

Spanish congress of Chemical Engineering Departments
Barcelona., September 9th 2011t.S

**Multiscale modeling of process dynamics in
large-scale bioreactors**

Spanish congress of Chemical
Engineering Department.

Matthias Reuss

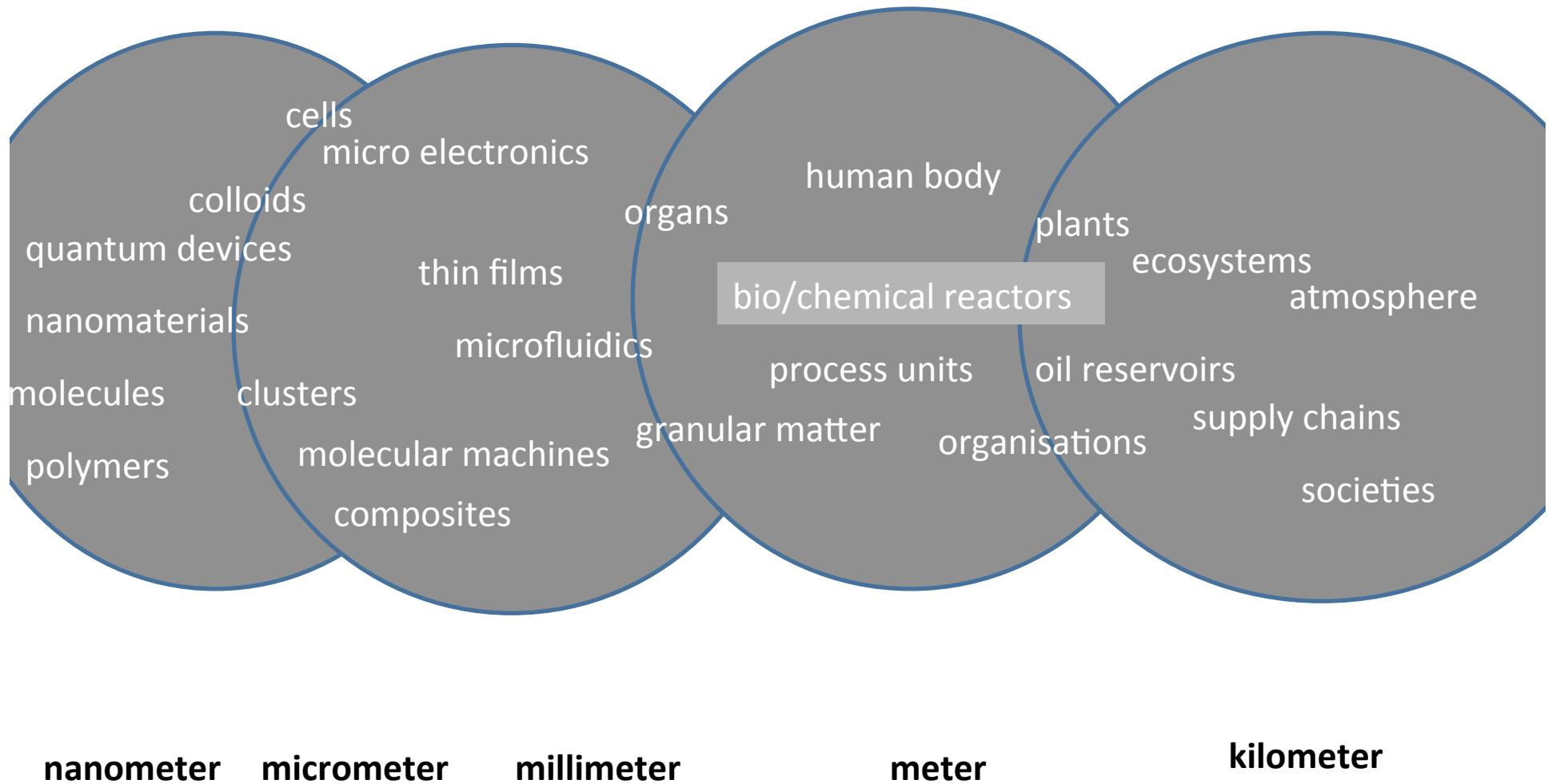
Centre Systems Biology (CSB), University Stuttgart, Germany
NGI Visiting Stipendiate, The Netherlands

**„Benchmarking the Research
Competitiveness of the
United States in Chemical Engineering“**

2007

Range of length scales covered by chemical engineering

(according to Julio M. Ottimo, AIChE Journal 2011)

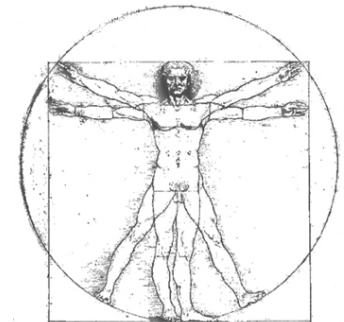
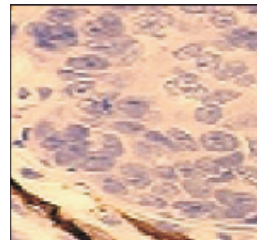
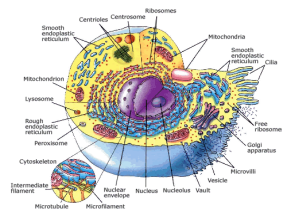
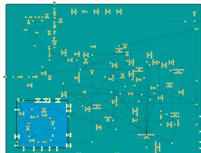
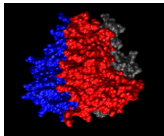


Multi-scale modeling in biomedicine

1nm

1m

10 orders of magnitude



proteins networks cells

tissues

organs

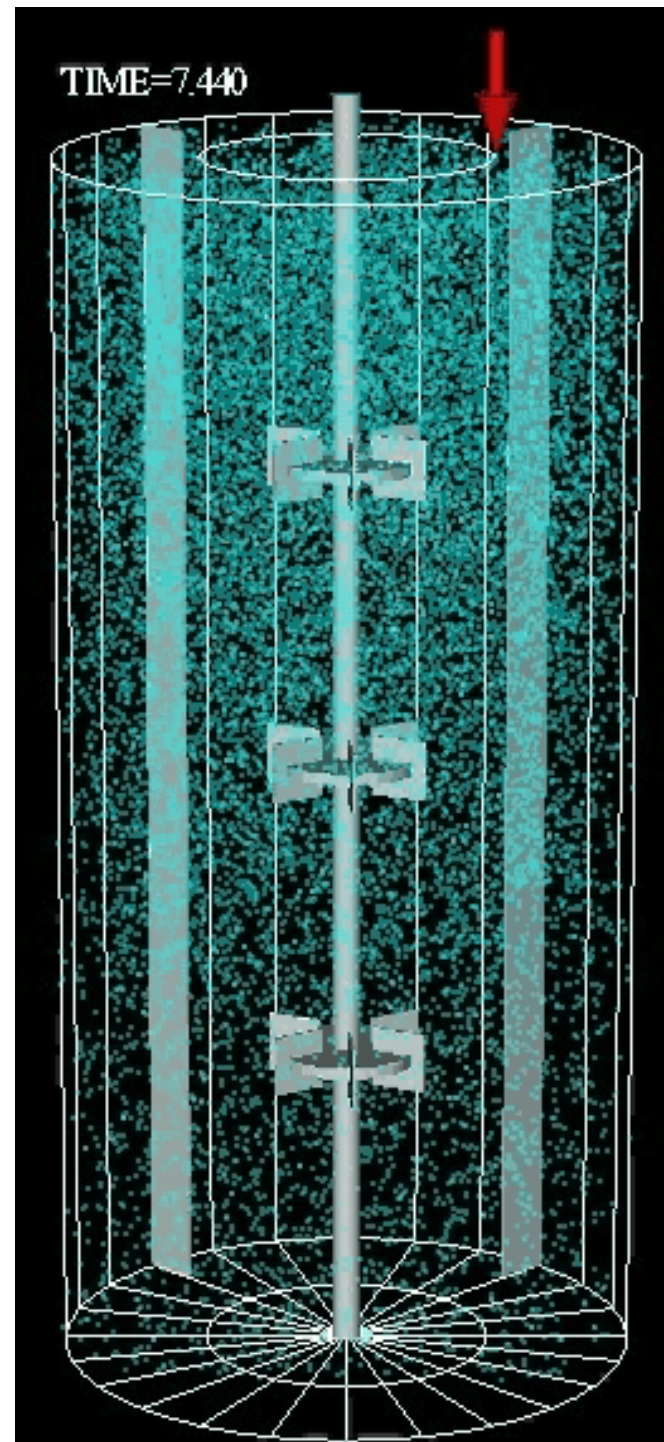
human body

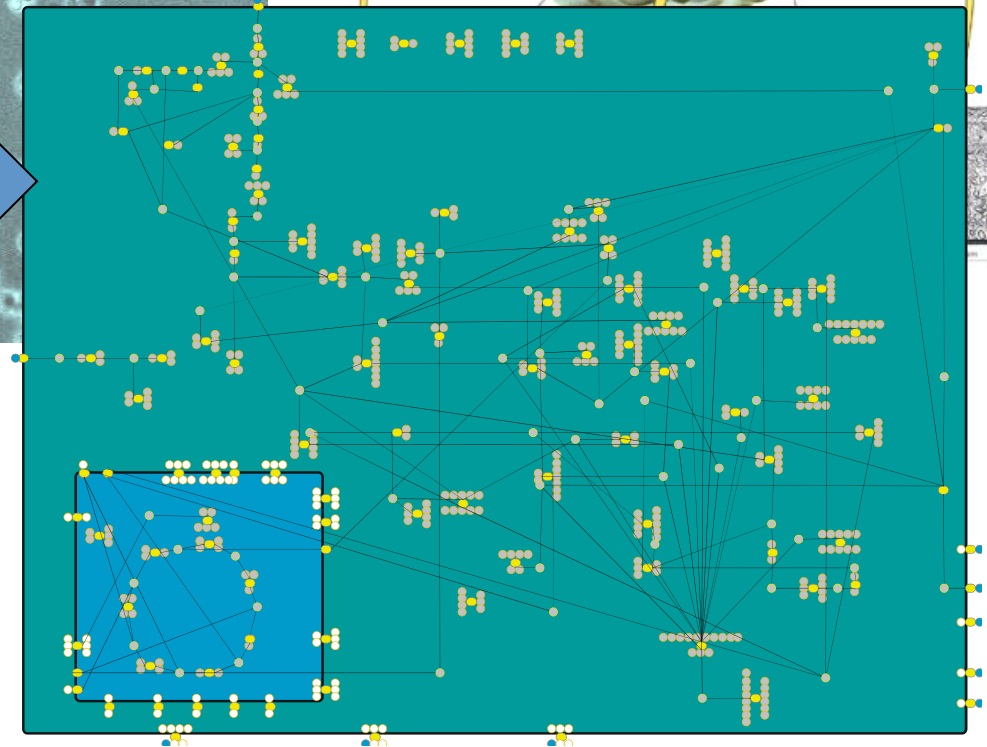
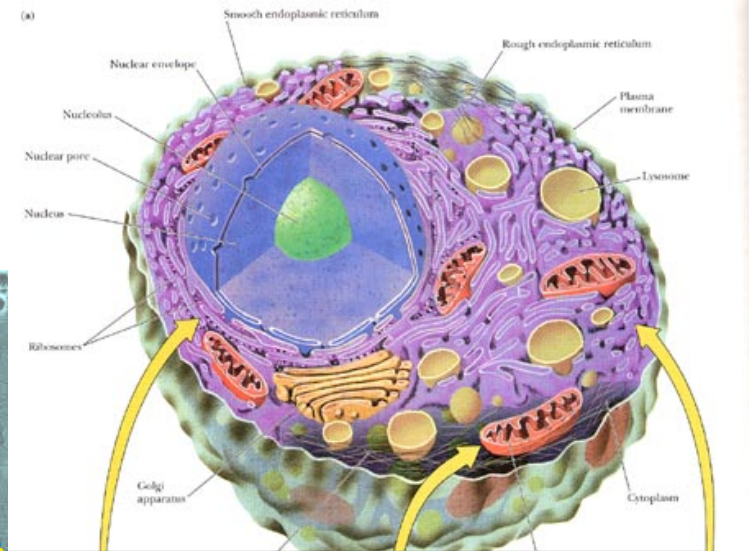
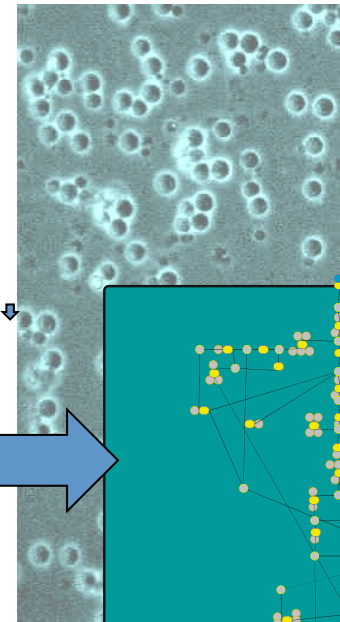
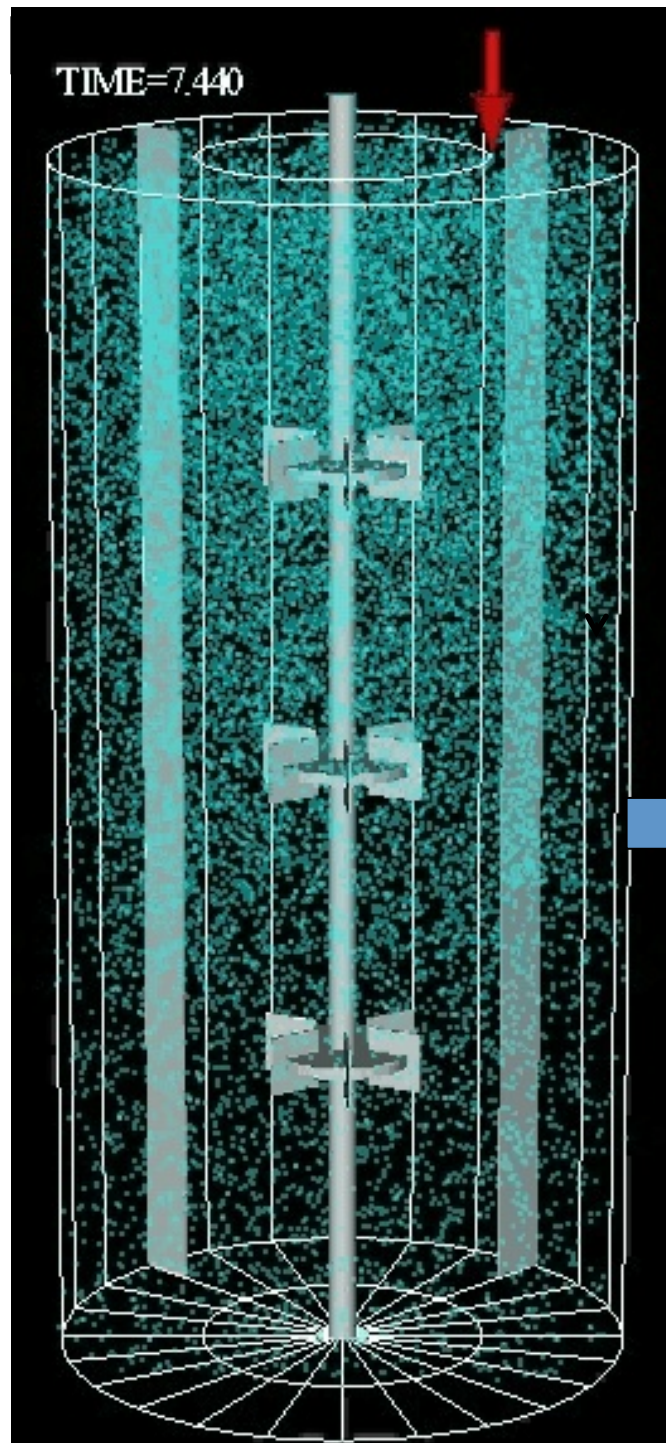
Temporal scales

$1\mu\text{s}$ (ion channel gating)

10^9s (human life)

