



CompuBioTic, a methodology for *de novo* design of vesicles performing programmed tasks

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Complex System Modelling and Engineering for Diagnosis
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Our goal

Design and build a **non-living** synthetic nanosystem,
« **programmed** » to perform *in vitro* or *in vivo*
diagnostic assays

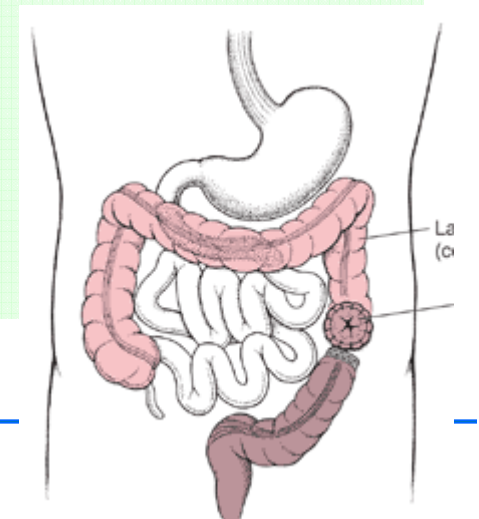
Colorectal cancer diagnosis and follow-up

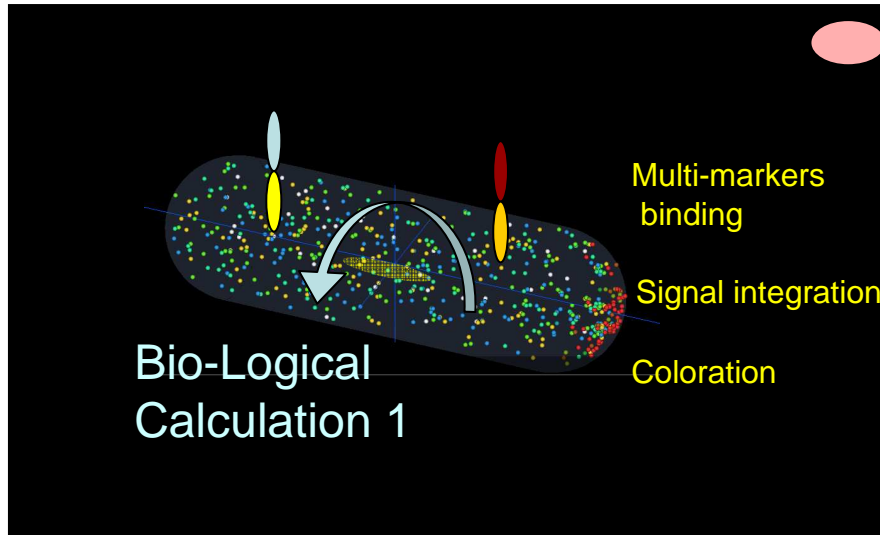
« Close to the patient » simple assay

Multi-parametric measure

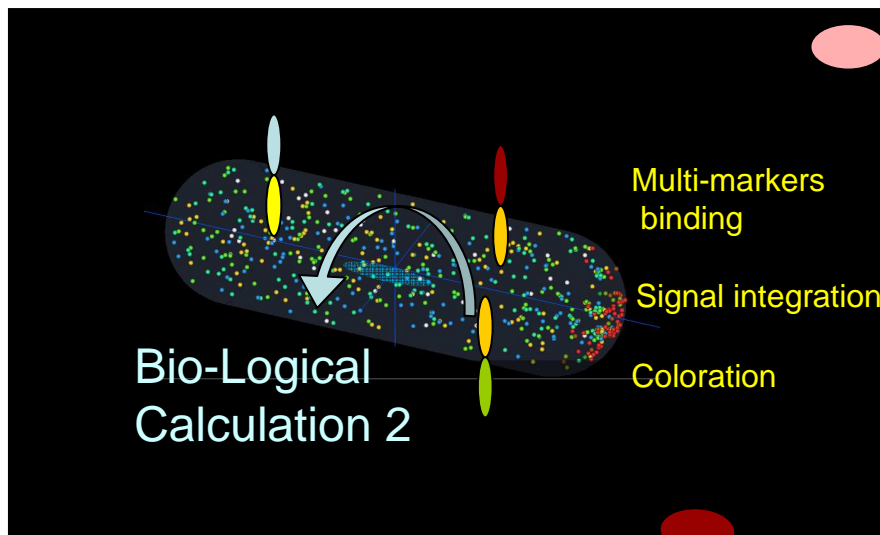
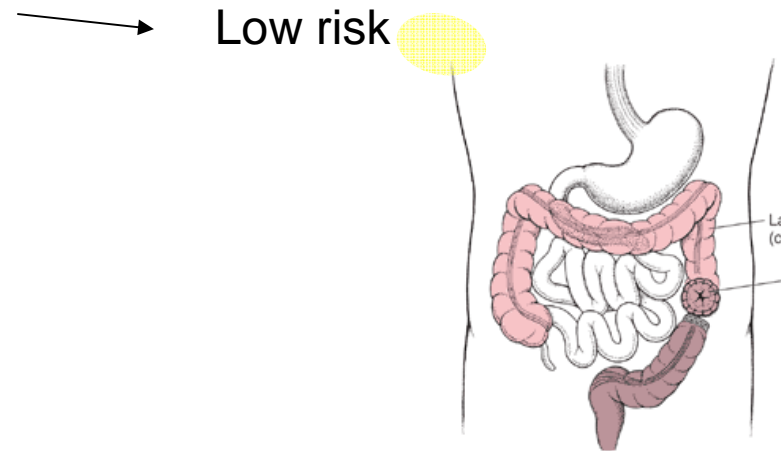
Sophisticated signal integration
