

European experience of running Infrastructure Systems Biology: perspectives for Russia

Systems Biological modelling as a client-oriented service

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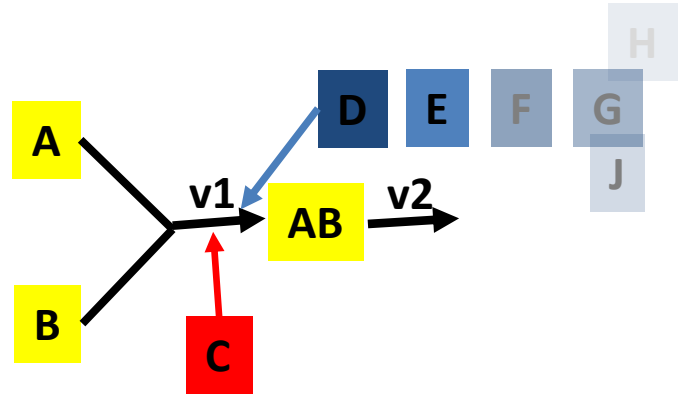
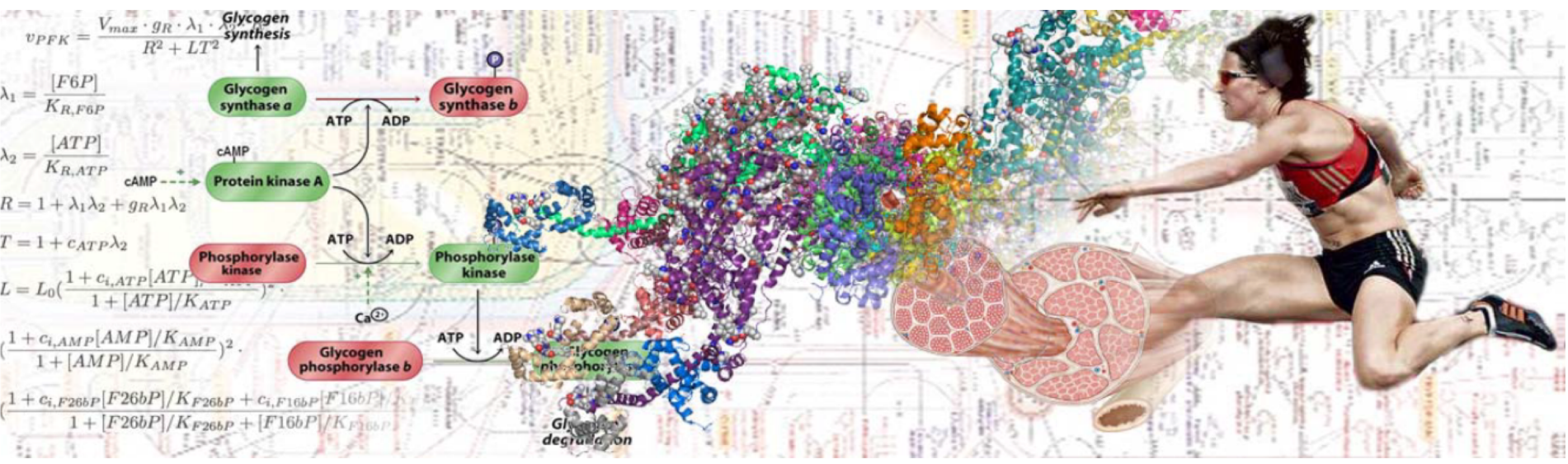
University Amsterdam



VU University

November 13, 2018

Systems Biology – reconstruction of biological emergence *in silico*

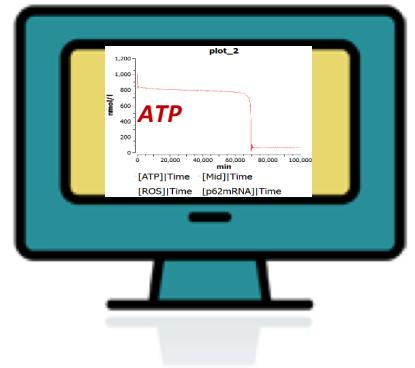


Solution through the systems biological approach:

Protein A + Protein B = Complex AB
 $v_1 = k_f[A][B][C] - k_b[AB]$
 $v_2 = \dots$
 $d[AB]/dt = v_1 - v_2 \dots$

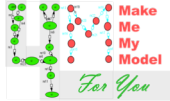
High state-dependency of component properties E.g. interactions between proteins A and B depends on

- Other components (C,D,E,F...)
- Hysteresis
- Flow of mass and energy through the system
- Initial and boundary conditions, etc.



ISBE

The Infrastructure Systems Biology Europe (ISBE) provides stewardship and help with biological and medical data, their acquisition, their analysis and their understanding. It consists of 5 interconnected infrastructure pillars:



- 1) The Make Me My Model (M4) pillar consists of a software infrastructure that helps customers to make their various types of data (genome sequence, transcriptome, proteome, metabolome, physiological, kinetic, etc.) predictive and understood via modeling (www.isbe.nl).