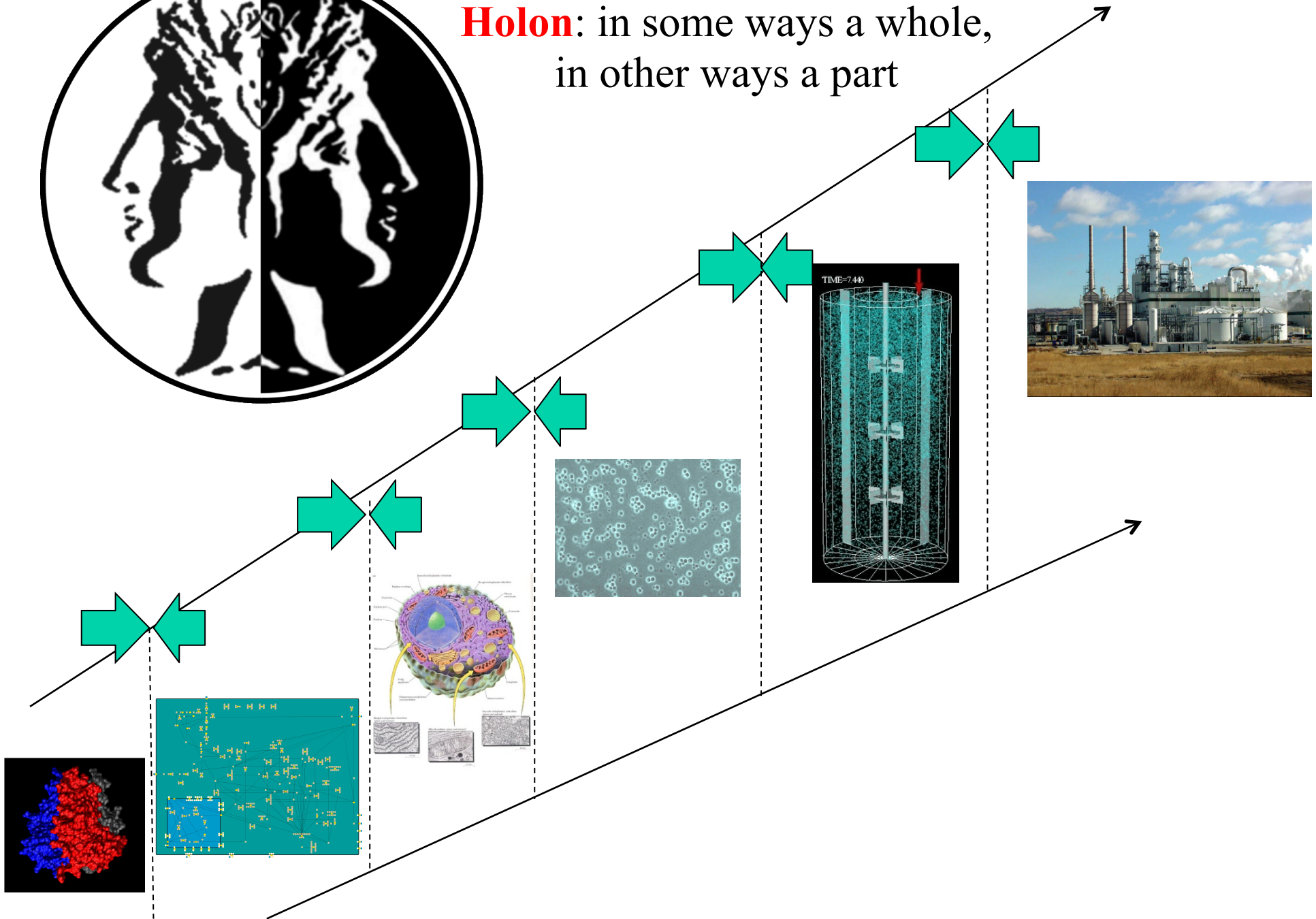
16 orders of magnitude

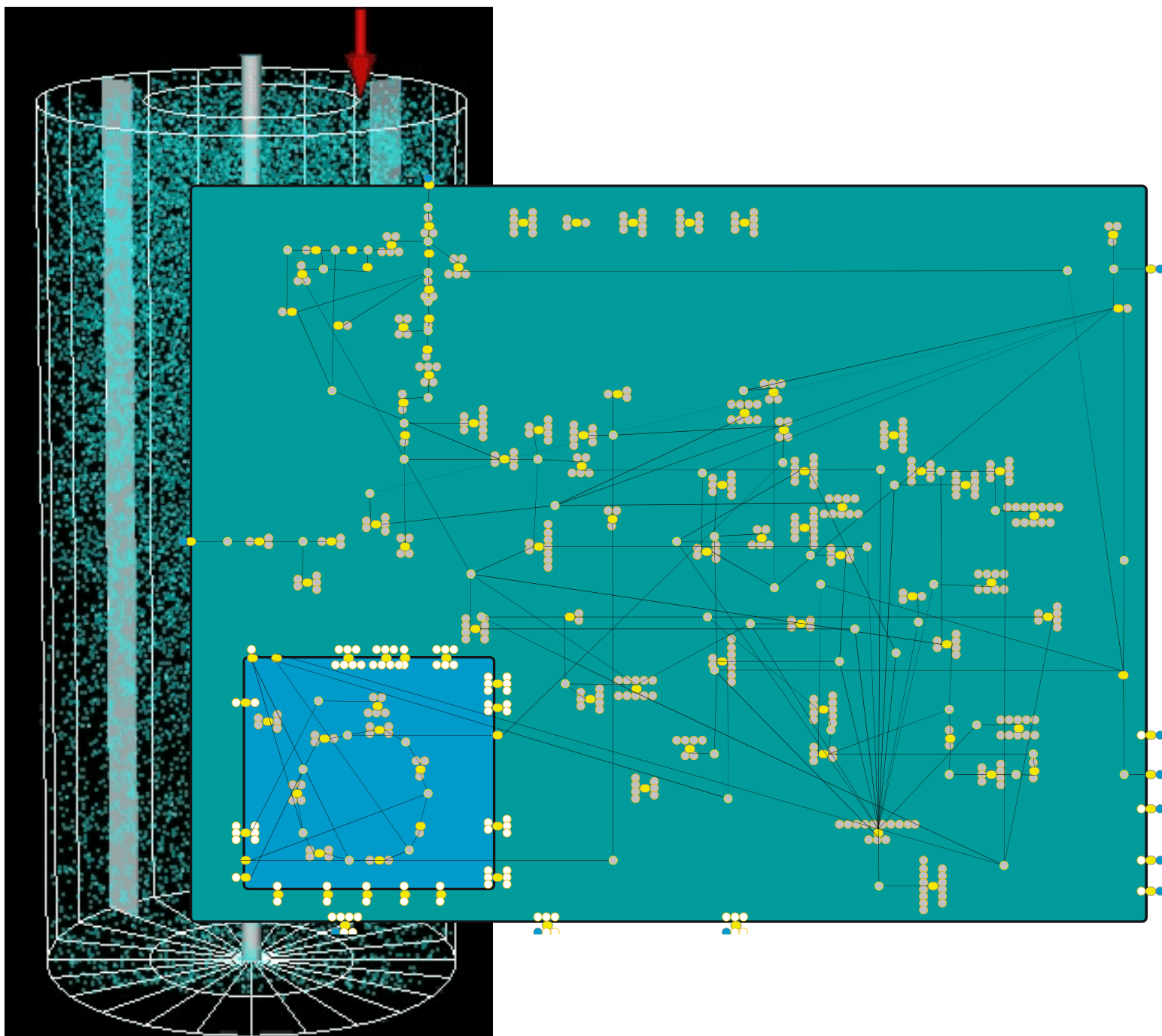






Holon: in some ways a whole,
in other ways a part





CFD

Computational Fluid Dynamics
&

CCD

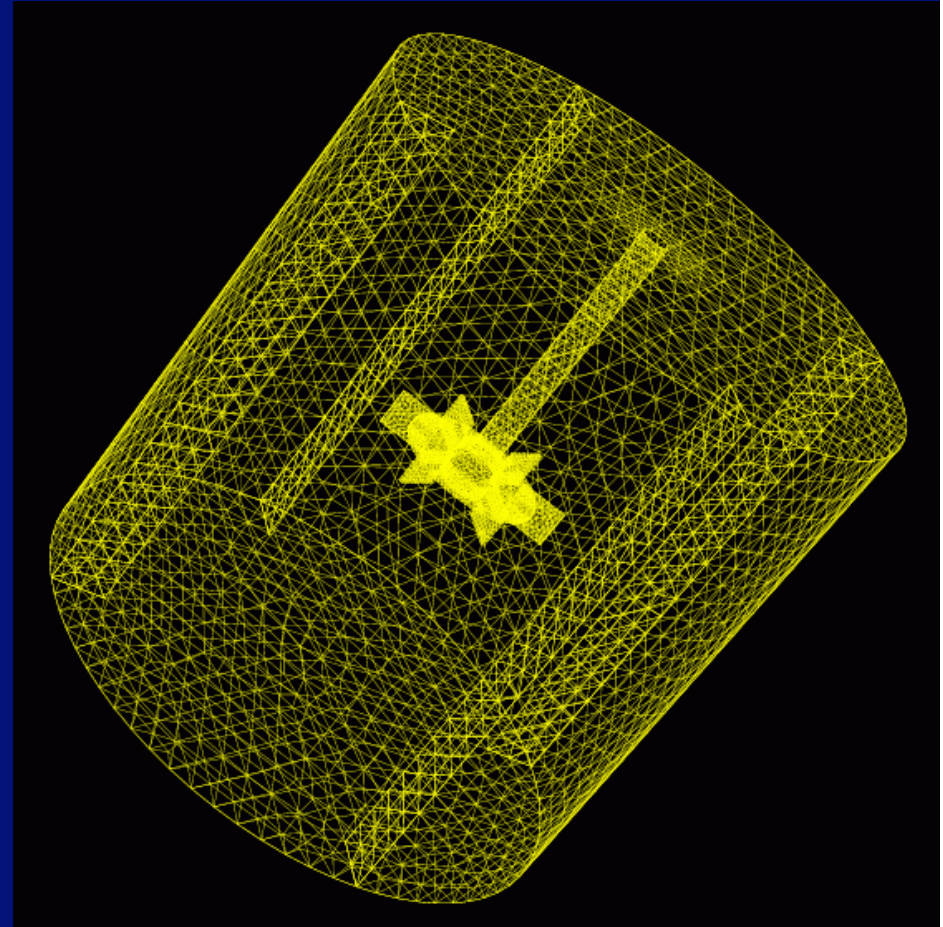
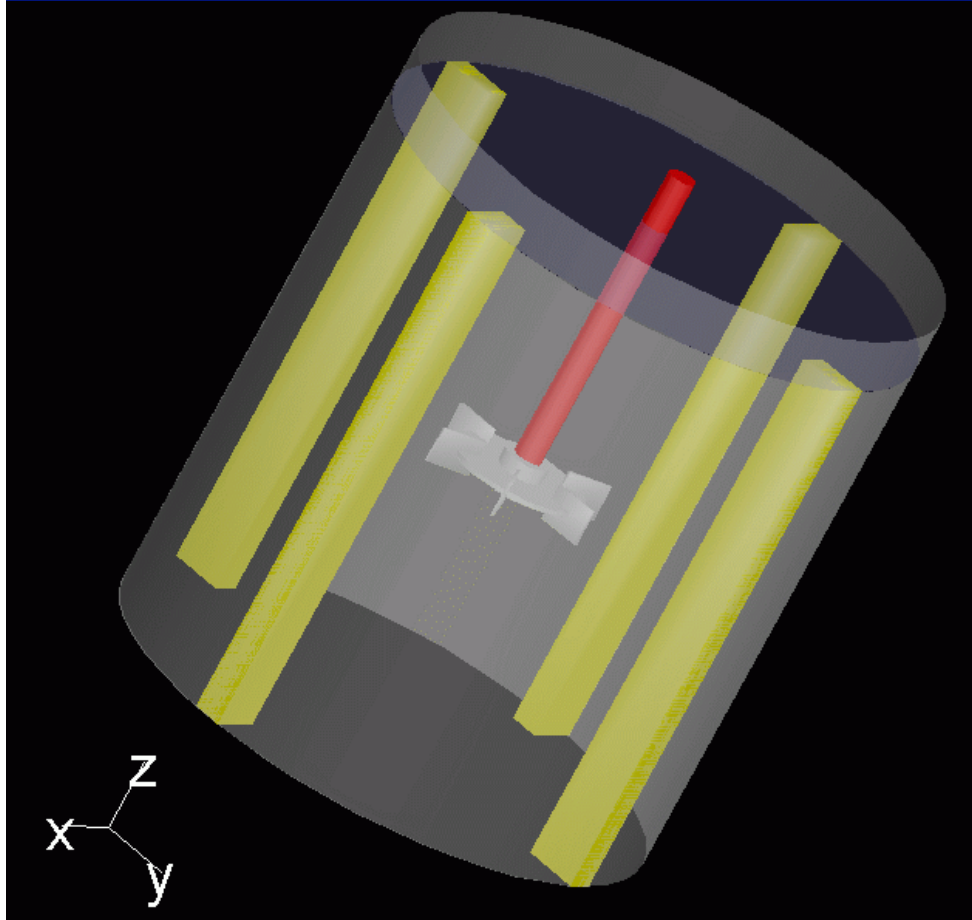
Computational Cellular Dynamics