(qualitative, quantitative, temporal, spatial etc.)

Result return in a simple way (local dyeing)





Local coloration depends on the response of multi-sensors to local biomarkers and the result of integrated « bio-logical calculation »





Our Strategy

in vitro Proof of concept

Design

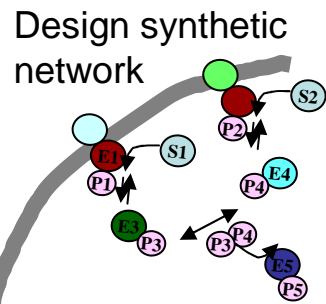
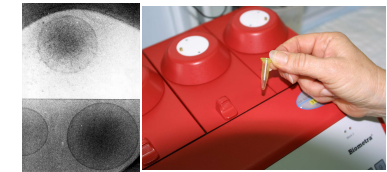
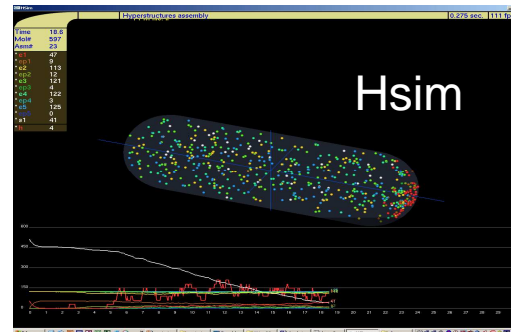
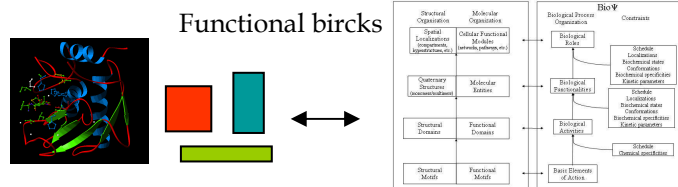
Simulation

Experimental validation

Identification and characterization of molecular compounds

Simulation

Stochastic Cellular automaton, Multi-Agent



BioPsi Formalization

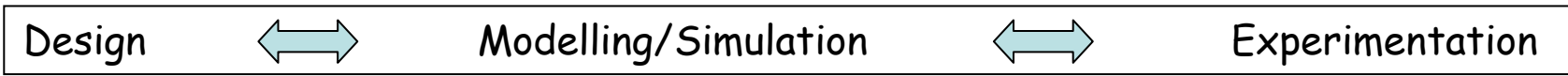
Modelling:
Control Th.
flux analyses
ODE
Elementary modes