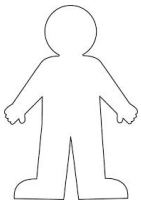
- 2) The Do Me an Experiment pillar is a distributed hardware-plus-service infrastructure that performs systems-biology quality assays as a service (M5; Make Me My Mass Spectra Measurements, enzyme kinetics, metabolomics, and epigenetics will be soon available. For the time being, there is a Systems Metabolomics service at ISBE.IT (www.sysbio.it/isbe): a complete metabolomic platform to perform ad hoc experiments, coupled with constraint-based modeling approach.



- 3) The Live Model Repository (LMR) of ISBE is a software infrastructure of interconnectable, systems-biology-quality kinetic models through JWS Online (<https://jjj.bio.vu.nl/>) and COSYS (sysbio.it/cosys/), where users can perform construction, modification, and simulation of kinetic models, and storage of curated ones (JWS), or define mathematical models of biological systems and perform constraint-based (e.g., Flux Balance Analysis) and mechanism-based dynamic simulations (either deterministic or stochastic), drastically accelerated by GPUs (COSYS).



- 4) The Data and Model Stewardship of ISBE called FAIRDOM (<http://fair-dom.org>). The FAIRDOM Project was started under the auspices of the ERA-NET programme ERASysAPP and ISBE, jointly funded by CH, DE, NL and UK. FAIRDOM assists researchers to be in control of collecting, managing, storing, and publishing data, models, and operating procedures. FAIRDOM takes responsibility for the Stewardship of research assets services of ISBE.



- 5) Help Me to Model (HMTM) provides training to customers wishing to make models themselves, in online or workshop tutorials (www.isb.nl).



Nodes

ISBE-light: Active nodes in:

Italy (Naples, Milano)

the Netherlands (Amsterdam, Leiden, Wageningen,
Eindhoven, Maastricht),

Slovenia (Ljubljana);

ISBE synergizes with other ESFRIs through Corbel

www.isbe.nl;

www.sysbio.it;

www.nh.cas.cz/isbelight

The Netherlands arm of the Infrastructure Systems Biology Europe (ISBE):



Alexey Kolodkin, Matteo Barberis, Ablikim Abdukerim, Zahid Hassan, Thierry Mondeel, Samrina Rehman and Hans V. Westerhoff



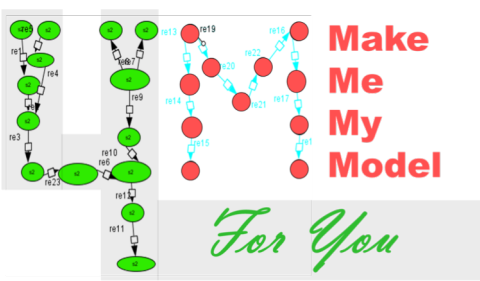
executive director



founding director

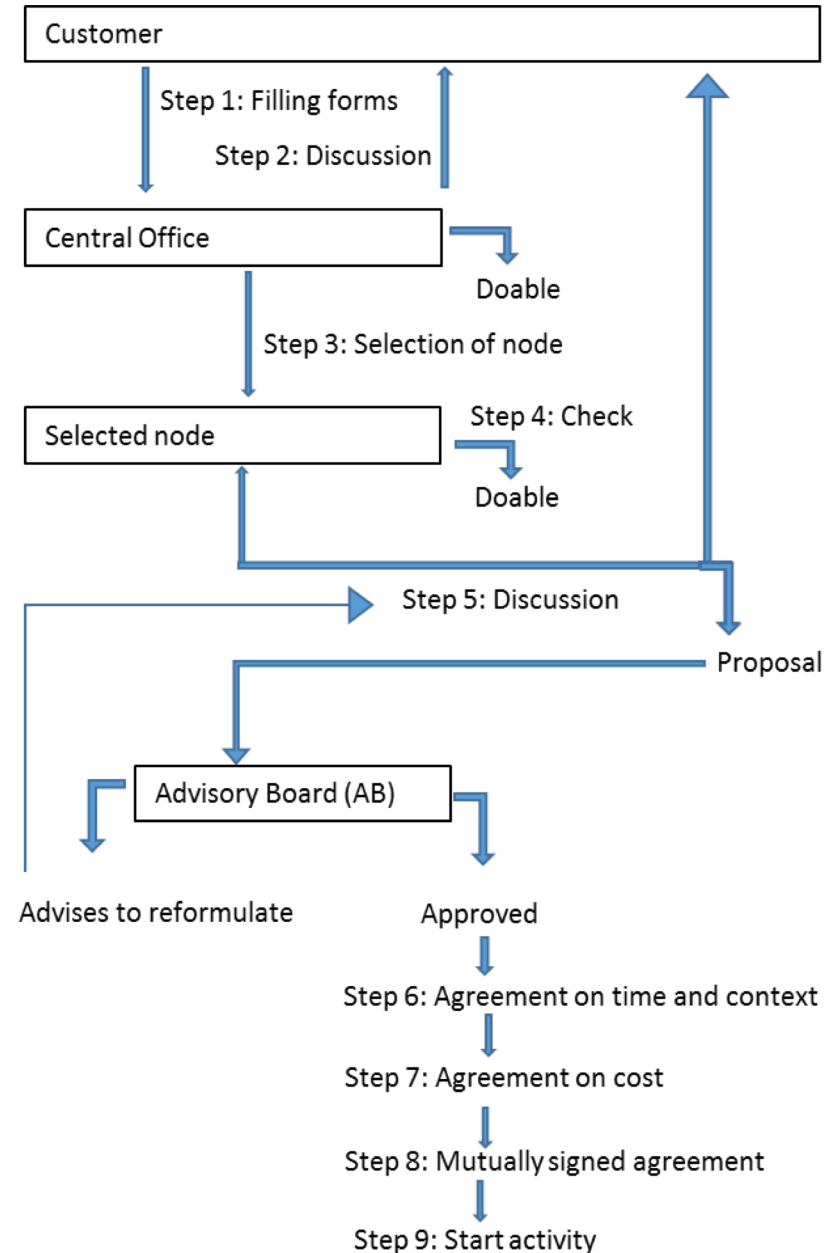
Stefania Astrologo, Ewelina Weglarz-Tomczak , YanFei Zhang, Jacky Snoep