CFD

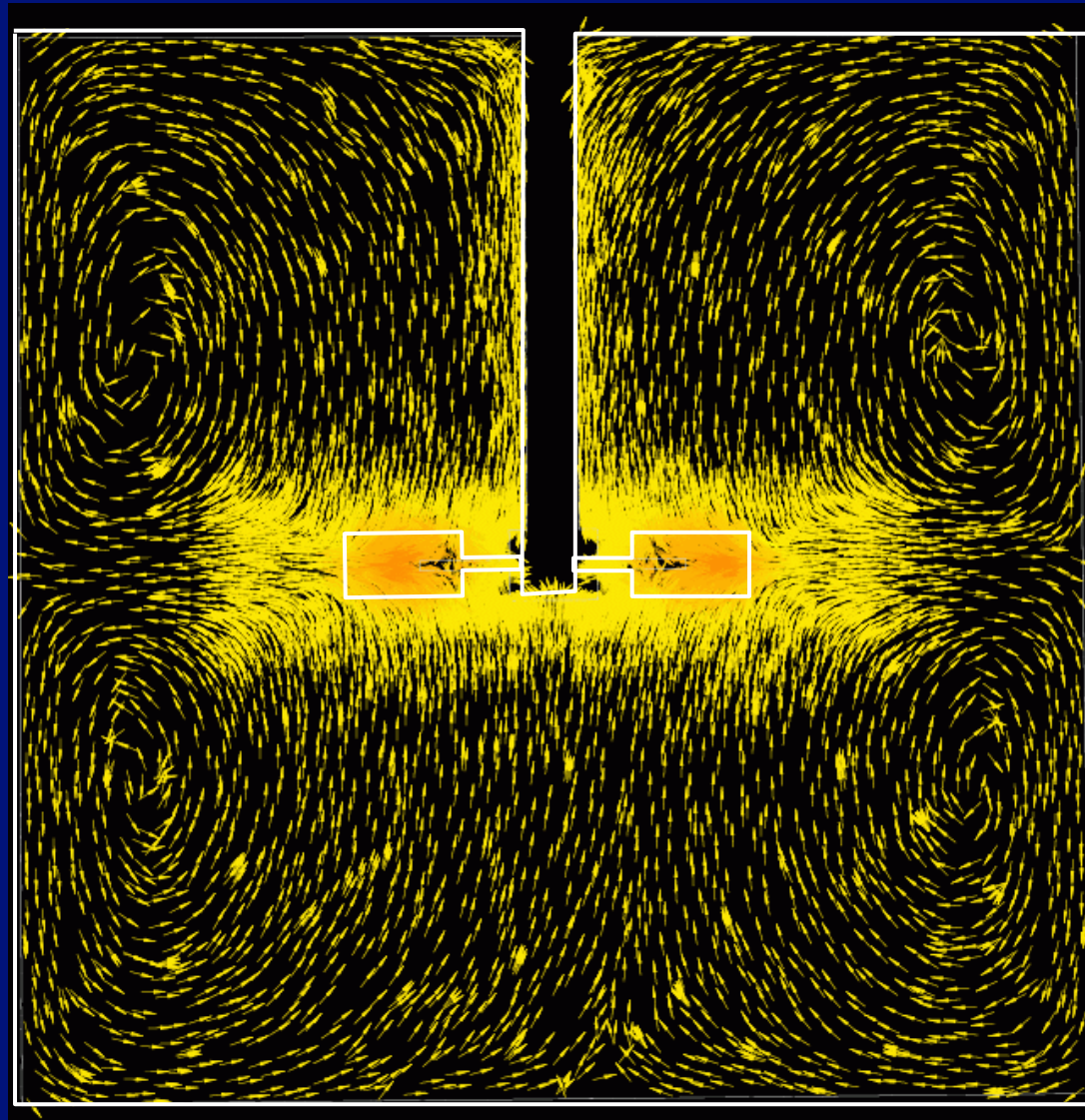
Computational Fluid Dynamics

Modeling of mixing tank

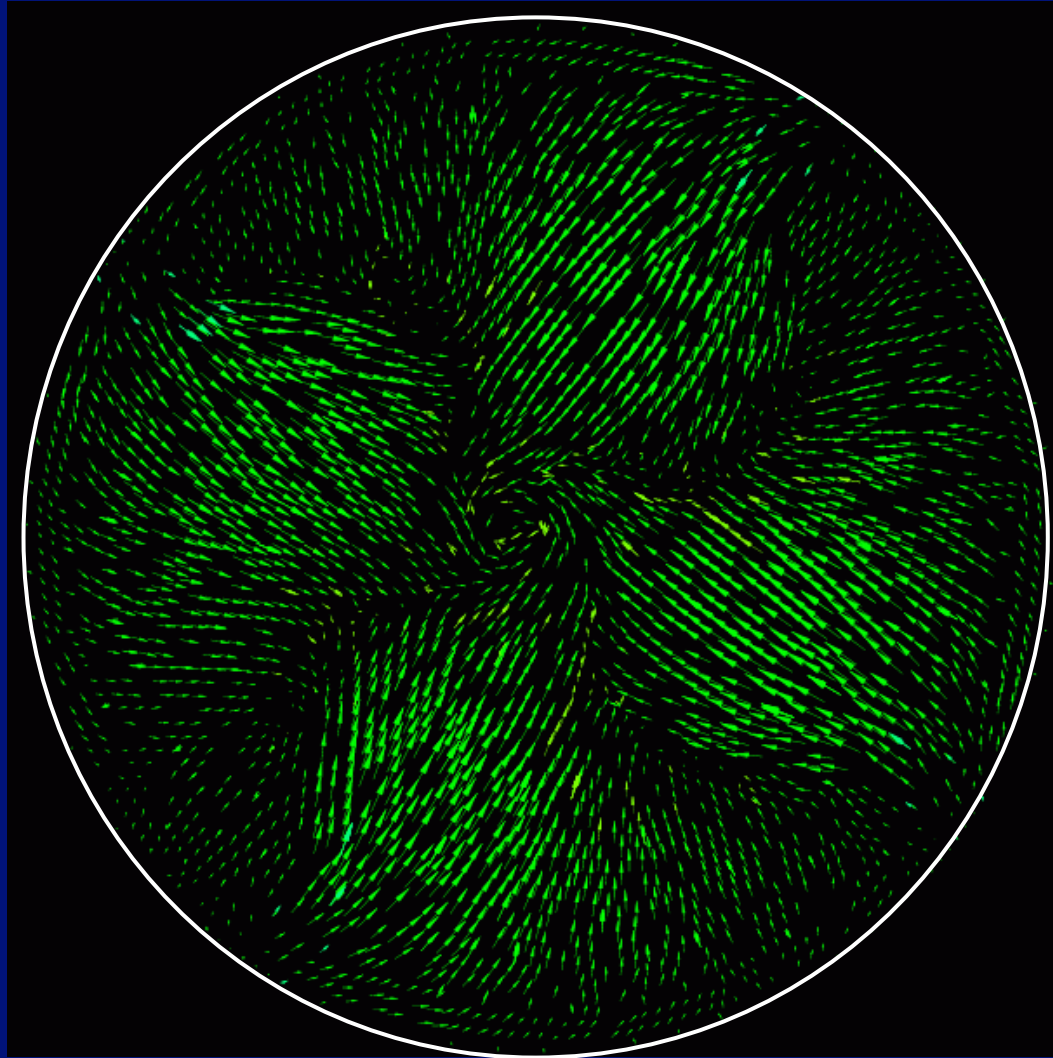
Software package:Fluent



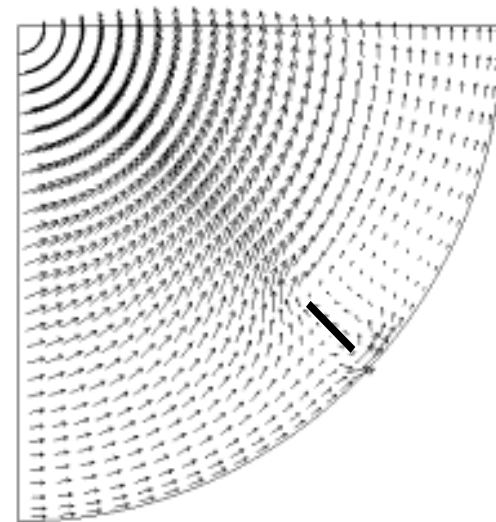
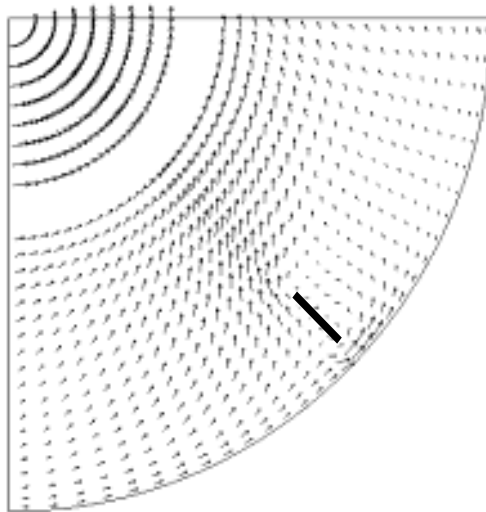
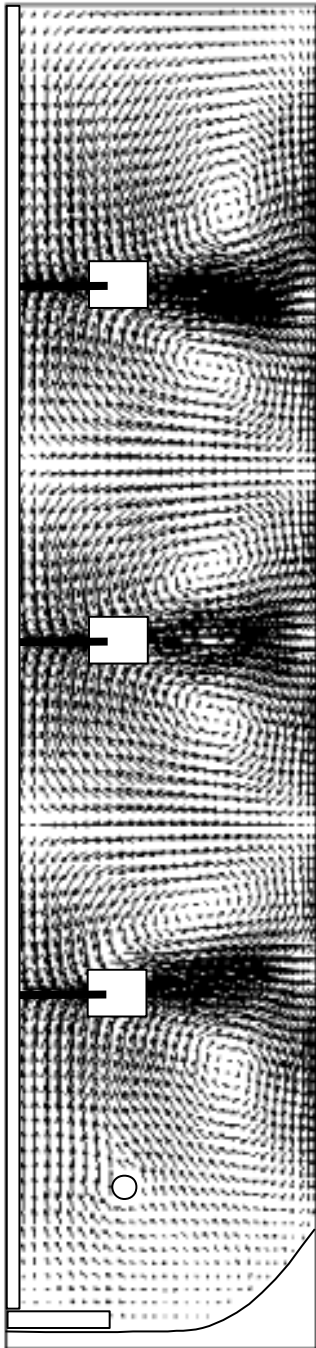
Velocity field (vertical section)



Velocity field (bottom)



Fluid dynamics in a 1500 l reactor with 900 l reaction volume



Simulation of metabolic overflow during fed batch operation

$$\frac{\partial C}{\partial t} + \frac{\partial(u_j C)}{\partial x_j} = \frac{\partial}{\partial x_j} \left(D_{eff} \frac{\partial C}{\partial x_j} \right) + S$$

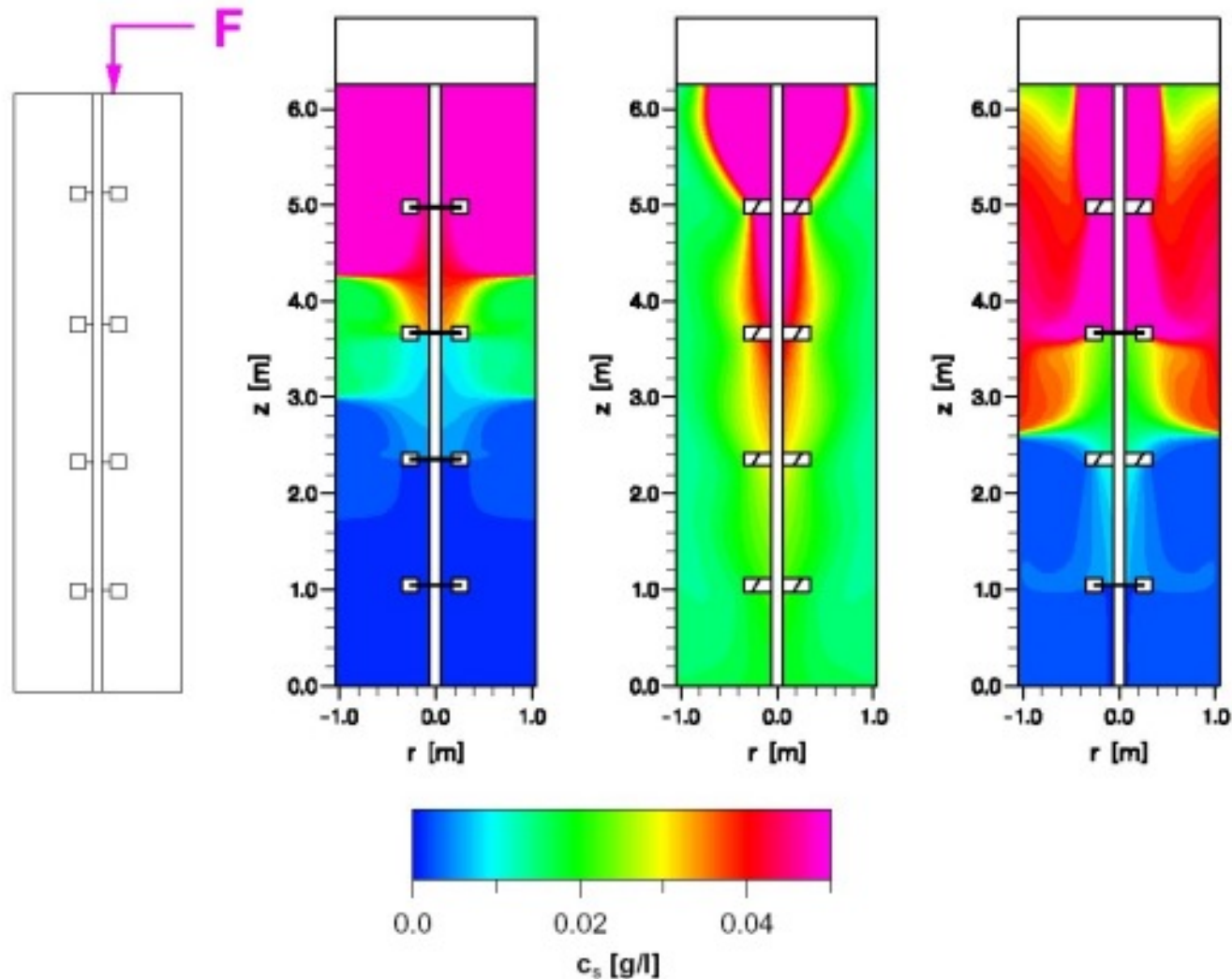
Substrate

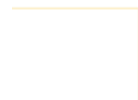
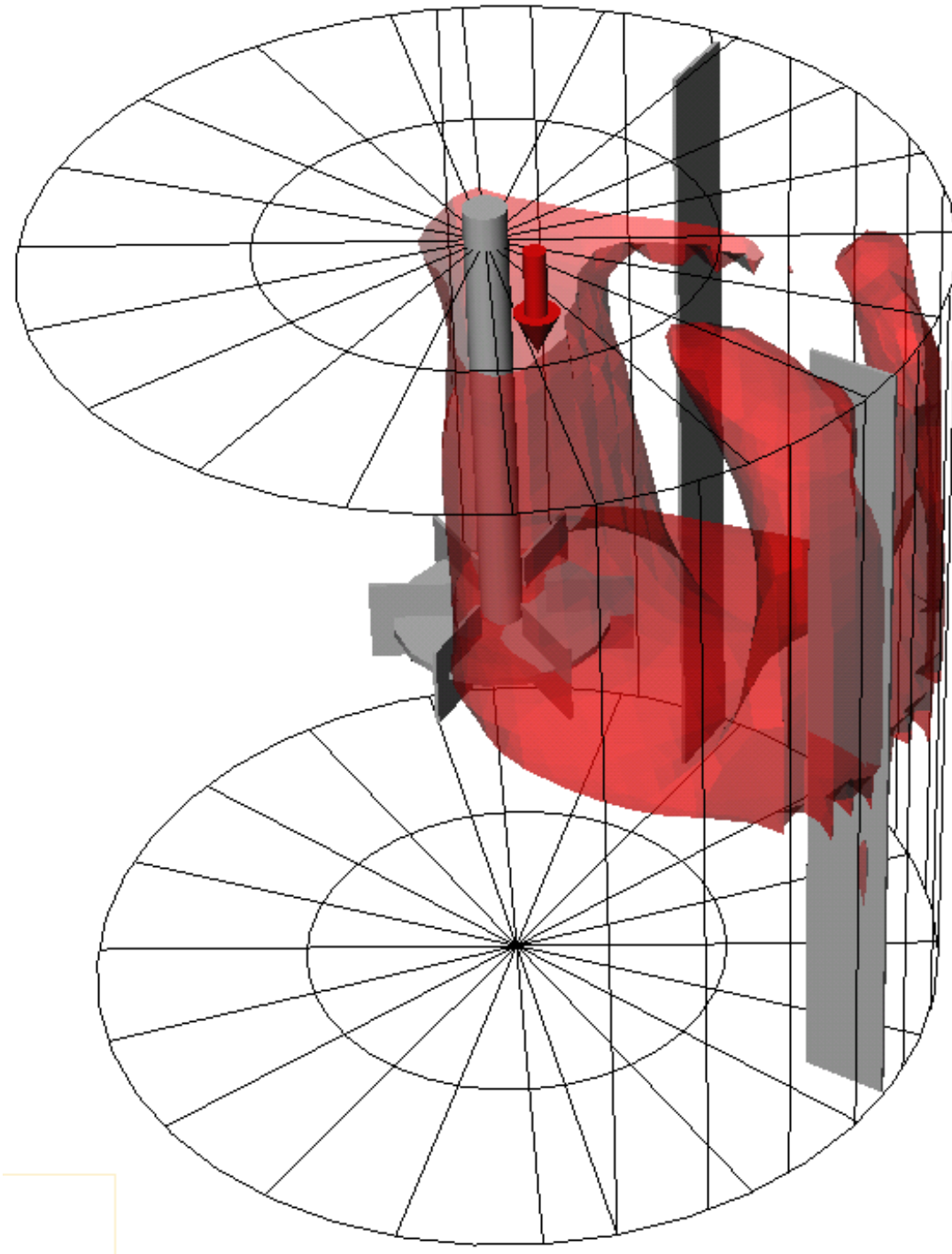
$$S = q_s^{\max} \frac{C_s}{K_s + C_s}$$

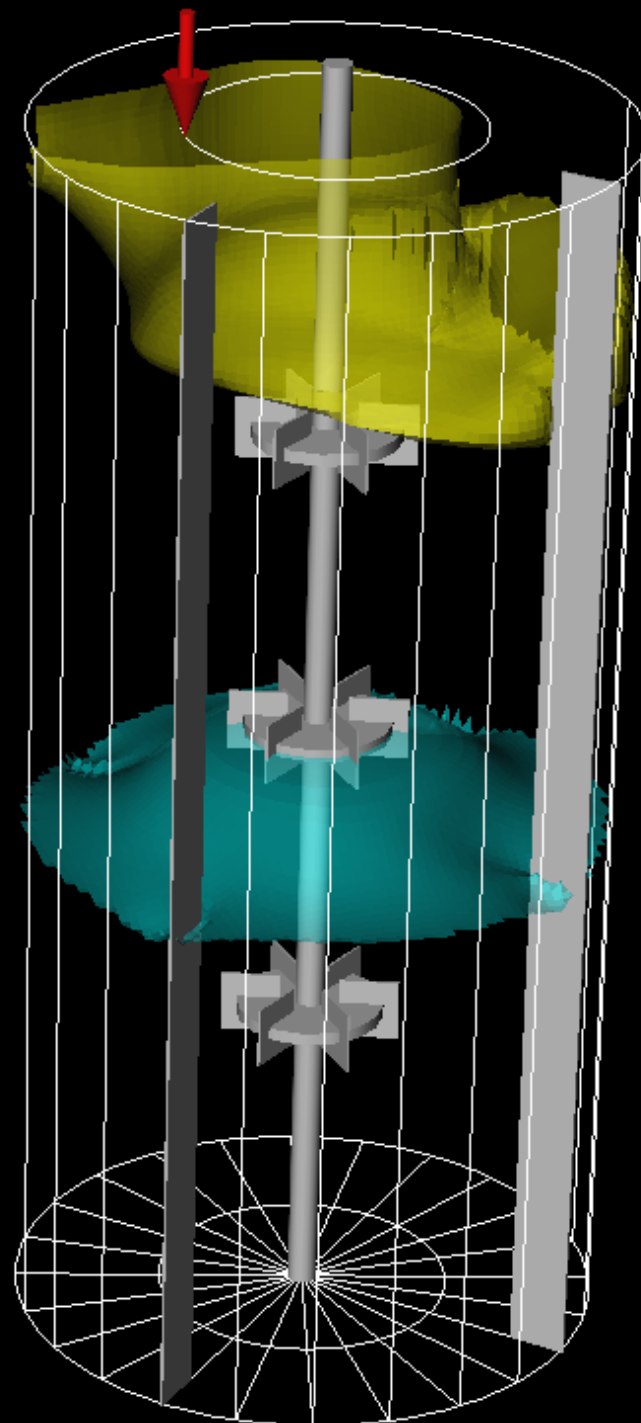
Product

$$S = q_s^{\max} \frac{\max(0, C_s - C_{s,krit})}{K_s + (C_s - C_{s,krit})}$$

Glucose concentration fields during fed-batch cultivation of *Saccharomyces cerevisiae*







Limitations

- **Analysis of systems behavior if biophase needs to be segregated into individual cells -> local interactions between microorganisms and their environment = $f(\text{intracellular state (history of the individuum)})$**