Biotechnology

**Auto-organisation/
Robustness**

Validation *in vitro*





Design

Standardized catalog of proteic biological compounds:

- Processes are formalized (ready for modelling)
- Biological behavior characterized experimentally (ready to use in a synthetic system)

- « compound » properties
- robustness
- stability
- functional diversity relative to context modifications

```

Generic BioPsi description
Biological Roles description
BREnzyme-linkedTransmembraneConditionalTest :=
BFTransmembraneReceptor(Ligand) IN transmembrane |
BF_Ligand(TransmembraneReceptor) IN extracell |
BFTransmembraneReceptor/Ligand(Substrate) IN transmembrane |
BFSubstrate(TransmembraneReceptor/Ligand) IN cytosol,
BFProduct IN cytosol

Biological Functionalities description
BFTransmembraneReceptor(Ligand) :=
  BA_Binding(Ligand)
  IN extracell

BFTransmembraneReceptor/Ligand(Substrate) :=
  BA_ConformationalChange,
  BA_EnzymaticActivity(Substrate) -> Product
  IN cytosol

BF_Ligand(TransmembraneReceptor) :=
  BA_ProteinBinding(TransmembraneReceptor)
  IN extracell

BFSubstrate(TransmembraneReceptor/Ligand) :=
  BA_ProteinBinding(TransmembraneReceptor)
  IN cytosol

BFProduct
  BA_ProteinBinding-1(TransmembraneReceptor)
  IN cytosol
    
```

SysDiag FRE3009 CompuBiotic project Contact

Reusable Molecular Elements Catalog

Catalog home | References | External links

Catalog Home

Actions at the Compartmental environment scale

- redox conditions keeper
- pH conditions keeper
- temperature conditions keeper

Actions at the System scale

- timekeeper
- switch
- killer
- scaffold controller
- fuel

Actions at the Module scale

- conditional sensor
- revealing
- amplifier
- inhibitor
- distributor
- revealing
- cargo
- conductor

Actions at the Molecule scale

BIOPsi processes used in the current module

Biological Role

- BR_TransmembraneConditionalTest

Biological Functionalities

- BF_TransmembraneReceptor
- BF_Ligand

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Actions at the Molecule scale

BIOPsi processes used in the current module

Biological Role

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Biological Functionalities

- BF_TransmembraneReceptor
- BF_Ligand

Catalog > Module scale > Conditional Sensor (CS) > Enzymatic Transmembrane CS

1 TransMembrane CS

1.1 Enzymatic TransMembrane CS

An enzymatic transmembrane sensor is a receptor, where the binding of an extracellular ligand leads to an enzymatic activity of the receptor himself, in the intracellular side.

cytoplasm

substrate(s)

product(s)

List of BioPsi processes used in the current module

Biological Role

- BR_TransmembraneConditionalTest

Biological Functionalities

- BF_TransmembraneReceptor
- BF_Ligand



Molecular compounds : Identification, characterization

- Modules with defined role

Redox conditions monitoring

pH conditions monitoring

Temperature conditions monitoring



Timekeeper (Time counter)



Switch On/Off



Killer (destruction component)