- Executive director Dr Alexey Kolodkin (UvA, VU, LCSB, Corbel)
- Postdoctoral fellow Dr Ewelina Weglarz-Tomczak (UvA)
- Part-time help from Westerhoff group members (VUA, UvA, UoM)
- Pending integration with Wageningen (Vitor Martins dos Santos et al)



M4 mission:

to service the Life Sciences community by facilitating the implementation of systems biology



Proposed ISBE M4 access management

ISBE.NL services

The services provided at present

The collage features 11 project posters, each with a title, description, and associated researchers:

- 1. ROS management in Parkinson's disease and cancer (LCSB, Luxembourg)**: Focuses on ROS management in Parkinson's disease and cancer. Researchers: Prof. Dr. Jean-Louis Dreyfus, Prof. Dr. Jean-Louis Dreyfus.
- 2. Modelling ROS management and mitochondrial dysfunction (Milan)**: Focuses on mitochondrial dysfunction and ROS management. Researchers: Dr. Armanita Corbelli, Prof. L. Lazzarini.
- 3. Institute of Cytology and Genetics (Wrocław, Poland)**: Focuses on multi-level integration of signaling pathways. Researchers: Dr. Hugo Rocha, Prof. Margarita E. Gomez.
- 4. CFTR maturation (Lisbon, Portugal)**: Focuses on CFTR maturation. Researchers: Dr. Hugo Rocha, Prof. Margarita E. Gomez.
- 5. Safety assessment of endocrine disrupting chemicals (Tarragona)**: Focuses on safety assessment of endocrine disrupting chemicals. Researchers: Raju Prasad Sharma, Dr. Vikas Kumar, Prof. Marta Schramm.
- 6. Institute of Experimental and Clinical Medicine Laboratory of Molecular Mechanisms of Free Radicals Processes (Radicals Processes by TC13 for Parkinson's disease therapy)**: Focuses on modulation of ROS management by TC13 for Parkinson's disease therapy. Researchers: Mousa Amirshahi, Azhar Chachkhava.
- 7. Charged peptide to charged membrane binding model (Munich, Germany)**: Focuses on charged peptide to charged membrane binding model. Researchers: Dr. Fritz Löffler.
- 8. Modelling substrate diffusion and metabolism in biofilms (Helmholtz Center Munich, Germany)**: Focuses on modelling substrate diffusion and metabolism in biofilms. Researchers: Dr. Martin Elsner.
- 9. Projects with Essen (Germany): Previous services: Cool design of Hot metabolism in Archaea**: Focuses on cool design of hot metabolism in Archaea. Researchers: Dr. Martin Elsner.
- 10. Modelling fibromatosis (Dupuytren's disease), Chronic Myeloid Leukaemia (CML), and uterine cancer (Manchester, UK)**: Focuses on modelling fibromatosis, Chronic Myeloid Leukaemia, and uterine cancer. Researchers: Dr. Saverio Robustelli, Prof. Philip Day.
- 11. Institute of Experimental and Clinical Medicine Laboratory of Molecular Mechanisms of Free Radicals Processes (Radicals Processes by TC13 for Parkinson's disease therapy)**: Focuses on modulation of ROS management by TC13 for Parkinson's disease therapy. Researchers: Mousa Amirshahi, Azhar Chachkhava.

Central hub: ODE model 1 output, ODE model 2 output, ODE model 3 output.