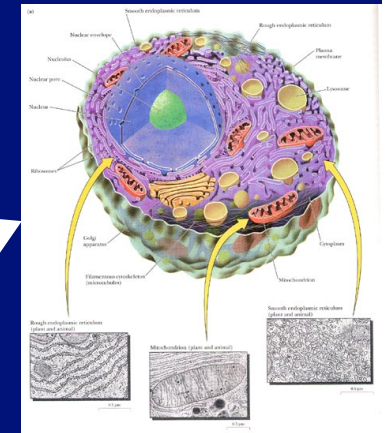
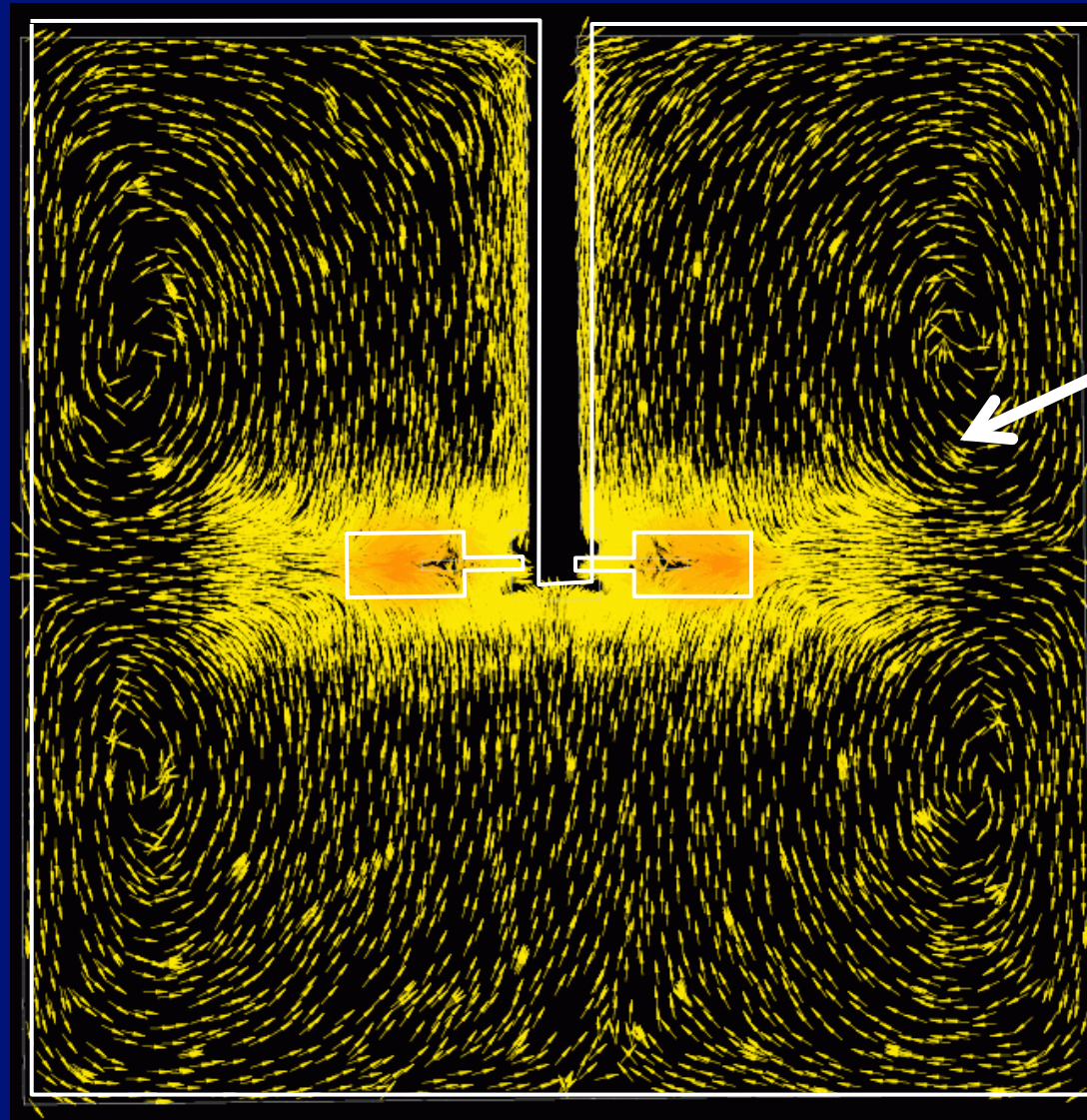


Lifelines of single cells in space and time



Dynamic behavior of the population

Velocity field (vertical section)



Lagrange-Euler-Approach

Fluid dynamics

- Predictions of *convective* movement of a single particle from simulated velocity field:

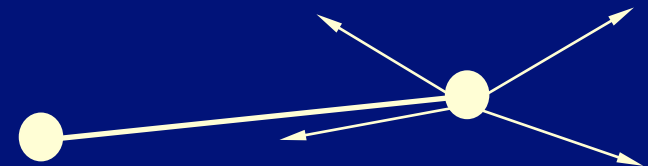
$$\frac{d\vec{x}}{dt} = \vec{V}$$

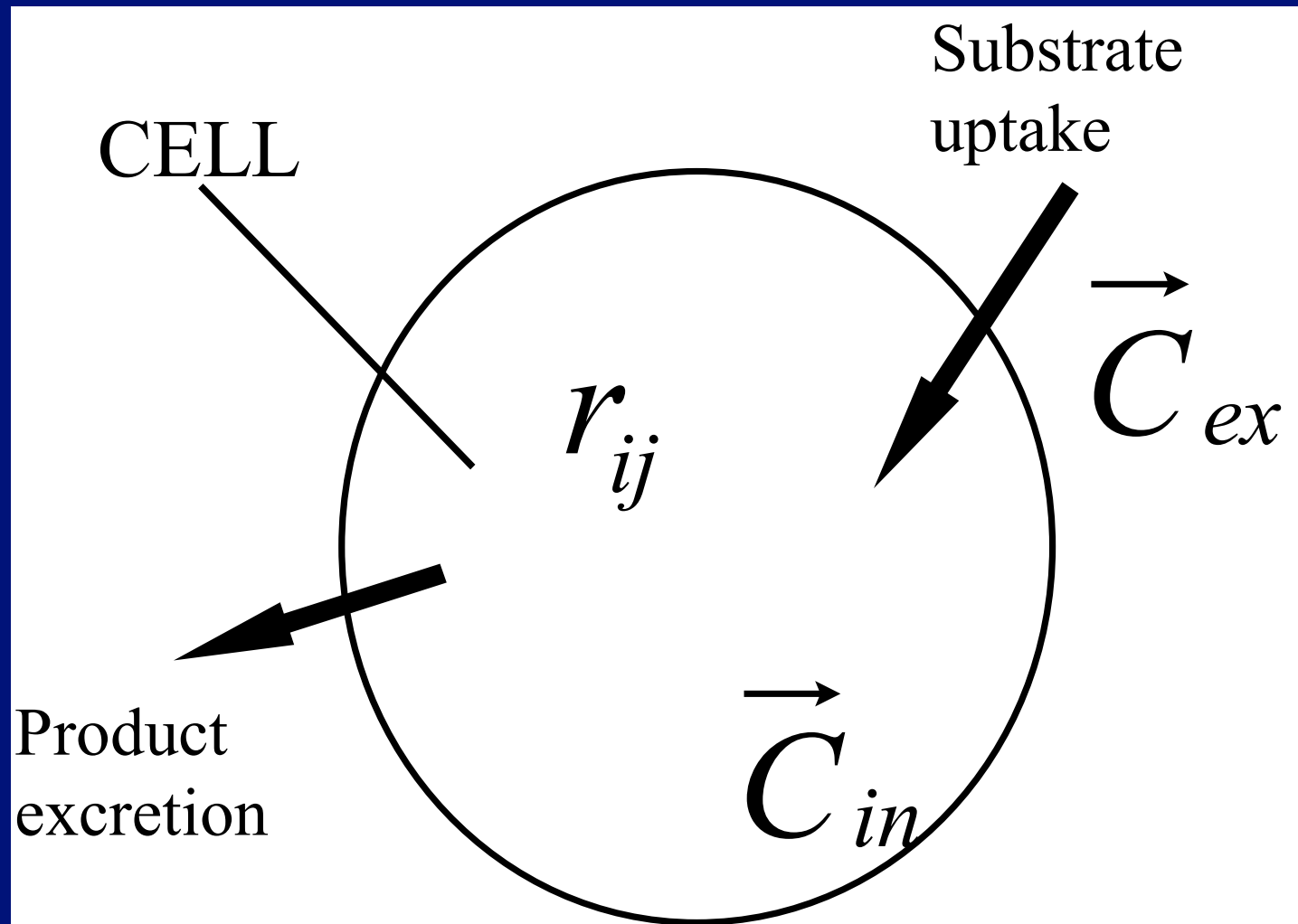

A diagram illustrating convective movement. It shows two yellow dots representing a particle at two different times. The first dot is labeled 't' and the second dot is labeled 't + Δt'. A straight arrow points from the first dot to the second dot, indicating the direction and distance of movement over the time interval Δt.

- Superposition of random movement caused by turbulent Dispersion (SDE):

$$\vec{x}(t + dt) = \vec{V}dt + (2D_Tdt)^{1/2} \vec{\xi}$$

ξ - Random number





- **Computation of intracellular state along the trajectory from integration of intracellular mass balance**

equations:

$$\frac{d\vec{c}_{in}}{dt} = \sum_n r_{ij} [\vec{c}_{in}(t), \vec{c}_{ex}(\vec{x}, t)]$$

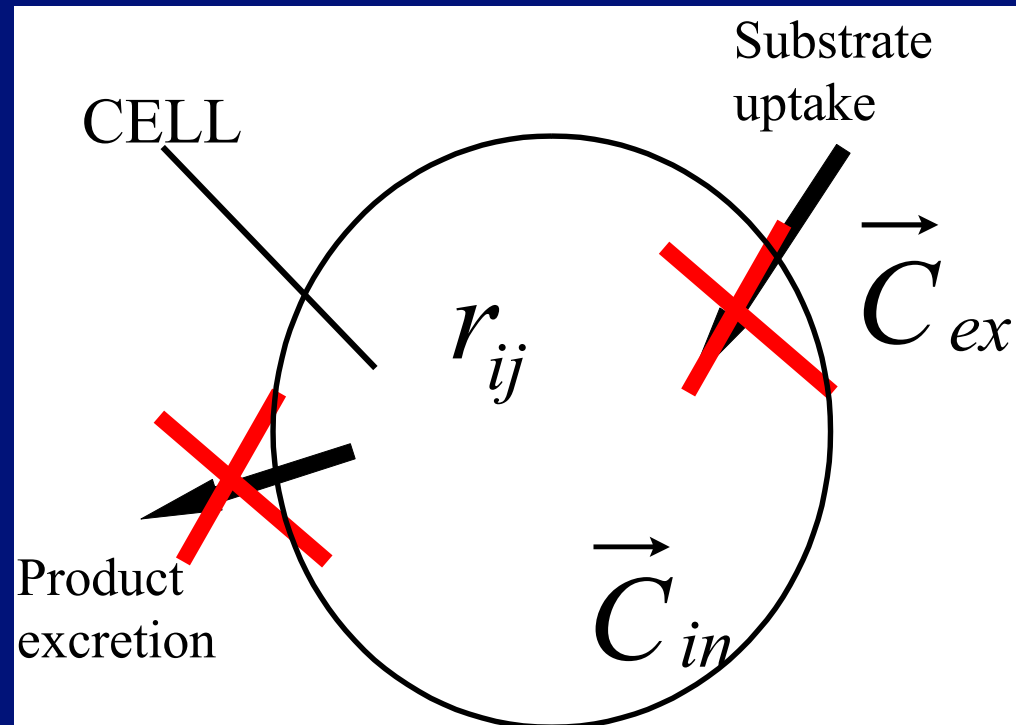
(substrate uptake and product excretion rates are included in r_{ij})

- **The individual exchange rates with the environment are taken into account within the Eulerian simulation of the extracellular state:**

$$\frac{\partial \vec{c}_{ex}}{\partial t} + (\vec{V} \nabla) \vec{c}_{ex} = \nabla (D_T \nabla \vec{c}_{ex}) + S_{exchange}(\vec{c}_{ex}, \vec{c}_{in})$$

($\vec{V}$ and D_T again from CFD-simulations)

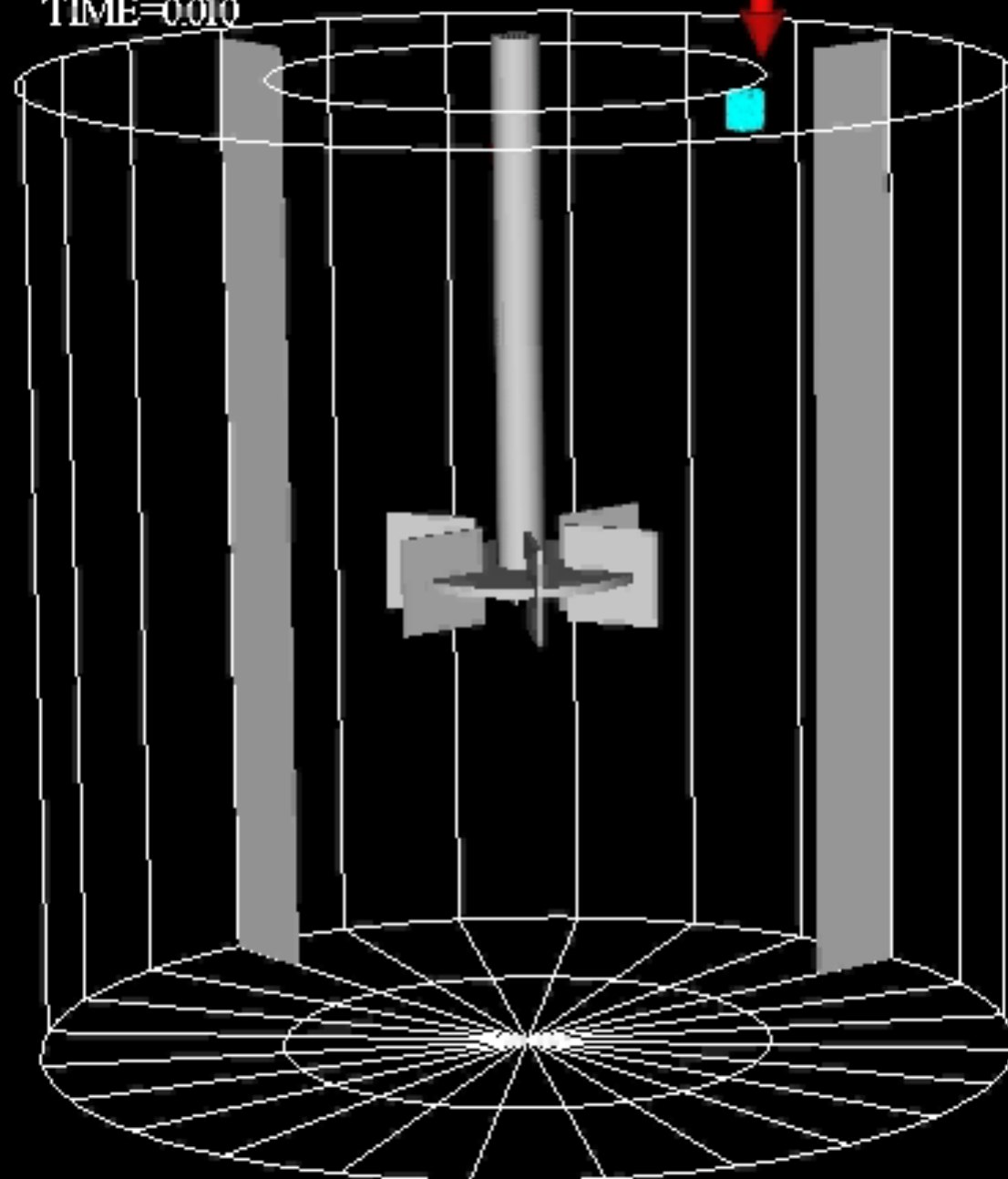
WITHOUT EXCHANGE (TRACER PARTICLE)



