



# From multiscale modelling to digital twins

Life Sciences at FIAS

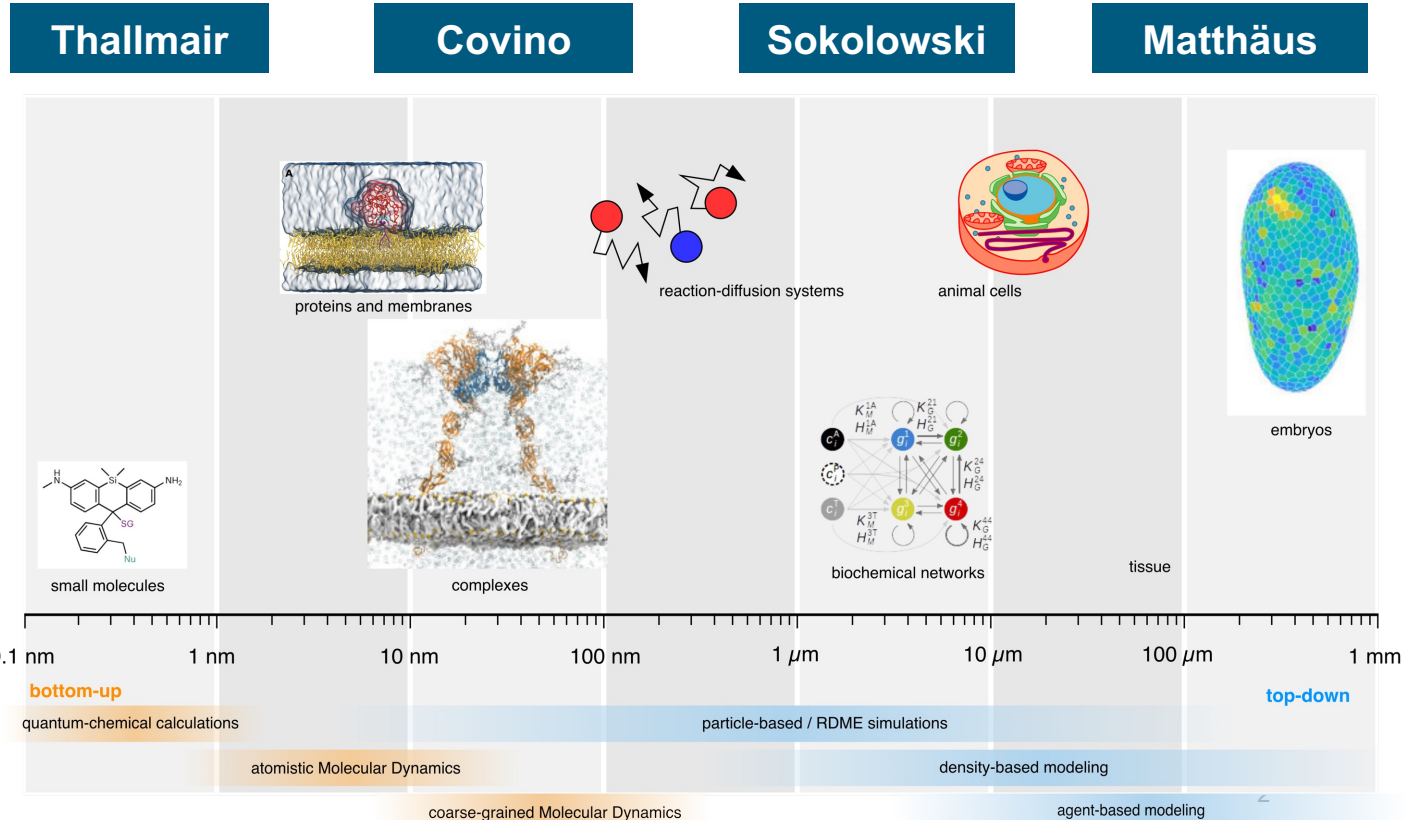
# Multi-scale modelling and simulations of biological systems