ISBE.NL M4 services (currently running and planned to be completed within the next 12 months and in negotiation phase):

Completed services

1. University of Duisburg-Essen, Germany: Cool design of hot metabolism: GAPN
2. Sanquin, Amsterdam: Modelling of acute and chronic inflammation

Active services

3. LCSB, Luxembourg: ROS management in Parkinson's disease and cancer
4. Milano-Bicocca, Italy: Modelling ROS management and mitochondrial dysfunction
5. Sheffield, UK: Mitochondrial perfect adaptation
6. Lisbon, Portugal: CFTR maturation
7. URV Tarragona, Spain: Safety assessment of endocrine disrupting chemicals
8. Institute of Experimental and Clinical Medicine, Novosibirsk, Russia: Modulation of ROS management by TC13 for Parkinson's disease therapy
9. Universidade Católica Portuguesa, Portugal: Molecular Insight into Autism Spectrum Disorder (ASD)
10. Jožef Stefan Institute, Slovenia: Protease signaling network in neurodegeneration

Prospective services

11. Munich, Germany: Charged peptide to charged membrane binding model
12. Helmholtz Center Munich, Germany: Modelling substrate diffusion and metabolism in biofilms
13. Institute of Cytology and Genetics of RAS, Novosibirsk Russia: Merged ODE and agent-based model for multilevel integration of signalling pathways
14. Manchester, UK: Modelling fibromatosis (Dupuytren's disease), Chronic Myeloid Leukaemia (CML), and urothelial cancer
15. University of Duisburg- Essen, Germany: The Yin-Yang of Metabolism; Endometatotoxicity (YYME)

Findable, Accessible, Interoperable, Reusable (FAIR) Model management for ISBE services in FAIRDOME

The screenshot shows a web browser window with the FAIRDOME website. The browser's address bar shows the URL <https://fairdomhub.org/programmes/27#projects>. The website header includes the FAIRDOME HUB logo, a search bar, and the user name Alexey Kolodkin. The main content area features a breadcrumb trail: Home / Programmes Index / Model repository for M4 (Make Me My Model) clients of ISBE. The title of the page is "Model repository for M4 (Make Me My Model) clients of ISBE". A description states: "ISBE-Light provides M4 service (Make Me My Model) where non modelers can request (assistance with) the making of a computational model of their biological system. These models are deposited here." Below the description, there are links for "Web page: <http://www.isbe.nl>" and "Programme Administrators: Alexey Kolodkin". A "Funding details" section indicates "No funding details specified". On the right side, there is an "Administration" dropdown menu, a "Change picture" button, and a "Storage Usage" section. A Skype notification is visible in the bottom right corner, showing "Venkata Satagopam is online".

Home / Programmes Index / Model repository for M4 (Make Me My Model) clients of ISBE

Model repository for M4 (Make Me My Model) clients of ISBE

Administration

ISBE-Light provides M4 service (Make Me My Model) where non modelers can request (assistance with) the making of a computational model of their biological system. These models are deposited here.

Web page: <http://www.isbe.nl>

Programme Administrators: Alexey Kolodkin

Funding details:

No funding details specified

Change picture

Storage Usage

skype

T Venkata Satagopam
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is follows

5 CORBEL supported services



VIP: Margarida D. Amaral - Cystic Fibrosis Transmembrane Regulator maturation

Anna Maria Colangelo - Modelling ROS management and mitochondrial dysfunction in models of Parkinson disease

VID: 3441 - Molecular Cell Physiology, Vrije University Amsterdam Awaiting Confirmation | Fitting the dynamic model to experimental data

CORBEL Track 2VID: 3444 - Chemogenomics (ChEMBL) at EMBL/EBI Awaiting Confirmation | Profiling of chemotypes for potential off-target effects, measured ADMET properties, in-vivo efficacies

CORBEL Track 2VID: 3447 - Advanced Light Microscopy Facility at EMBL Awaiting Confirmation | Fluorescence resonance energy transfer (FRET) - EMBL HD Awaiting Confirmation | Electron microscopy - EMBL HD

Vikas Kumar EDC-SysTox: Approaches towards Systems Toxicology model via coupled PBPK/PD-system biology benchmarking dosimetry for safety assessment of Endocrine Disrupting Chemical

VID: 3421 - Molecular Cell Physiology, Vrije University Amsterdam

CORBEL Track 2VID: 3422 - Chemogenomics (ChEMBL) at EMBL/EBI

CORBEL Track 2VID: 3423 - Screening and medicinal chemistry at Leibniz-Institute for Molecular Pharmacology (FMP)

Veronika Stoka Jožef Stefan Institute (Slovenia) - Protease signaling network in neurodegeneration