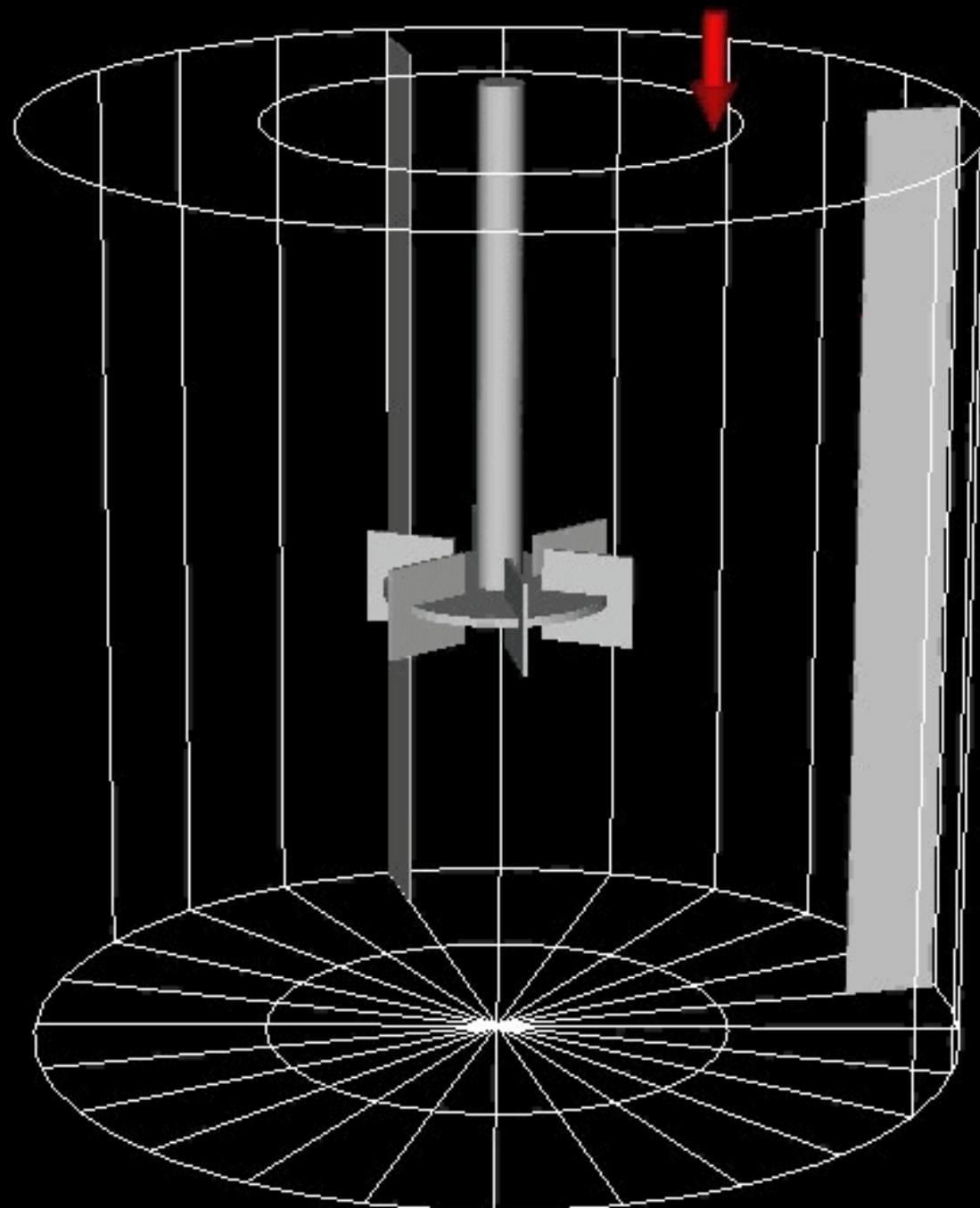
SIMULATION OF TRACER EXPERIMENTS

MIXING CHARACTERISTICS

TIME=0.010

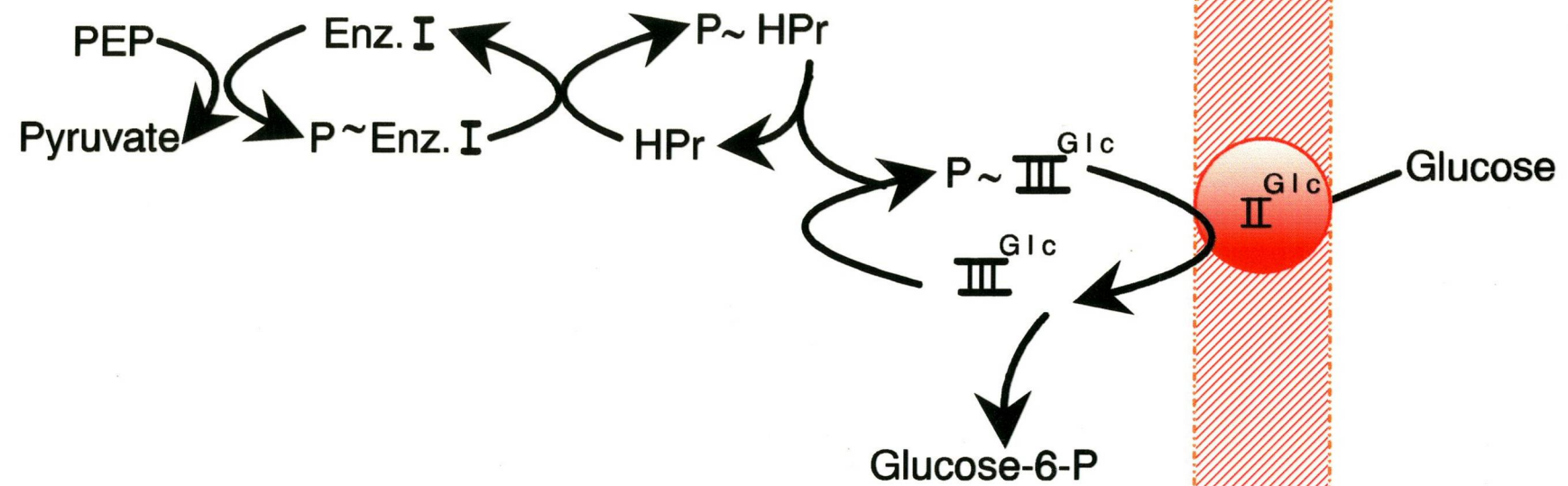


TIME=0.000

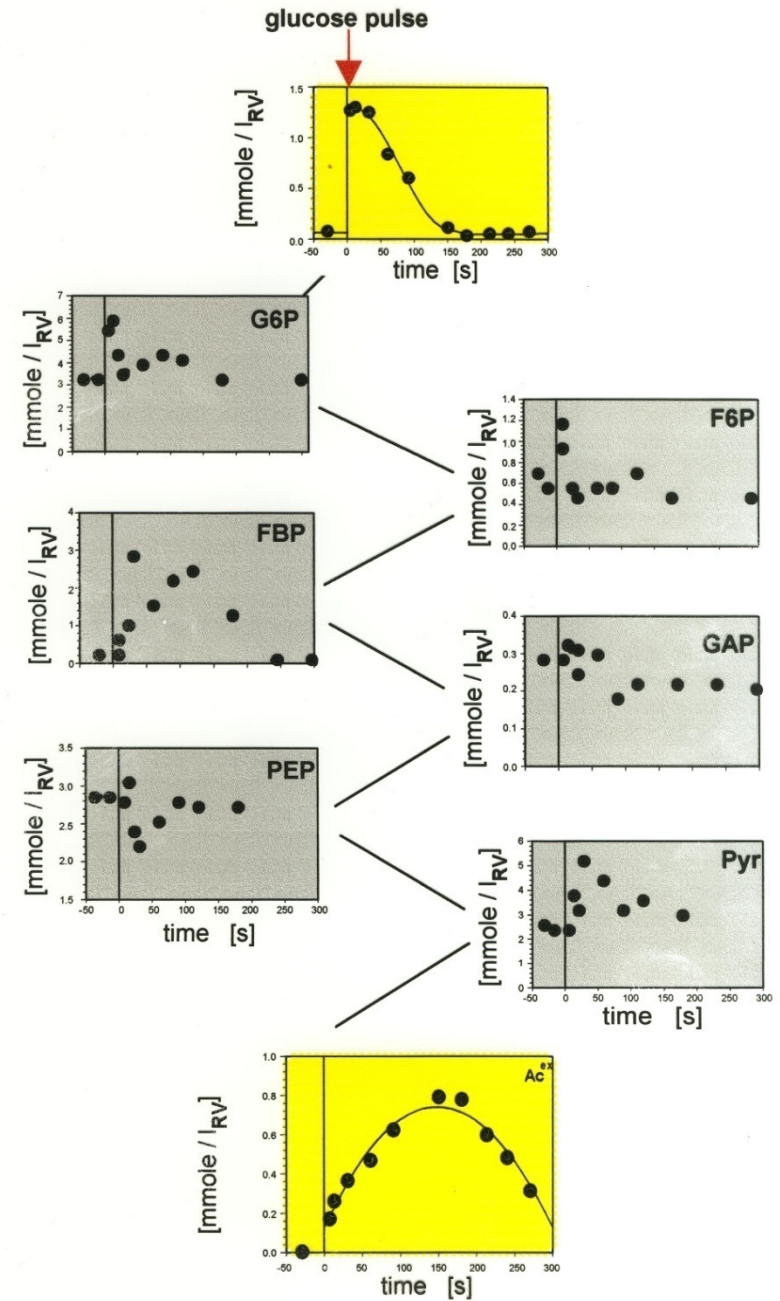


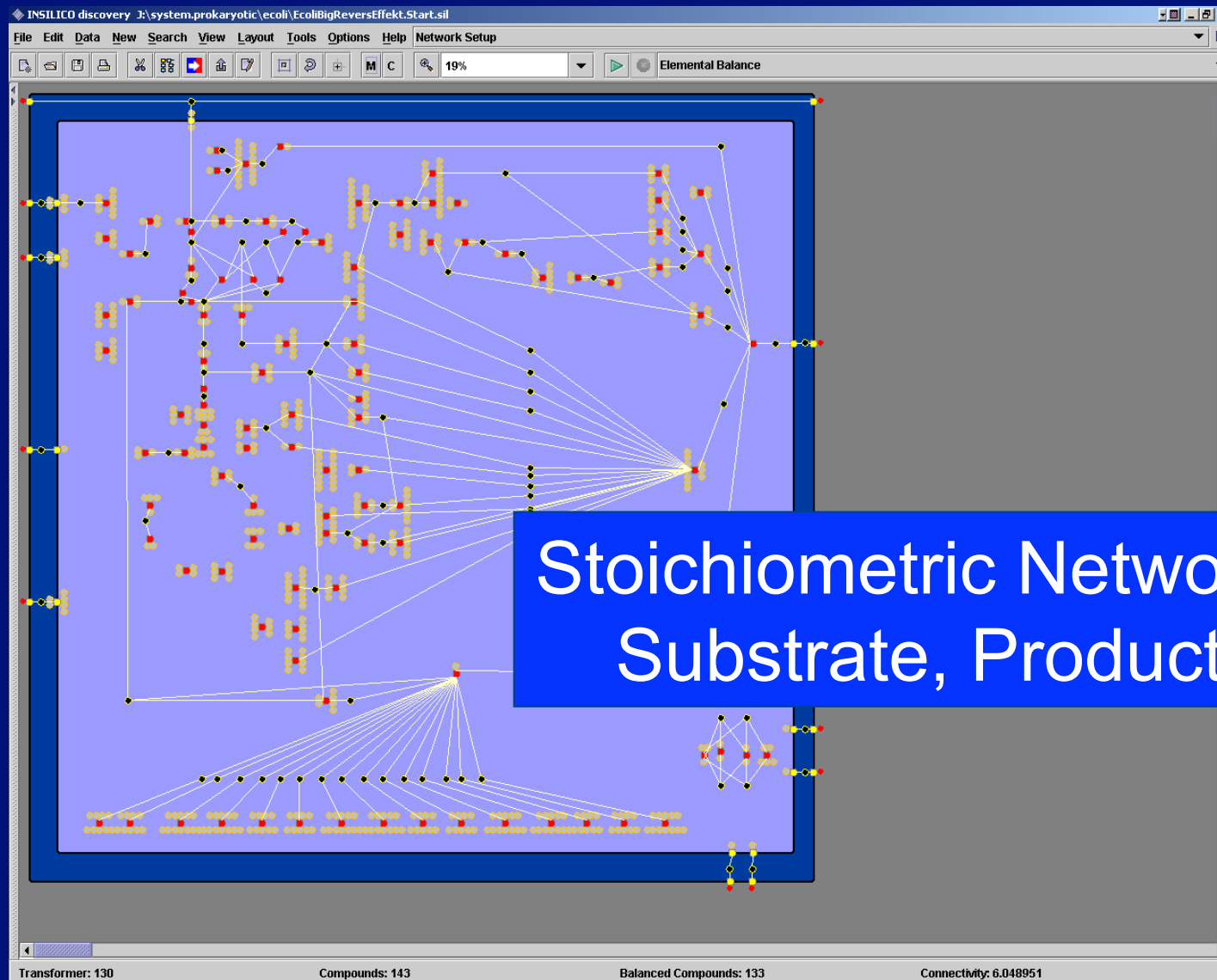
Glucose uptake into *Escherichia coli*:

Traveling along the lifelines of single
cells

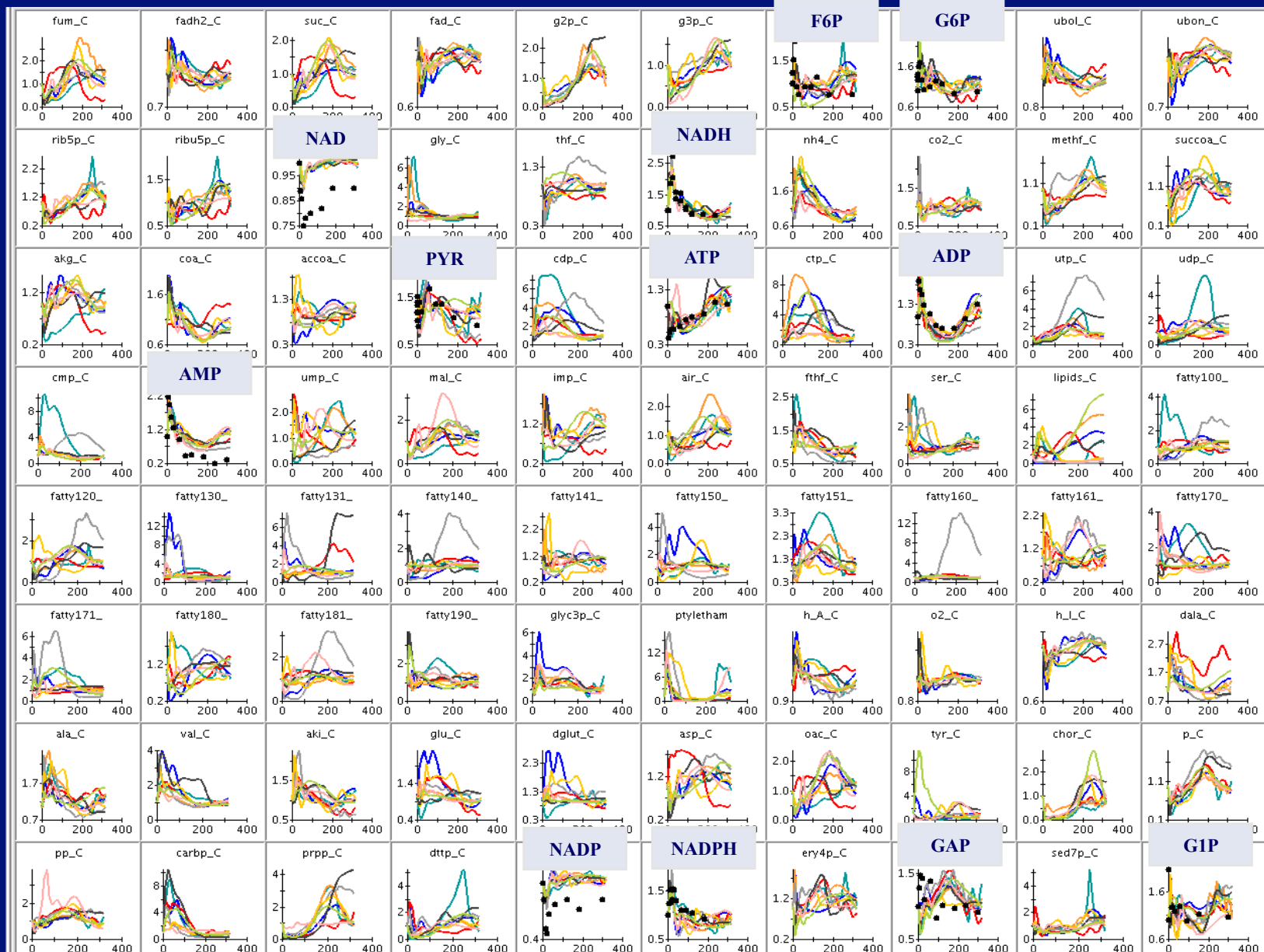


STIMULUS-RESPONSE METHODOLOGY



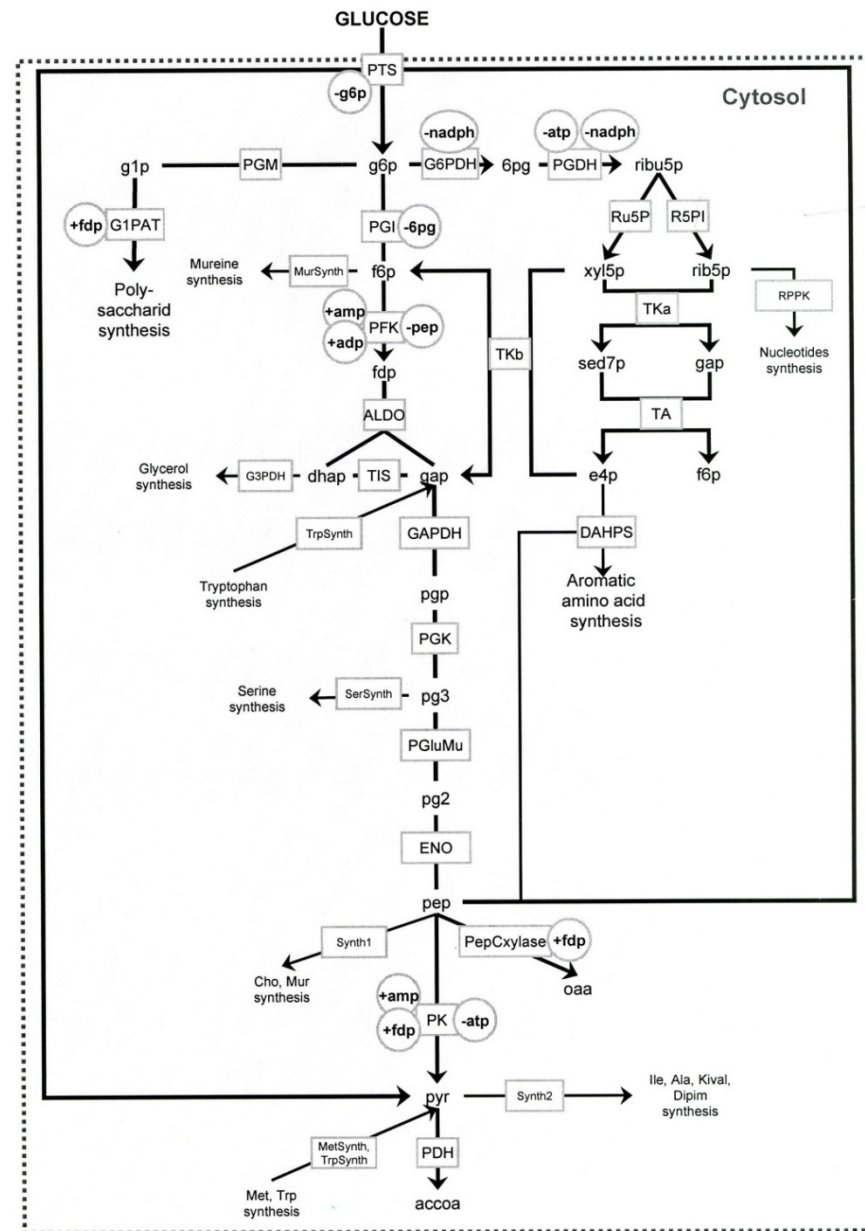


Escherichia coli



Escherichia coli

Reuss, Aguielera-Vazquez, Mauch (2007). Topics Current Genetics



Chassagnole, Noissomit-Rizzi, Schmid, Mauch, Reuss (2002) Biotechnol. Bioeng.79, 53-73