Scaffold controller



Fuel

Conditional Sensor

Amplifier

Inhibitor

Distributor

Revealing



Cargo

Conductor

- Proteins and small molecules useful for synthetic biology

Ex : Peroxidase



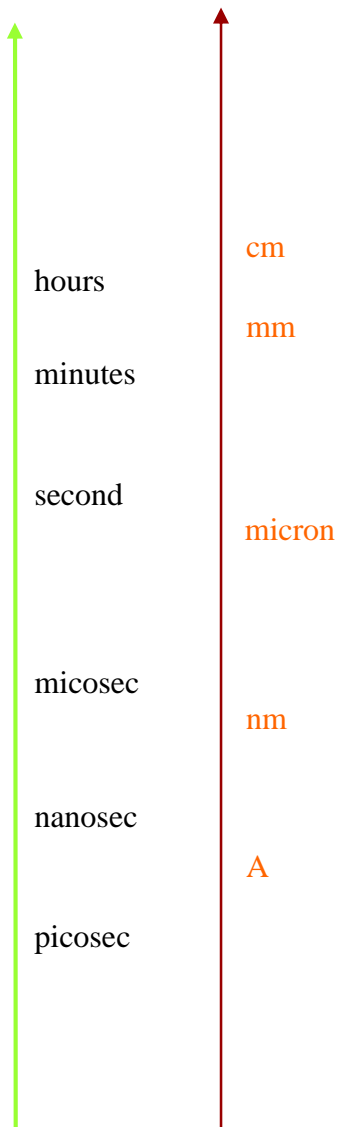
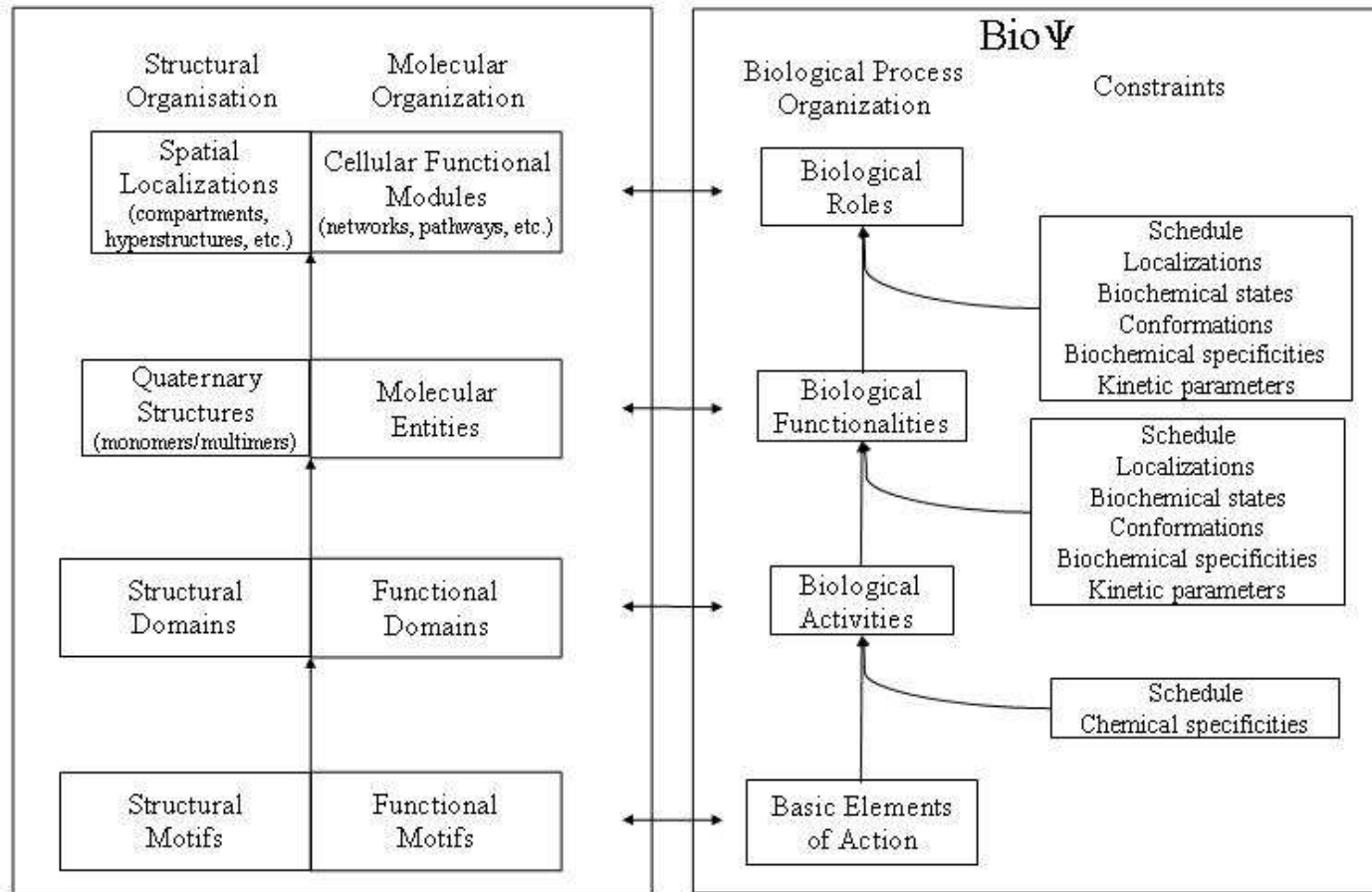
Revealing role