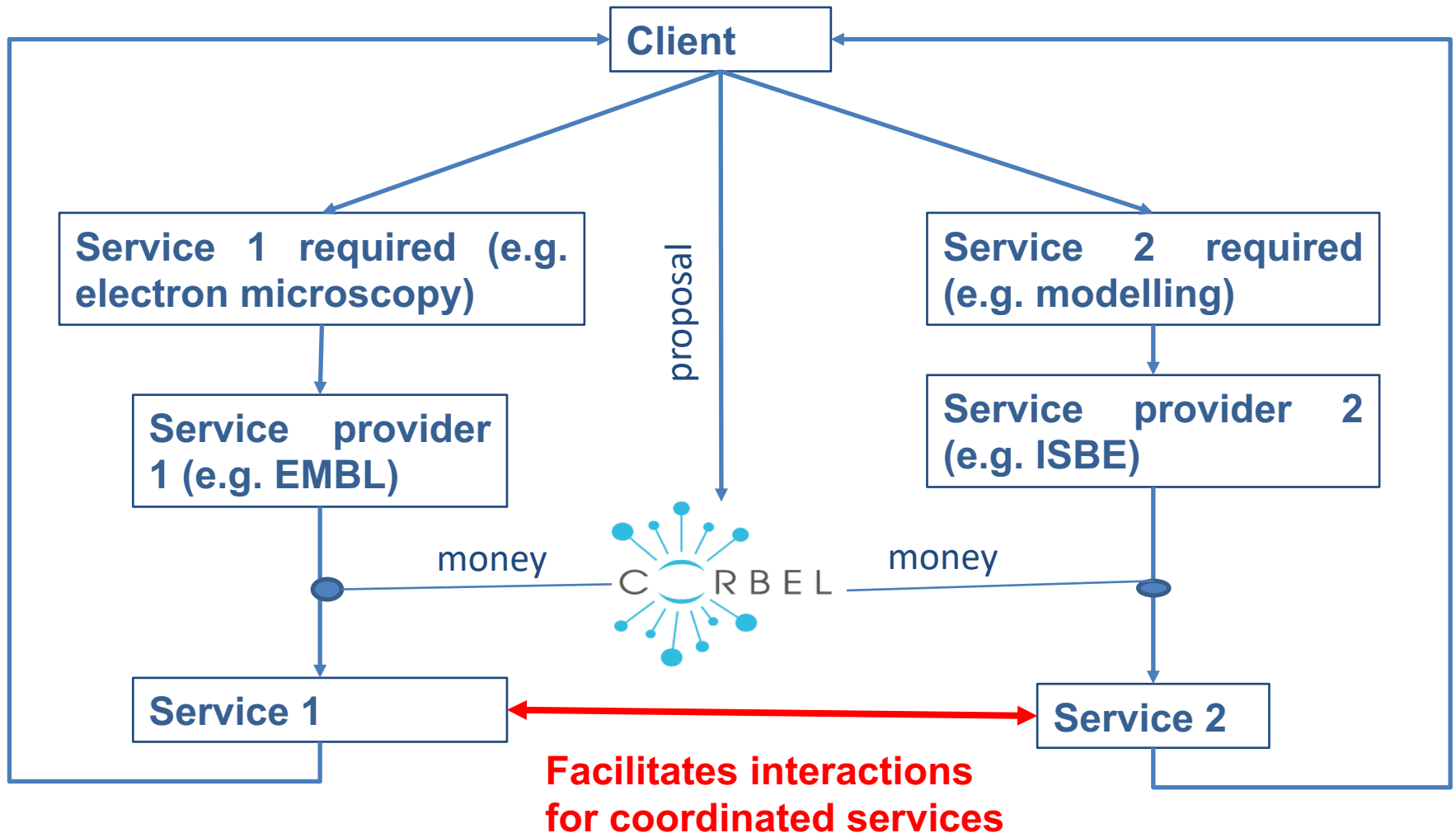
With Chemogenomics (ChEMBL) at EMBL/EBI; Advanced Light Microscopy Facility at EMBL; - Screening and medicinal chemistry at Leibniz-Institute for Molecular Pharmacology (FMP)

Maria Correia - Molecular Insight into Autism Spectrum Disorder (ASD)

1VID: 3248 - Biological Research Foundation Academy of Athens BRFAA Awaiting Confirmation | Genomics and transcriptomics services

CORBEL Track 1VID: 3530 - Molecular Cell Physiology, Vrije University Amsterdam

Experience from CORBEL supported services: Integration of Systems Biological Infrastructures



ISBE at Novosibirsk (Institute of Cytology and Genetics)

Teaching (February 2017, October 2017)

“Systems Biology and Personalised Medicine” course

72 hours (lectures, seminars, tutorials individual work and exam)

Lectures: systems biology aims to reconstruct *in silico* the emergence of biological function in terms of interactions between biomolecules

Principles

Training in building mechanism based dynamic model (with ODE)

Approaches of model analysis

Application of modelling for personalized medicine

principles of modeling, building simple models on the back of envelope

using special softs (COPASI and Cell Designer) to build toy models, using already developed model