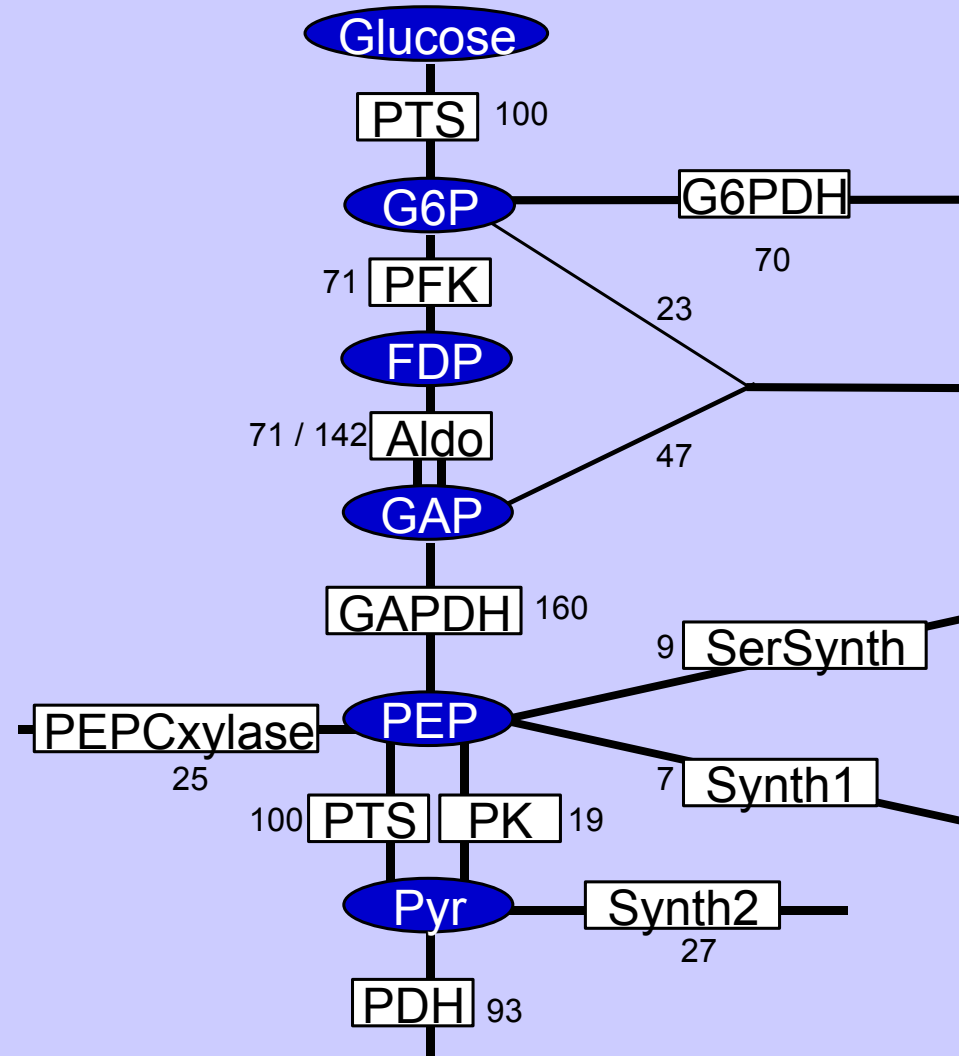
Bridging scales

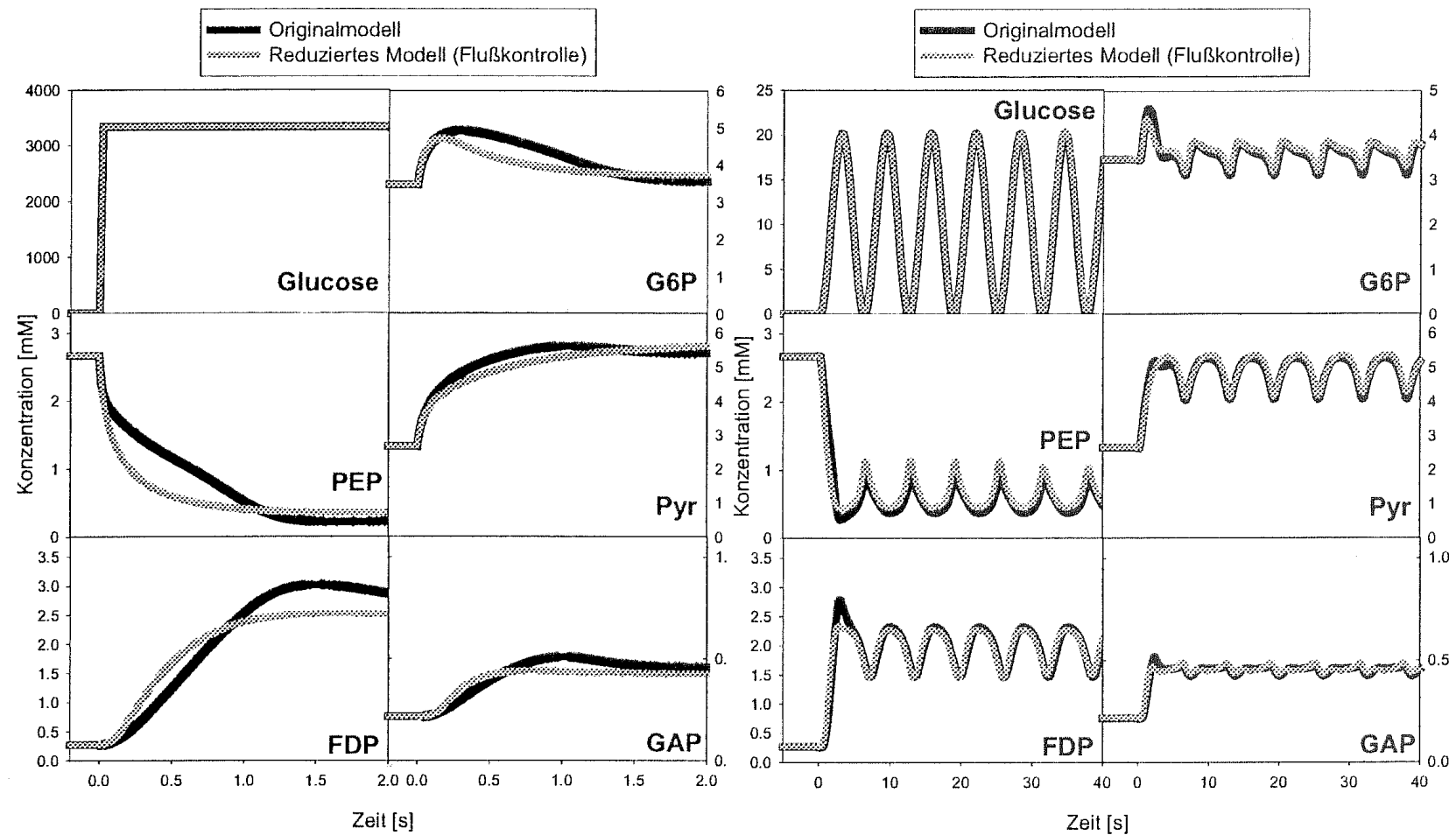
Model reduction

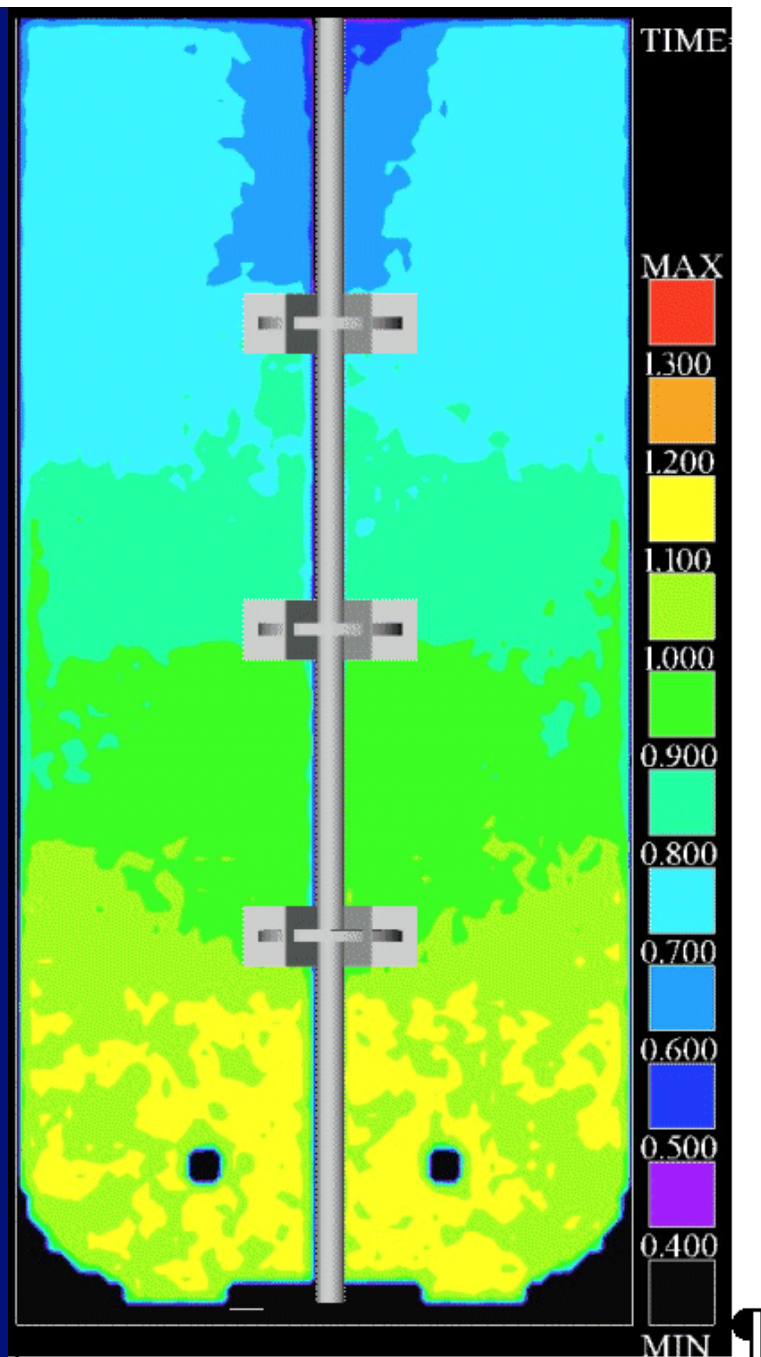
Flux control coefficients

$$C^J = \frac{E}{J} \frac{dJ}{dE} = \frac{d \ln J}{d \ln E}$$

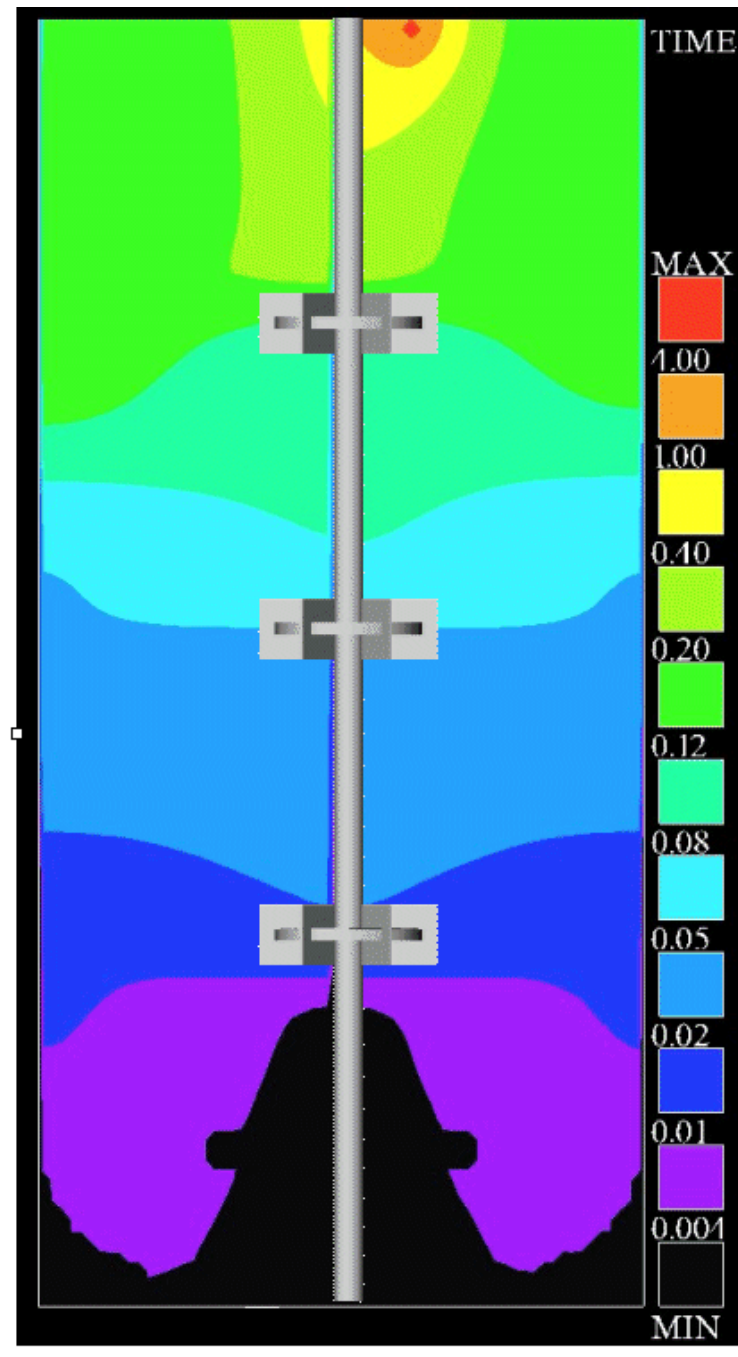
Glucose uptake system (*E.coli*: PTS)



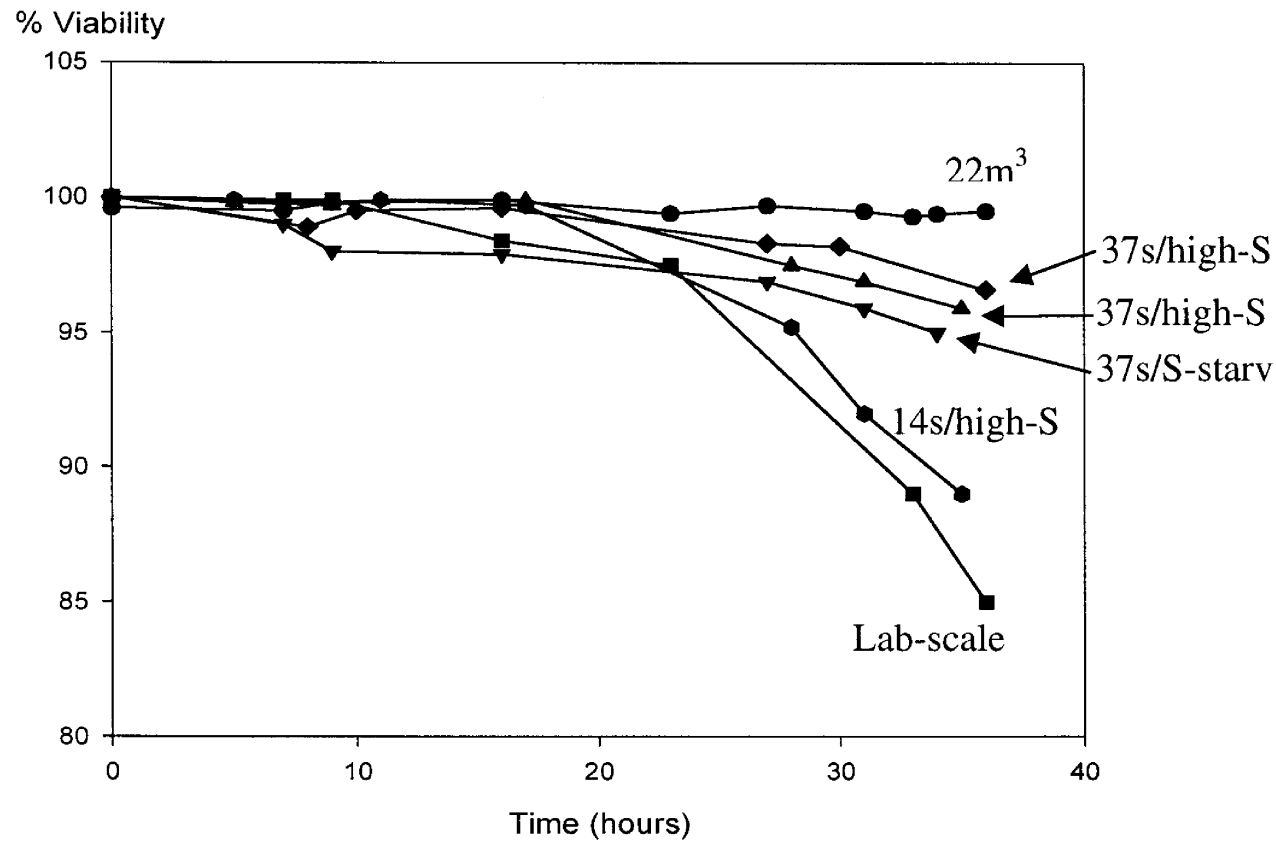




Pep/Pyr

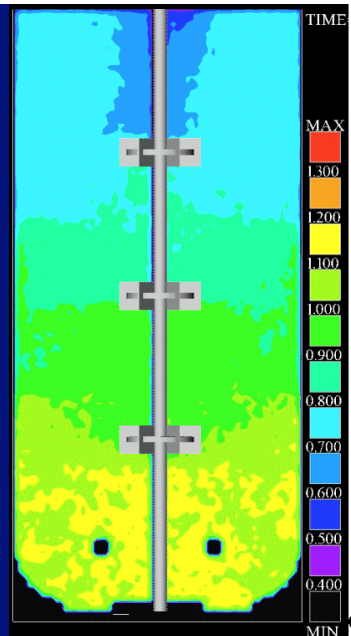


Glucose (Segregated)

