

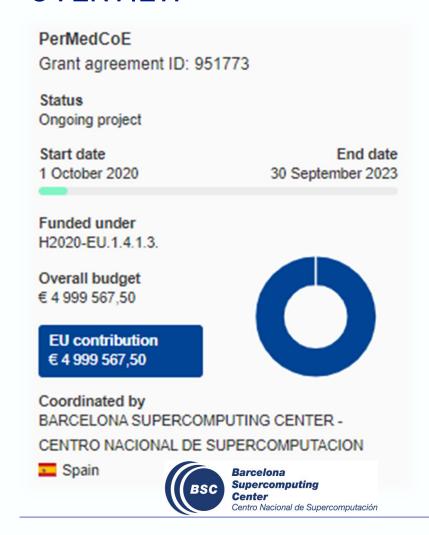
HPC/Exascale Centre of Excellence in Personalised Medicine

PROJECT OVERVIEW



The PerMedCoE project has received funding from the European Union's Horizon 2020 research and innovation programme under the grant agreement N°951773

OVERVIEW







PARTNERS





University of Ljubljana

























MISSION

PerMedCoE is the HPC/Exascale Centre of Excellence for Personalised Medicine in Europe

Our motivation

The performance of current simulation software is still insufficient to tackle medical problems such as tumour evolution or patient-specific treatments.

Simulation of cellular mechanistic models are essential for the translation of omic data to medical relevant actions and these should be accessible to the end-users in the appropriate environment of the PerMed-specific big confidential data.

Our vision

The challenge is to develop a sustainable roadmap to scale-up the essential software for the cell-level simulation to the new European HPC/Exascale systems.

Our goal is to provide an efficient and sustainable entry point to the HPC/Exascale-upgraded methodology to translate omics analysis into actionable models of cellular functions of medical relevance.







PerMedCoE aims to integrate PerMed into the new European HPC/Exascale ecosystem

Physiological-level models (Fluid dynamics – CompBioMed)

Cell-environment interactions

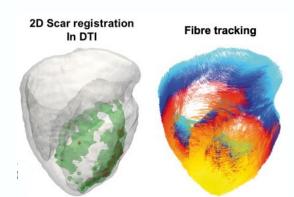
Cell-cell interactions

Metabolic pathways

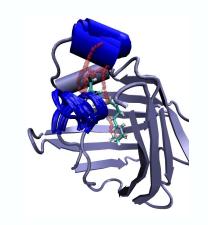
Signalling pathways

Cell-level models

Atomic-level models (Molecular dynamics – BioExcel)



Europace. 2019 May, 21(5): 822–832.

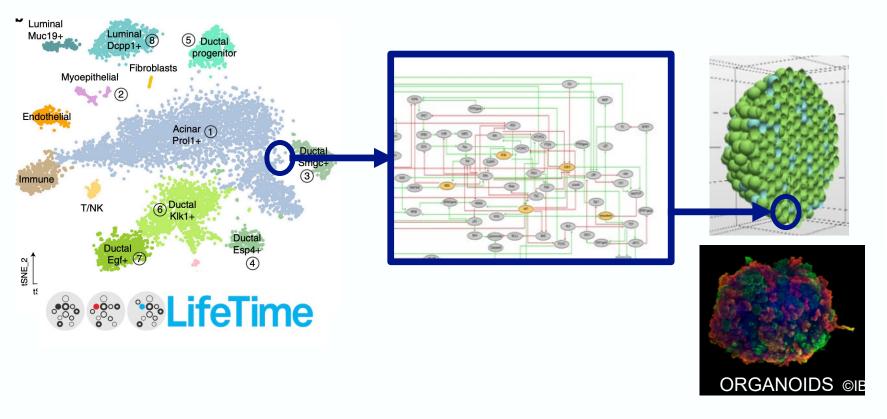


J. Chem. Theo. Comp. 2005, 6, 1304-1311

From Single Cell Data to Models



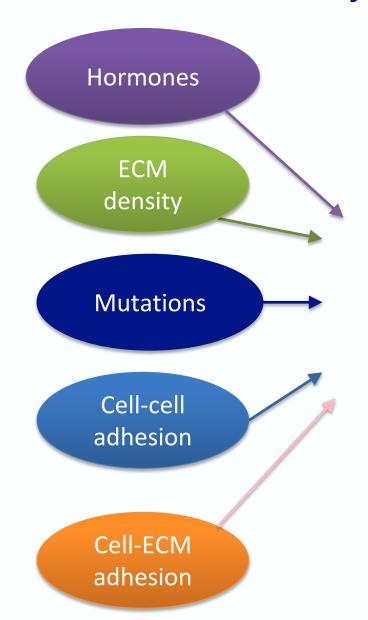
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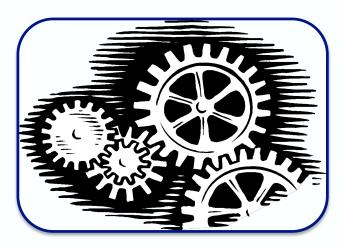




"Replace the current generation of bioinformatics methods with cellular models, providing mechanistic descriptions and testable hypotheses, instead of current statistical approximations and intuitive descriptions"

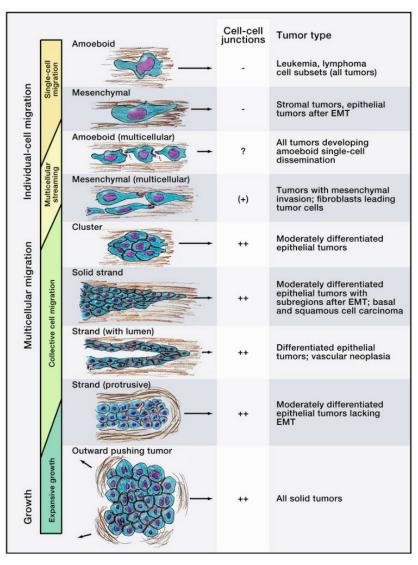
The motivation: Genotype-to-phenotype modelling