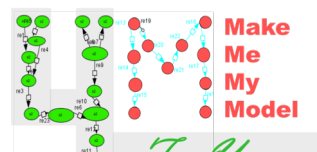
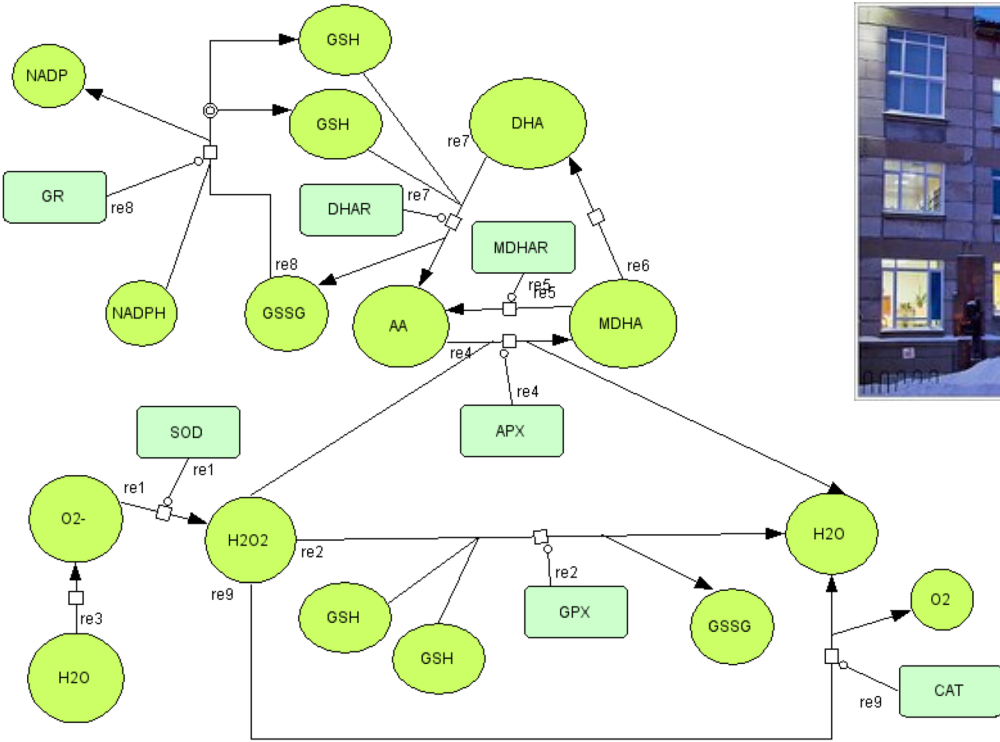
using already developed models, design principle studies, MCA

Practical individual work

From course to modelling research

ROS model for plants instantiation

Institute of Cytology and Genetics (Novosibirsk, Russia): Cold resistant wheat

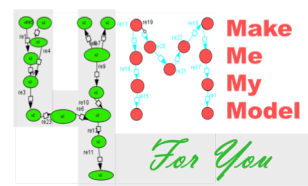


Aleksandr Bobrovskih



Alexey Doroshkov

CFTR maturation (Lisbon, Portugal) With student from NSU/ICG



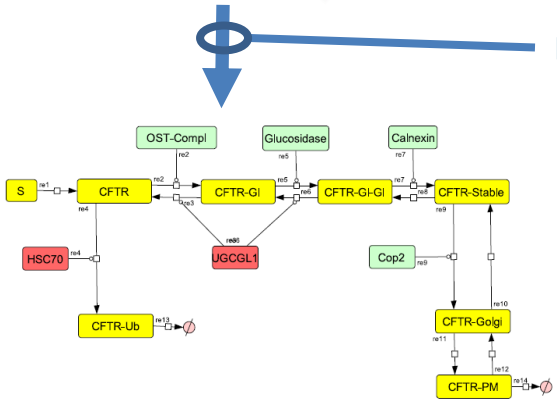
Cystic Fibrosis (CF)- lethal autosomic disease
CFTR - Cystic Fibrosis Transmembrane Regulator



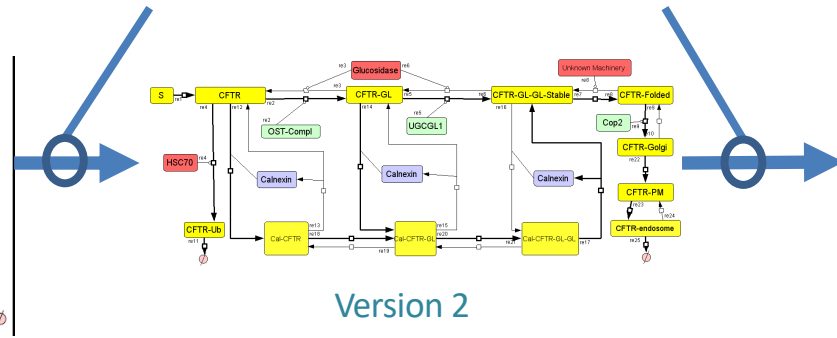
Dr. Hugo Botelho

Prof. Margarida D. Amaral

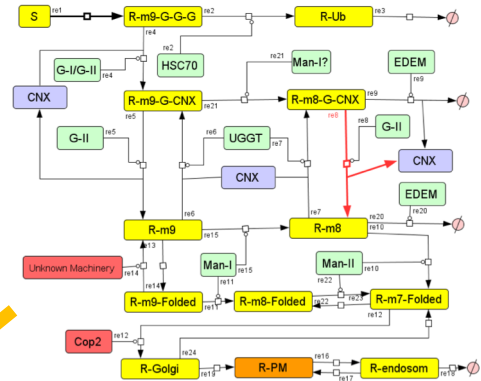
Daria Astapenko (NSU-ICG)