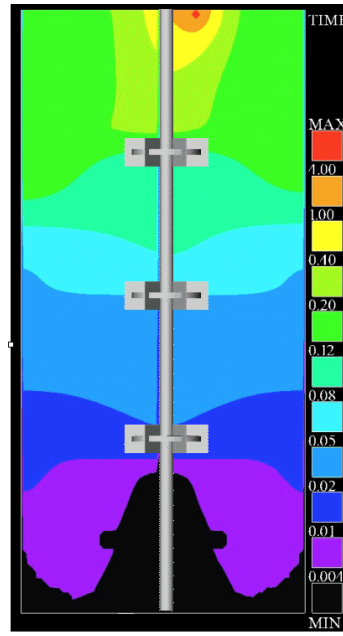


E. coli viability (flow cytometry) in lab-scale, at 22 m³-scale and in a scale-down reactor with 14-37 sec passages through a glucose starvation or glucose excess (oxygen limitation) zone

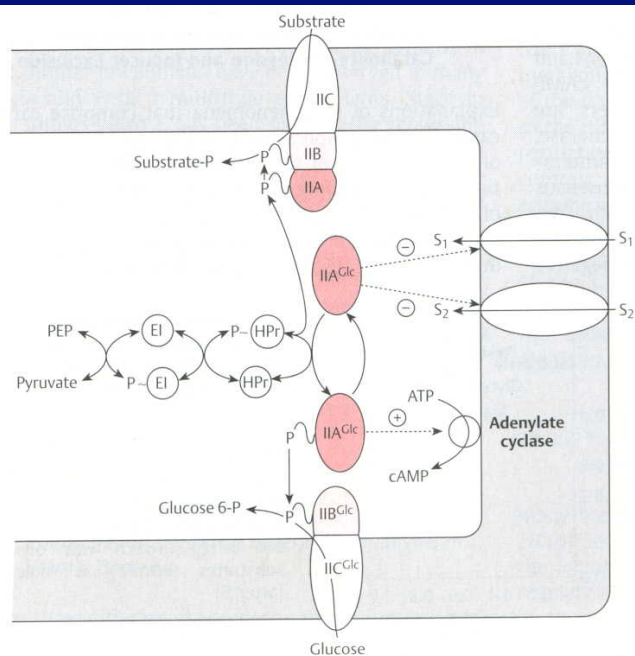
Hewitt et al. 1999



Pep/Pyr

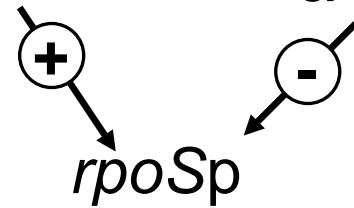


Glucose (Segregated)



ppGpp

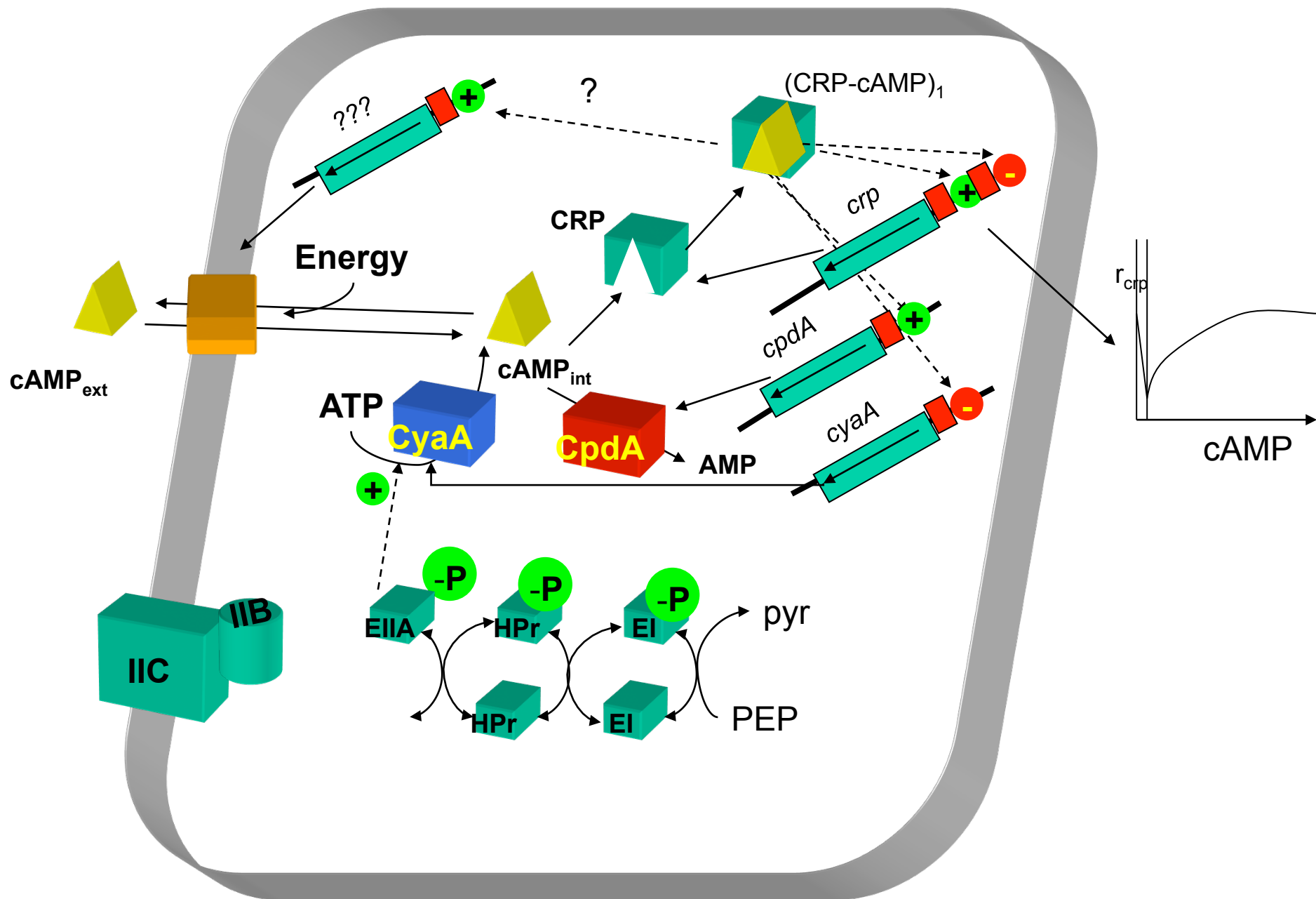
cAMP-CRP



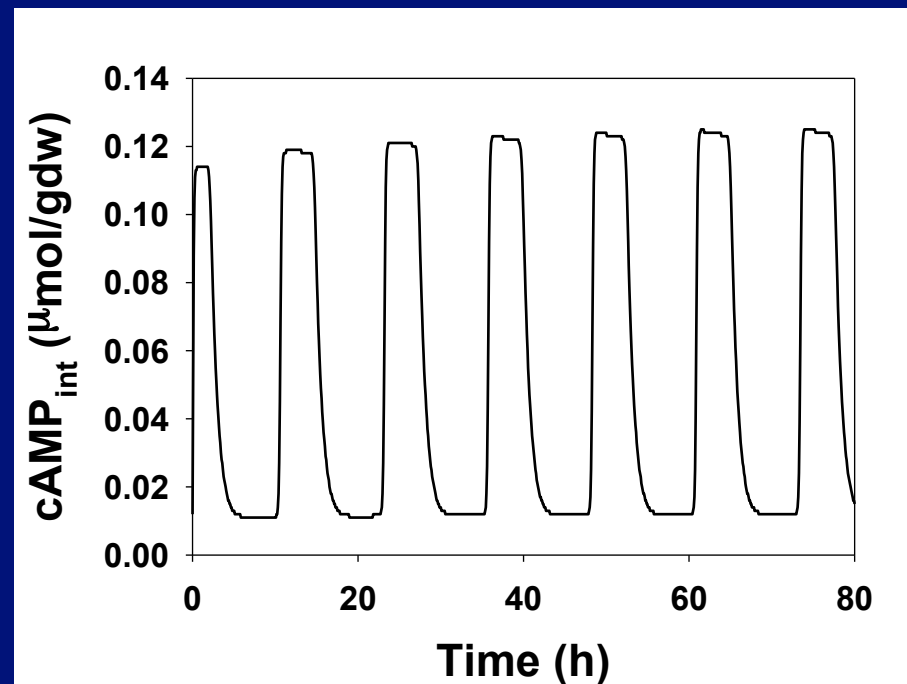
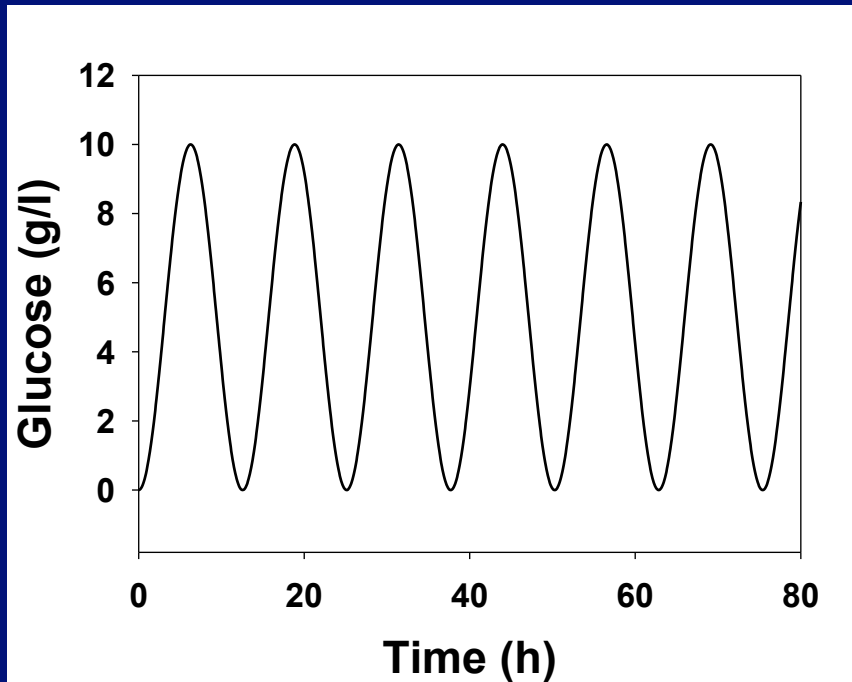
rpoSp

rpoS

σ^S

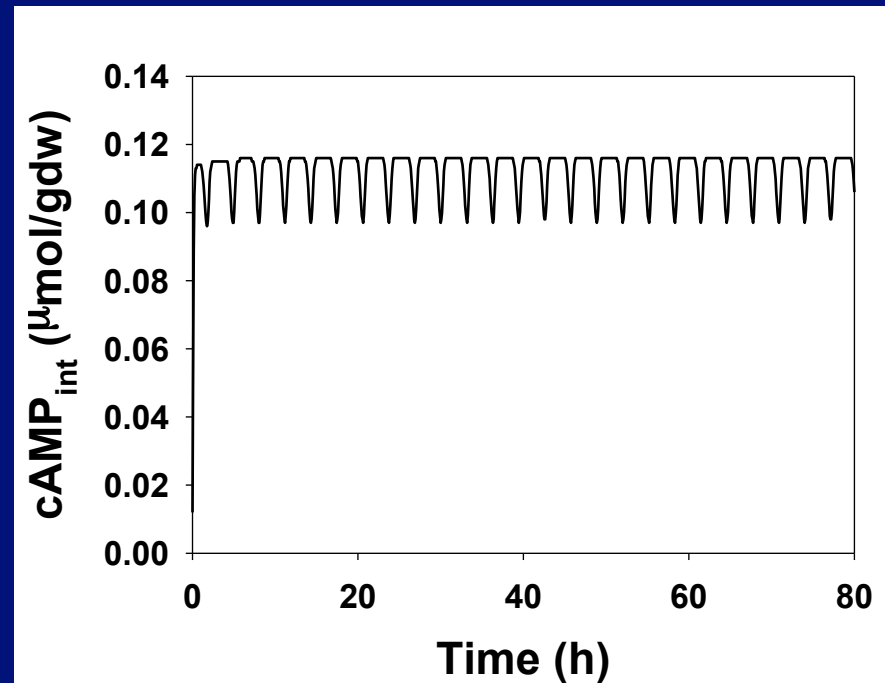
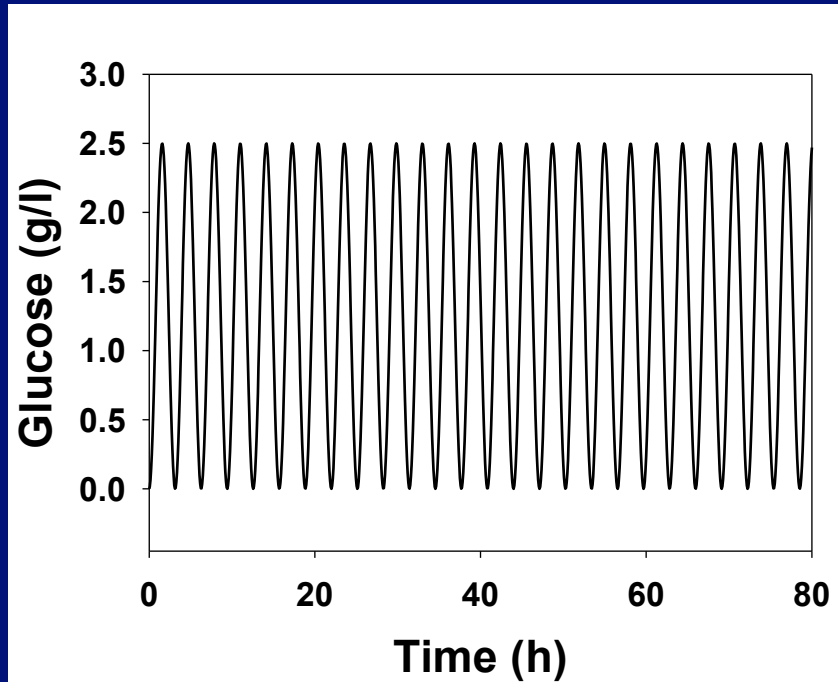


