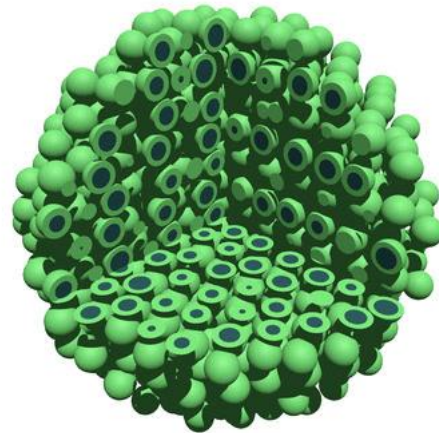
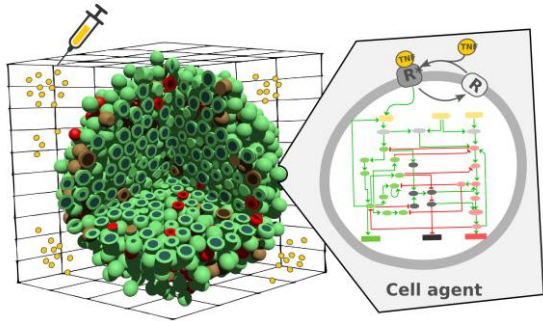
Different combinations of drugs



Emergent prediction of drug synergy from a model calibrated on single-agent responses.