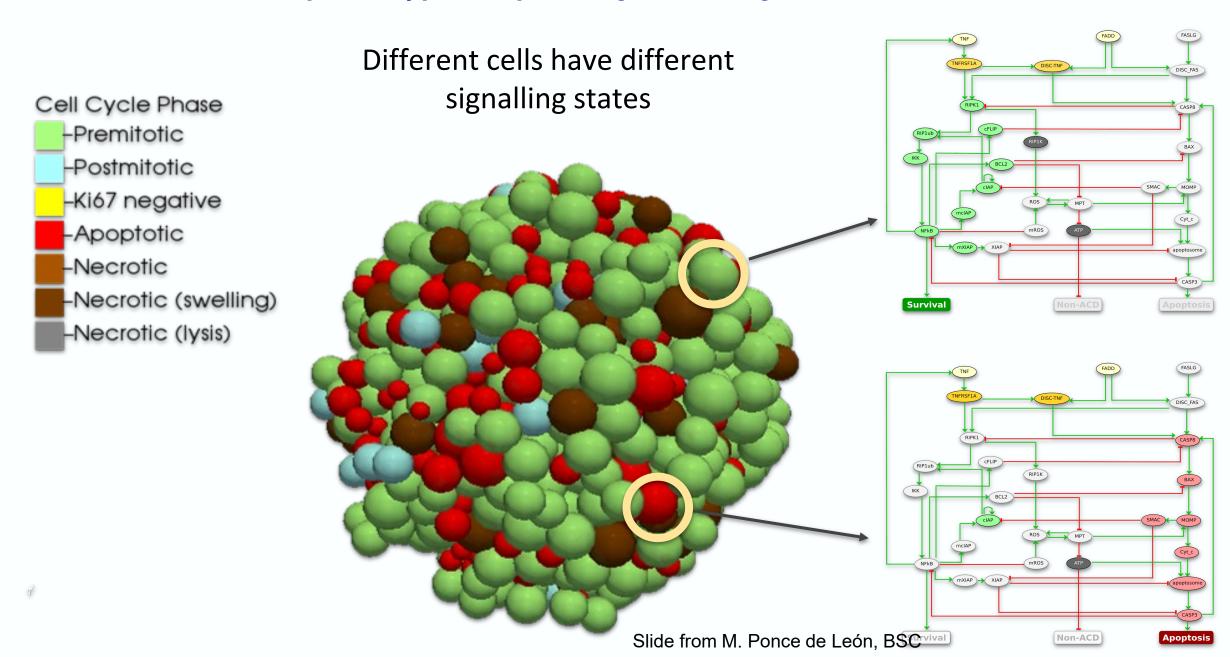
Multiscale modelling

Gene mutations
Signalling pathways
Cell - environment
ECM modification



Friedl and Alexander, Cell, 2011

Cells have different phenotypes depending on their genes' activation

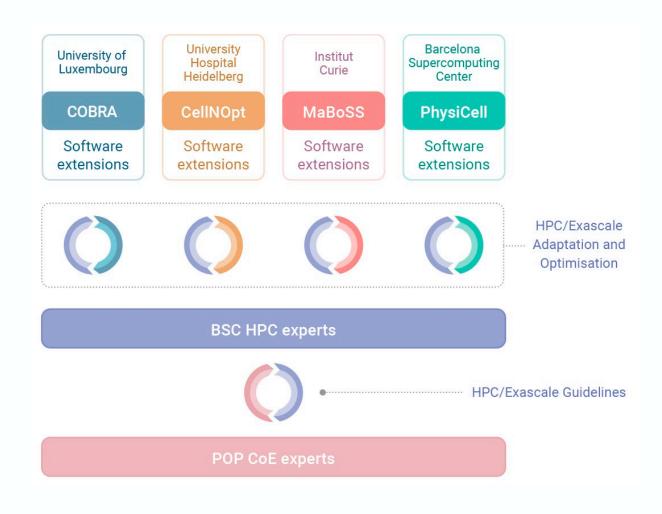


CORE APPLICATIONS

PerMedCoE optimises key software for cell-level simulations to the new preexascale platforms

The PerMedCoE four core applications are:

- COBRA for the simulation of cellular metabolism at genome-scale
- CellNOpt for modelling signal transduction networks
- MaBoSS for stochastic simulations of Boolean models
- PhysiCell an agent-based modelling framework for simulating cell-cell interactions





USE CASES

PerMedCoE works on five use cases to drive the development of cell-level simulations



Tumour Evolution Based on Single-Cell Omic and Imaging



USE CASES

PerMedCoE works on five use cases to drive the development of cell-level simulations



COVID-19
Multiscale
Modelling of
the Virus and
Patients' Tissue



Cancer Diagnosis
Based on Omics
Information



Tumour Evolution Based on Single-Cell Omic and Imaging



Drug Synergies for Cancer Treatment



Personalised
Modelling of Groups
of Rare-Disease
Related Patients



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COVID-19
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Cancer Diagnosis
Based on Omics
Information



Cancer Research

Biotech & Omic to

Organois based



Tumour Evolution Based on Single-Cell Omic and Imaging



Clinical Trials
/
Pharma

Drug Synergies for Cancer Treatment



Personalised
Modelling of Groups
of Rare-Disease
Related Patients

Rare Diseases
/
EU Health Systems &
Patient Organisation



THANK YOU

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