Low frequency



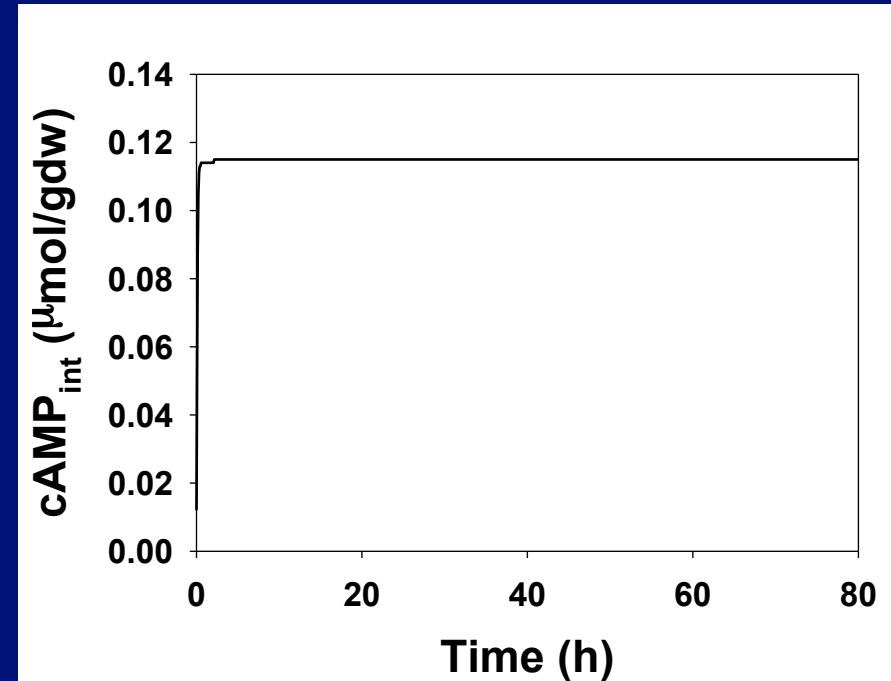
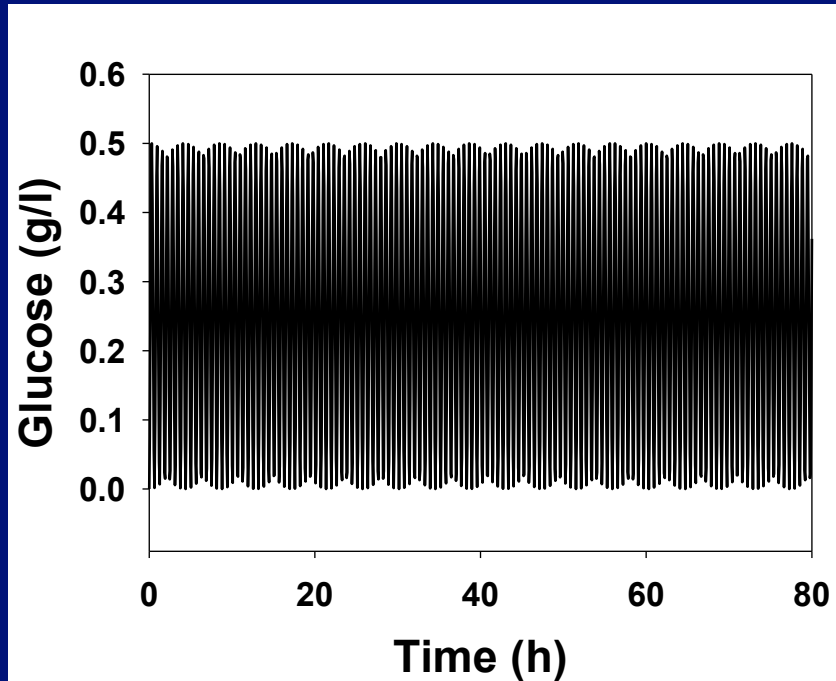
Sevilla, Reuss: to be published

Medium frequency



Sevilla, Reuss: to be published

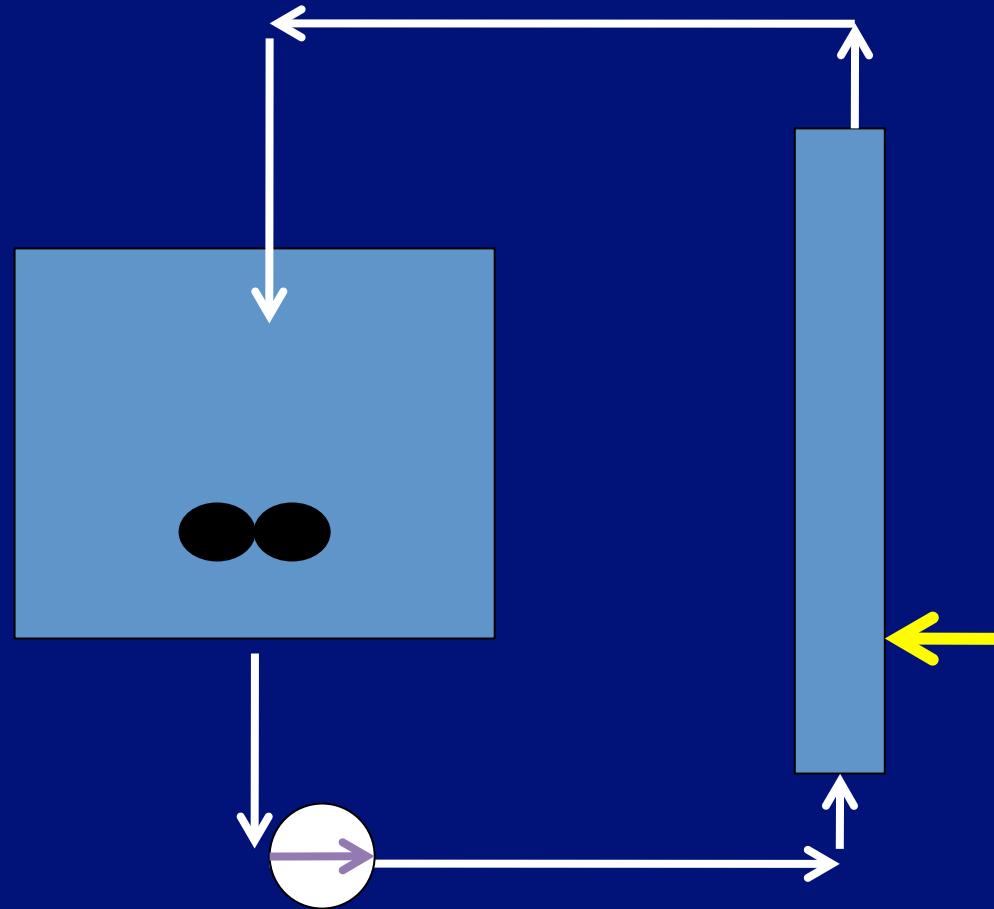
High frequency



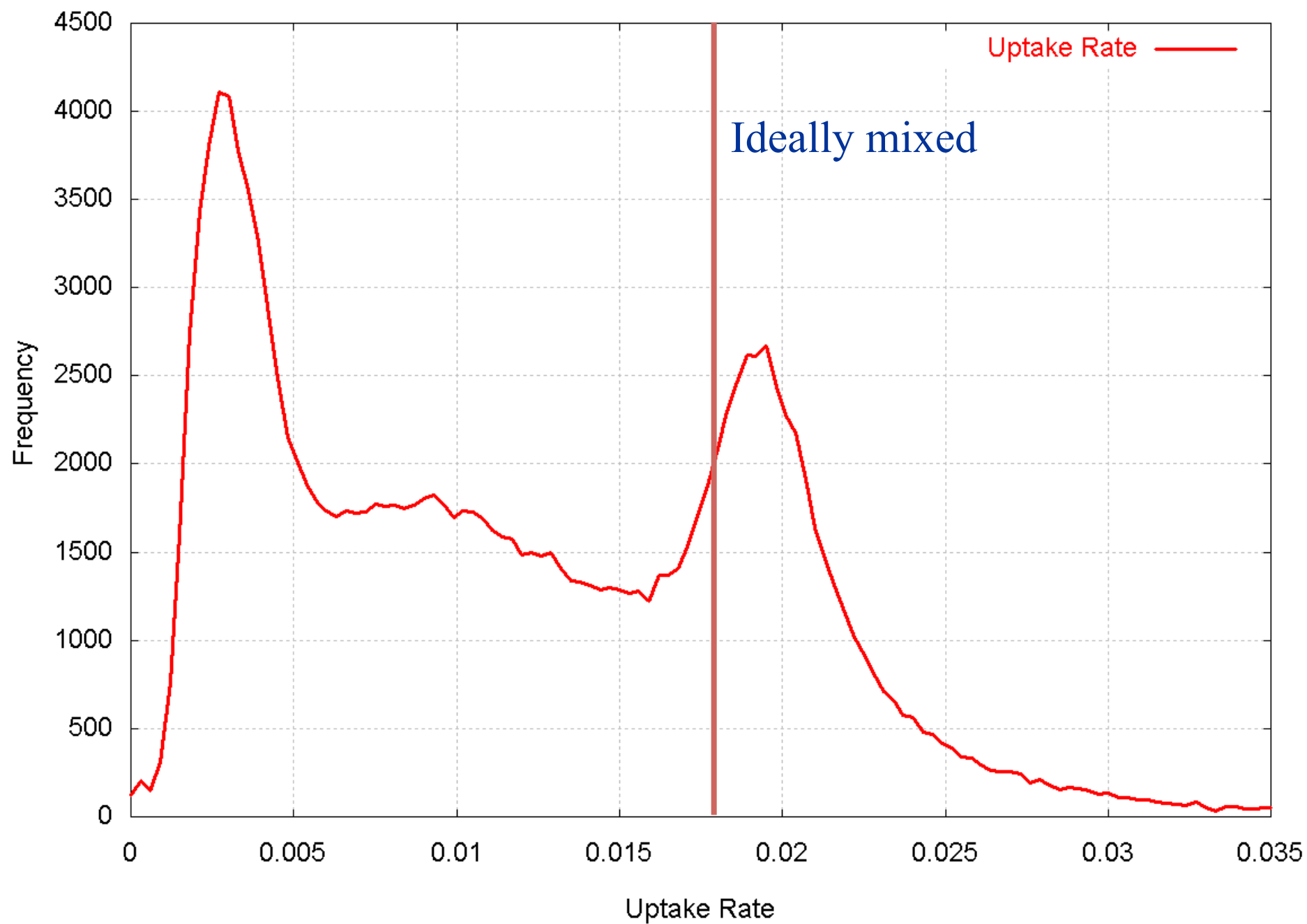
Sevilla, Reuss: to be published

Delvigne, F., Boxus, M., Ingels, S., Thonart, P (2009)
Microb. Cell Factories 8: 1-17 prpoS:GFP reporter gene

Delvigne, F., Boxus, M., Ingels, S., Thonart, P (2009)
Microb. Cell Factories 8: 1-17 *prpoS*:GFP reporter gene



expression of *rpos* is reduced and distribution is bimodal



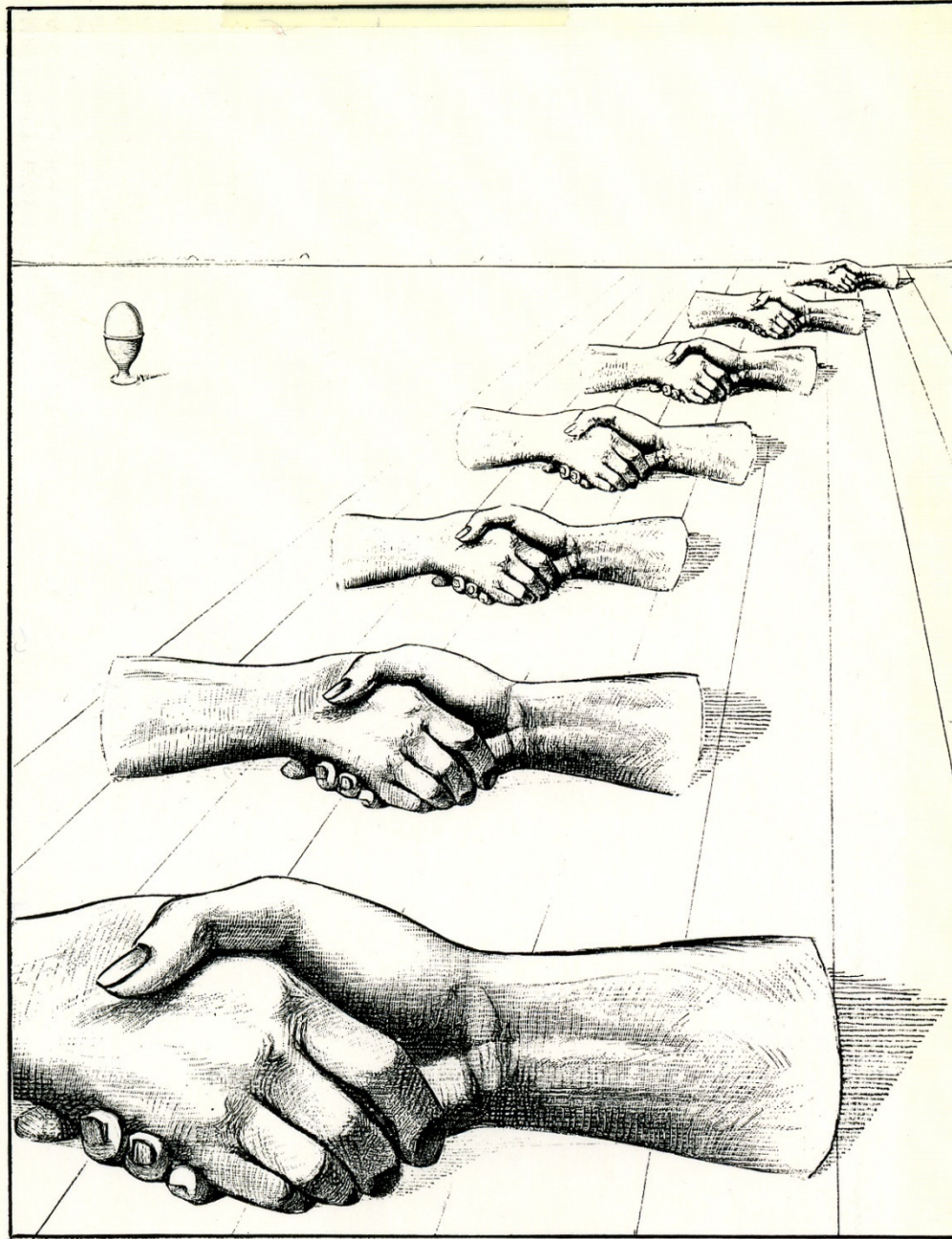
CONCLUSIONS

- Formidable challenges stand in the way of progress in multiscale modelling and simulation research. These challenges involve resolving open problems associated with simulation methods, appropriate quantitative data, model validation and verification and vizualization.
- Sustained research in multiscale modeling with lasting impact on bioprocess technology requires long term strategic planning. Progress in the field will require the creation of multidisciplinary teams that works together.
- Significantly, one of the challenges is cross disciplinary education to improve interdisciplinary communication between biologists, bioinformatitions, computer scientists, mathematicians ,physicists and engineers.

Acknowledgements

Alexej Lapin
Joachim Schmid
Klaus Mauch

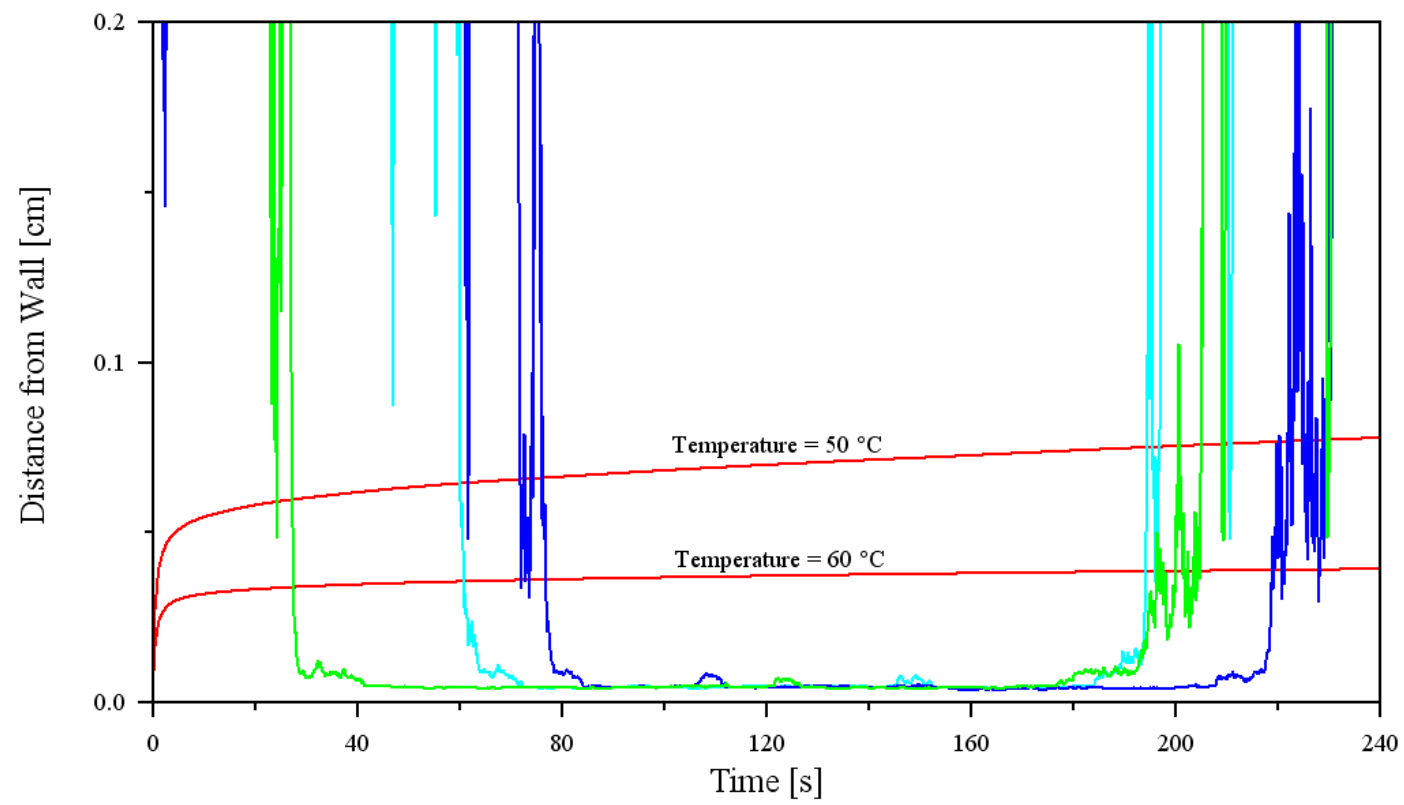
Funding: German Science Foundation (DFG)



MAX ERNST (1891-1976)

$37 \rightarrow 42^{\circ}C$





Modal Analysis:

$$c_i(t) = \bar{c}_i + \delta c_i(t)$$

$$\tau_i = \frac{1}{|\operatorname{Re}(\lambda_i)|} \quad i = 1, \dots, n$$