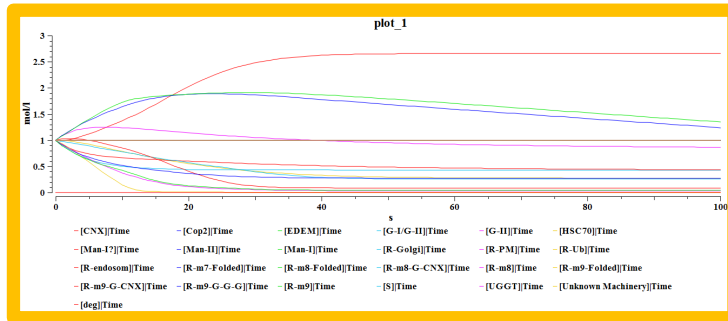
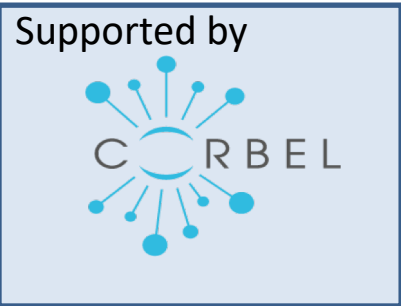
Version 1



Version 2



Version 3



Version 4

ISBE at Novosibirsk (Institute of Cytology and Genetics)

Since February 2017

LETTER OF INTENT

with respect to the collaboration between the Federal Research Centre “the Federal Research Centre Institute of Cytology and Genetics of the Siberian Academy of Science”, Russian Federation and the Centre of Systems Biology “Systems Biology Amsterdam”, the Netherlands

The Federal Research Centre “the Federal Research Centre Institute of Cytology and Genetics of the Siberian Academy of Science” hereafter called «ICG SB RAS», represented by its director Kolchanov Nickolay Alexandrovich and bound by the rules of aforementioned organization, on the one side, and the Centre of Systems Biology “Systems Biology Amsterdam” (hereafter SysBA¹) represented by its director Westerhoff, Hans Victor, on the other side, called together as «Parties» are herewith signing this agreement about the following:

- Parties, i.e. the Division of Systems Biology «ICG SB RAS» (the head of the division is Kolchanov N.A.) on the one hand and «SysBA» on the other hand declare their willingness to collaborate with respect to the following:

630090
 Phone: (383) 363-49-80 Fax: (383) 333-12-78
 University of Amsterdam
 Science Park, 904 (room C2.103) NL1098 XH
 Amsterdam (the Netherlands), EU



Director of ICG SB RAS
 Kolchanov Nikolay A.
 Responsible person from ICG SB RAS
 Head of sector
 Lashin Sergey A.

Director of SysBA
 Westerhoff Hans V.
 Responsible person from SysBA
 Executive director of ISBE.NL
 Kolodkin Alexey N.

August 2017

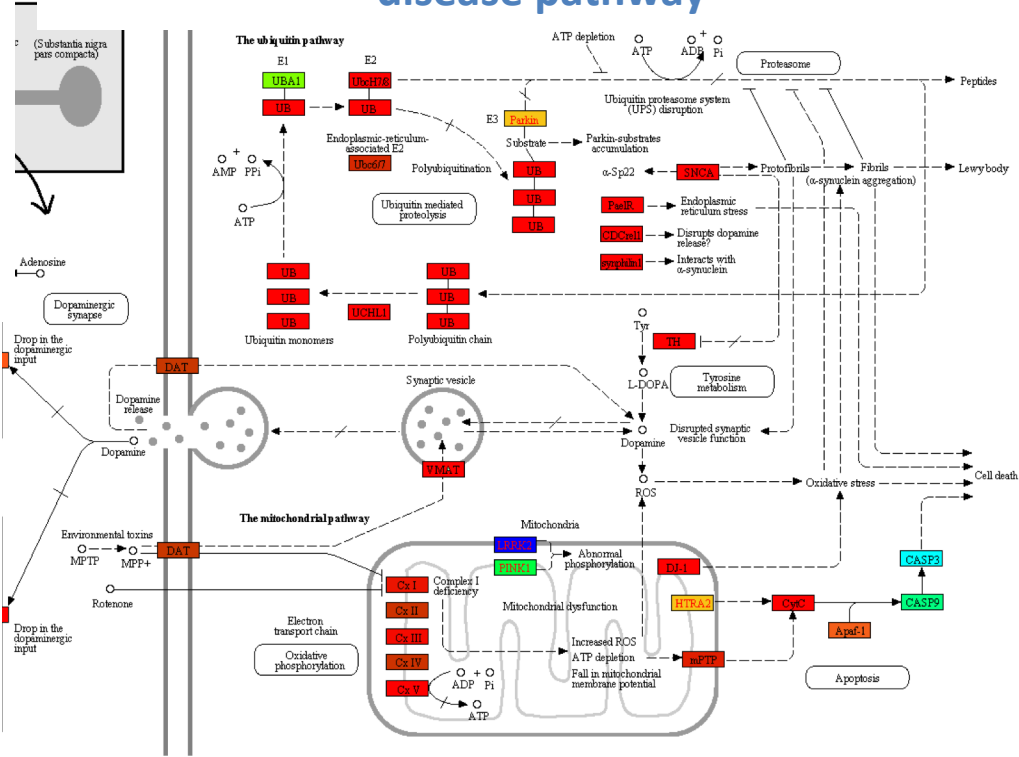
Vrije Universiteit Amsterdam
 Faculty of Sciences
 De Boelelaan 1083
 1081 HV Amsterdam
 The Netherlands