**CMMS**

Multi-scale modelling

Integration of theory and experiments

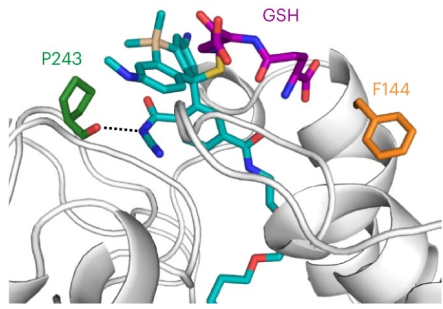
High-performance computing



# Thallmair: Controlling and inhibiting protein function

## Development of chemical probes

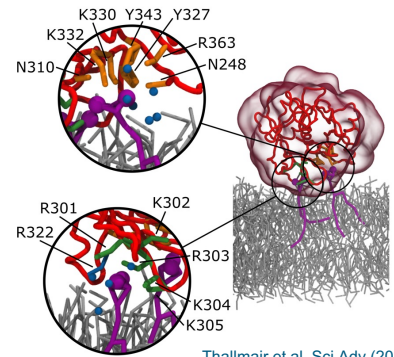
Development and characterization of a glutathione-sensing platform for live-cell imaging



Emmert et al. Nat Chem (2023)

## Protein-Lipid interactions

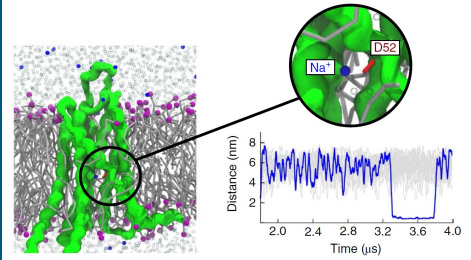
Membrane binding mechanism for decoding of spatio-temporal phospholipid signals in cells



Thallmair et al. Sci Adv (2022)

## Coarse-grained force-field development

Reparameterization of the Martini force field for general purpose coarse-grained molecular dynamics

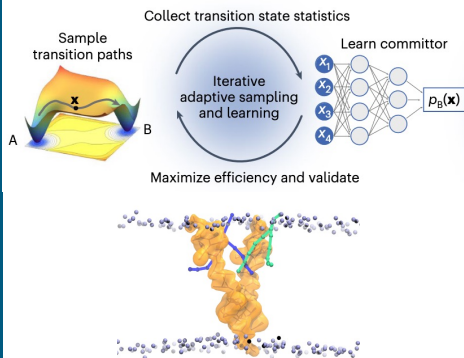


Souza et al Nat Meth (2021)

# Covino: AI for biomolecular simulation and analysis

## Physics-based simulators and AI

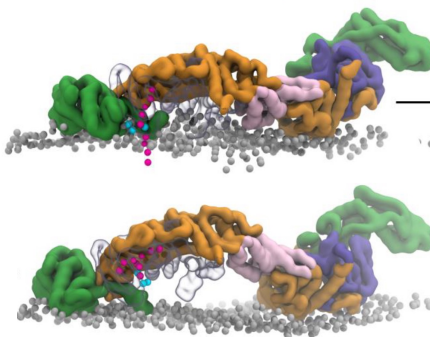
Integration of AI and MD to enhance simulations of structural reorganization



Jung et al. Nat Comp Sci (2023) Lazzeri et al. JCTC (2023)

## High-Performance Simulation

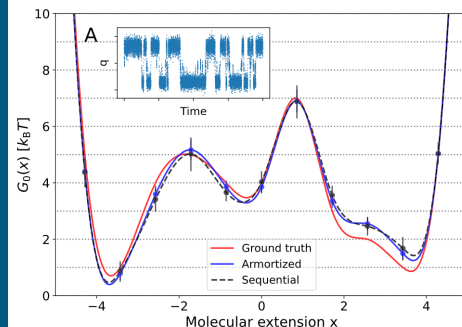
Atomistic and coarse-grained MD of molecular mechanisms in proteins and membranes



Nishimura et al. Sci Adv (2023)

## AI & Bayesian inference of exp.

Extracting mechanistic models from spectroscopy and microscopy experiments

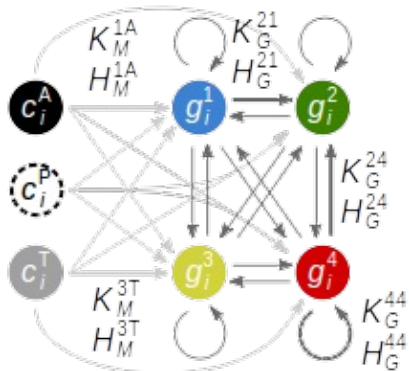


Dingeldein et al. Mach Learn Sci Tech. (2023)