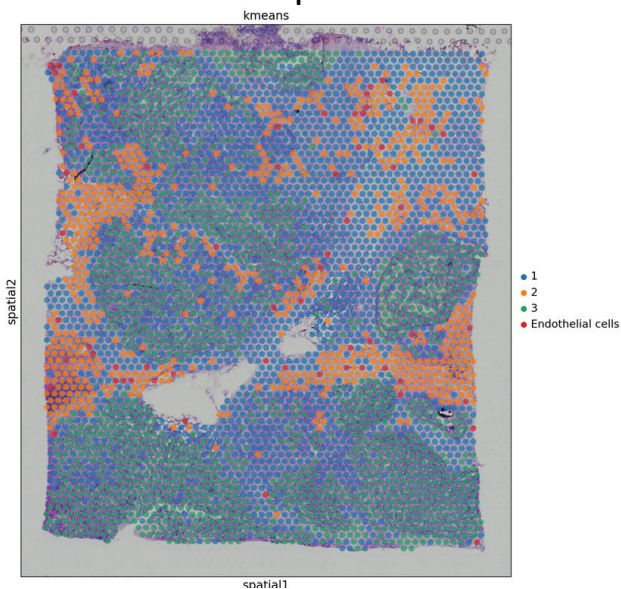
Hayoun-Mya et al. *under preparation*

PhysiBoSS allows for finding optimal drug regimes

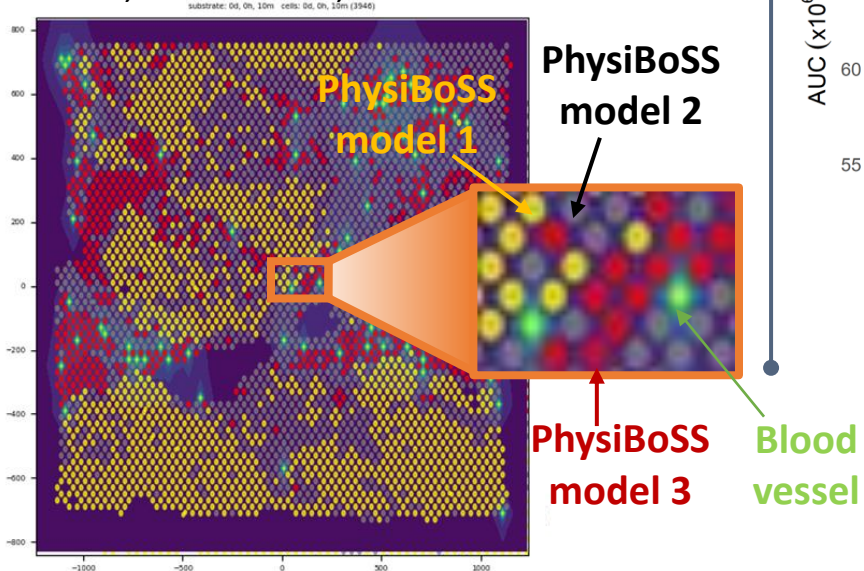


Integrating spatial omics in PhysiBoSS

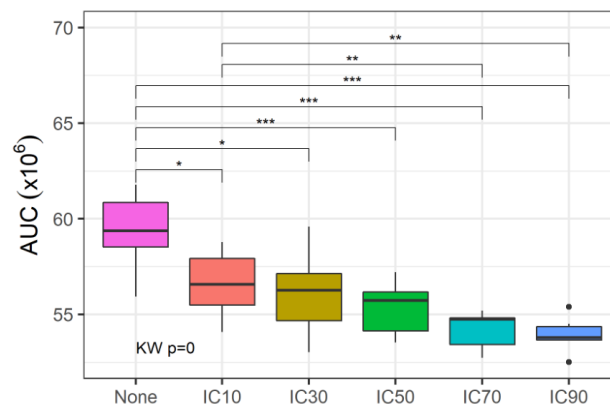
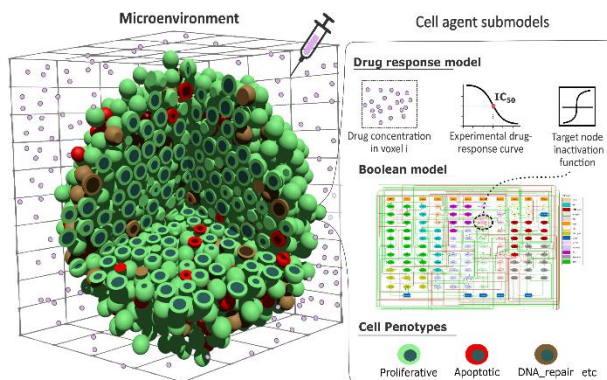
Breast cancer spatial omics



ECM, O2 sources, different cell clusters

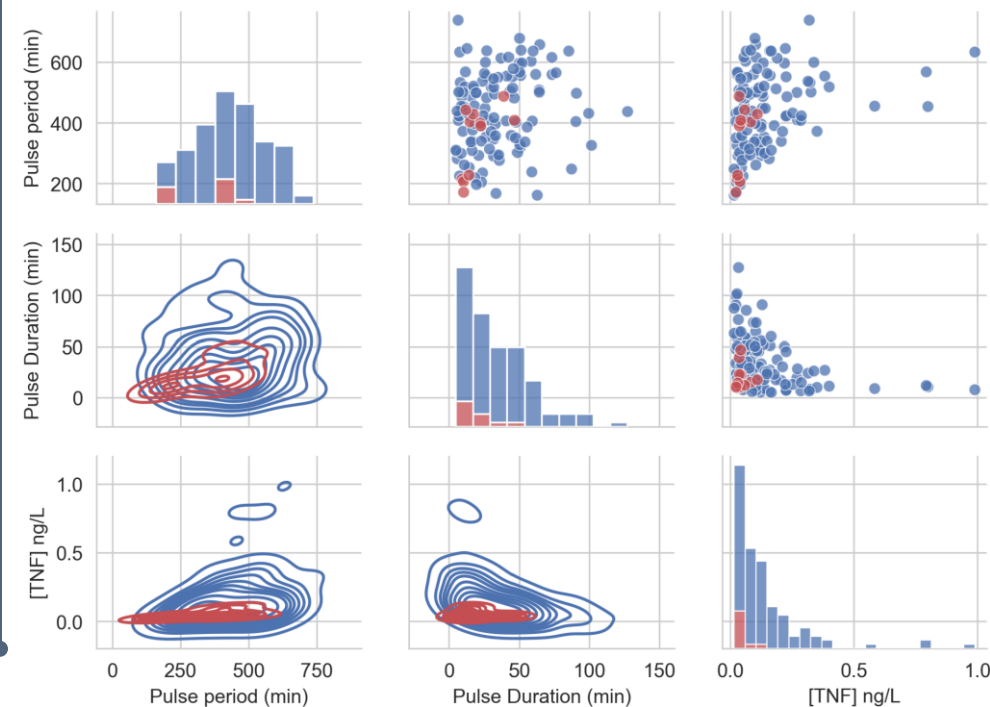
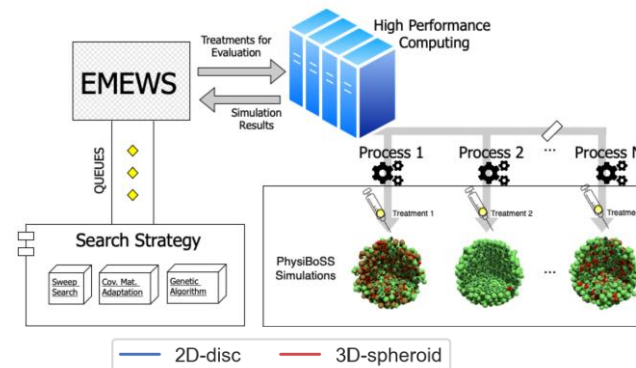


Patient-specific drug response curves



Ponce-de-Leon et al. (2023)
npj Syst Biol Appl, 9:54

High-Throughput Computing to parametrise the simulations



Akasiadis et al. (2022) *Computational Intelligence*, 38(4), 1–23
Ponce-de-Leon et al. (2022) *Frontiers in Molecular Biosciences*, 9, 836794

20 YEARS



**Barcelona
Supercomputing
Center**
Centro Nacional de Supercomputación

Thank you!

Dept. Life Sciences, BSC
over 250 scientists, engineers
and students



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