05012 1/2014
 (c) Kazelissa Laboratories

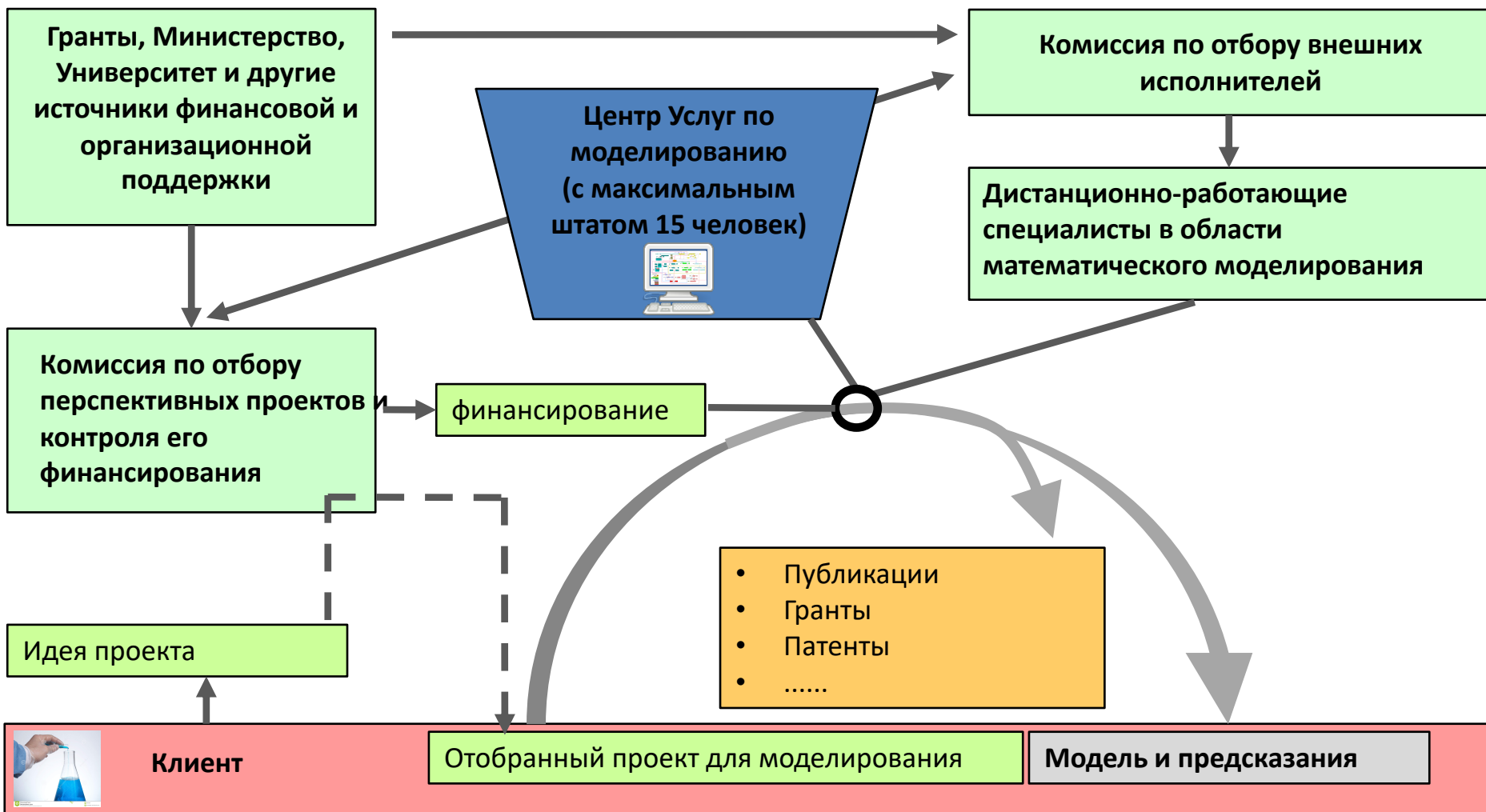


Nickolay A. Kolchanov Sergey A. Lashin

Several projects, e.g. DI analysis of Parkinson's disease pathway



Развитие потенциального Симстемно-Биологического центра: Компьютерное моделирование как сервисная услуга



Благодарю за внимание!