# Sokolowski: Noise-control in cells and tissues

## Biophysics Theory

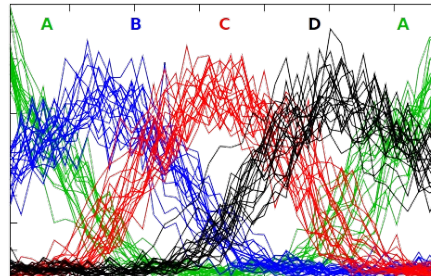
- Statistical physics
- Information theory
- Biochemical networks



Sokolowski, Gregor, Bialek, Tkačik (2023)

## High-Performance Simulation

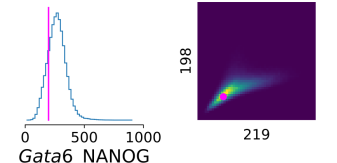
- Spatial-stochastic models
- Event-driven algorithms
- Tissue development



Majka, Becker, ten Wolde, Zagorski, Sokolowski (2023)

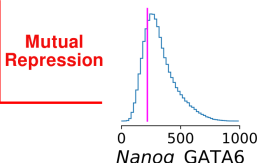
## Optimization, Inference & AI

- Simulation-based inference (SBI)
- Stoch. optimization
- Rare-event sampling



Mutual Repression

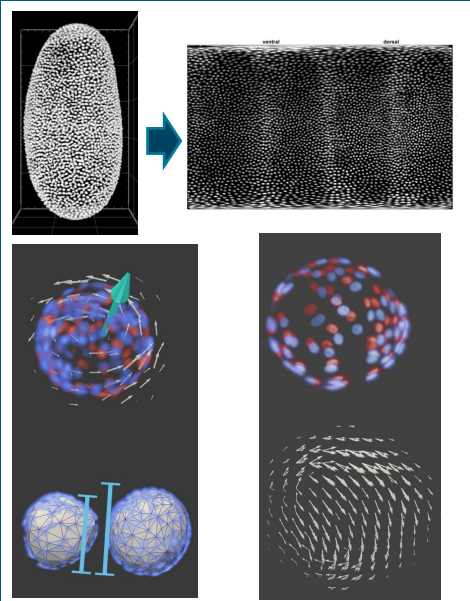
Stoch. mouse embryo model inferred via SBI



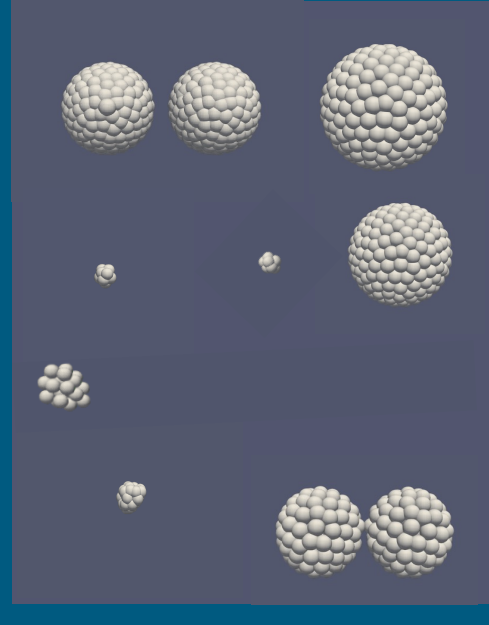
Ramírez-Sierra & Sokolowski (2024)

# Matthäus: Modeling of multi-cellular systems

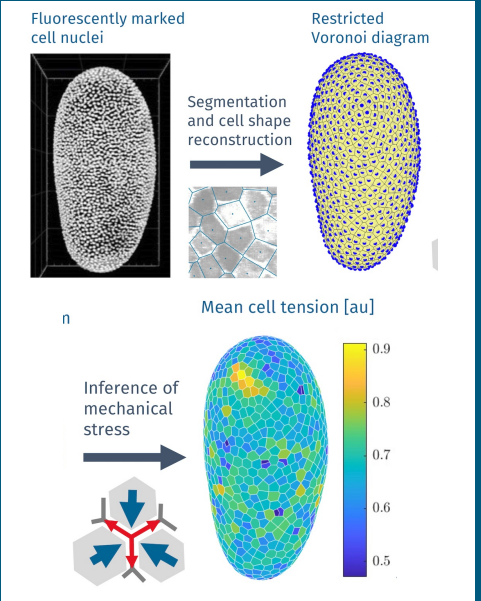
## Analysis of 3D-time-lapse microscopy



## Agent based modelling



## Force inference





# A community of passionate scientists

High scientific productivity (last 3 years):

- Publications: 37
- Preprints: 11

High-impact and international visibility:

- Sci Adv, Nature Methods, Neuron

**Enthusiastic and dedicated community of early career researchers**



## Research consortia

Cluster of Excellence “SCALE”

SFB “Membrane-Associated Protein Assemblies, Machineries and Supercomplexes”

IMPRS “Cellular Biophysics”

SFB “Molecular principles of Light-control in complex environments” (evaluation)

SFB “Acute and chronic liver failure” (preproposal)

LOEWE Schwerpunkt “Lipid Space” (preproposal)

**Deep integration in the Frankfurt landscape**

# Next generation multi-scale modelling: the digital twin

Quantitative in silico cell biology experiments

- Verifiable predictions
- Design of in vitro and in situ experiments

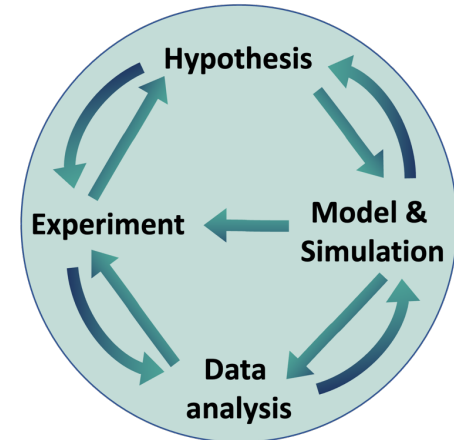
**Strategy:** developing **Digital Twins**

- Bridging multi-scale descriptions
- Simulation intelligence: AI, HPC, and physics-based modelling
- Doctoral research school for methodological development

**Full scientific and strategic support to the Cluster of Excellence SCALE**

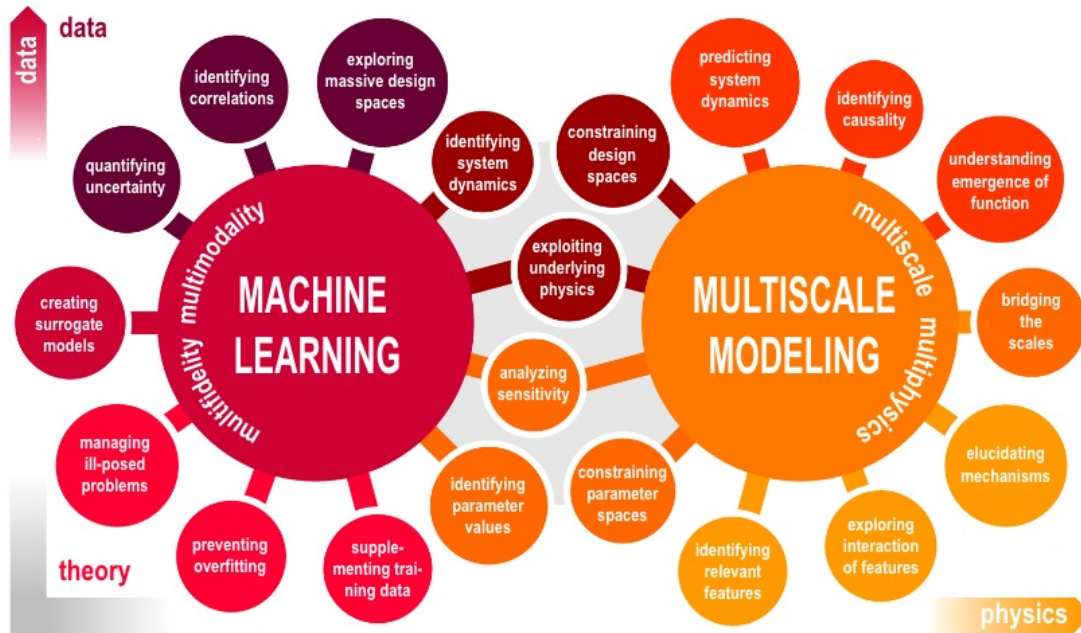


**Flagship project:  
DT of Subcellular Compartments**





# Primed for the next phase of digital biology



Aber et al. Nature Digital Medicine (2019)

## Digital Twin

Integration of pipelines, datasets, visualization techniques, physics-based simulations, and AI4science.

FIAS is **uniquely** equipped to tackle the methodological challenges that will lead us to developing **quantitative in silico biology**