The BioΨ language

Molecular description

Processes description





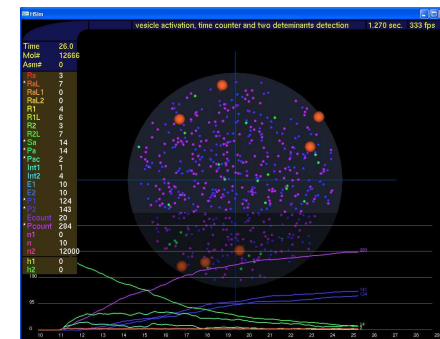
Bio Ψ what for ?

Modelling and simulation of biological processes

Elementary bricks of process at each level can be considered as primitives for formal language construction.

Genericity of processes description is an advantage for a non deterministic modelling

Easy to use with multi-agents approaches

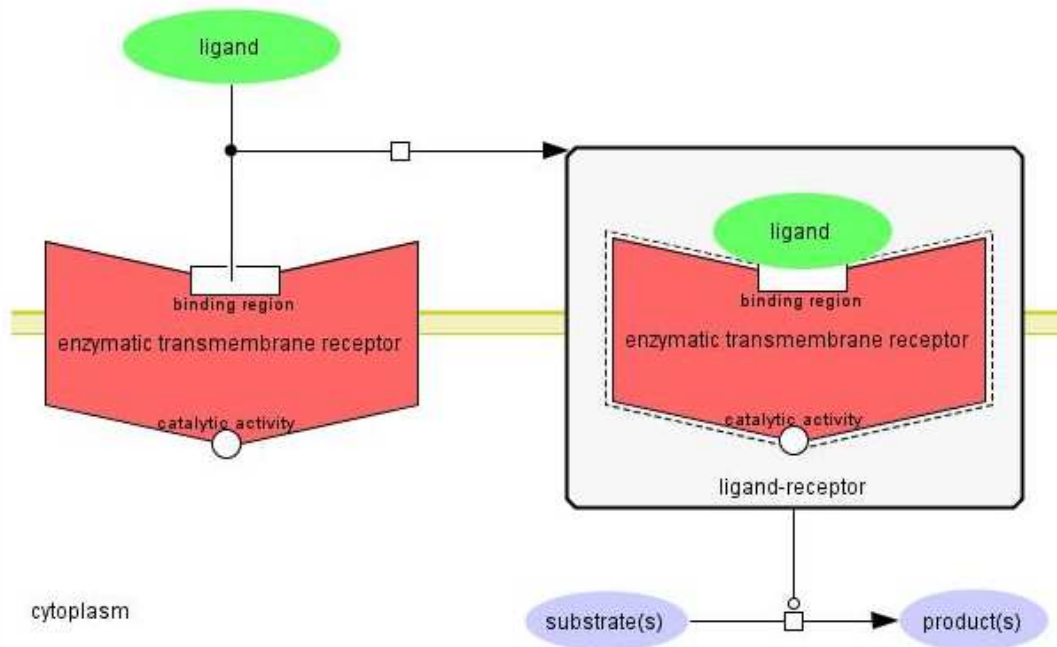


Conditional sensor

1 TransMembrane CS

1.1 Enzymatic TransMembrane CS

An enzymatic transmembrane sensor is a receptor, where the binding of an extracellular ligand leads to an enzymatic activity of the receptor himself, in the intracellular side.



List of Bio Ψ processes used in the current module

Biological Role

- BR_TransmembraneConditionalTest

Biological Functionalities

- BF_TransmembraneReceptor
- BF_Ligand

Generic BioΨ description

Biological Roles description

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BREnzyme-linkedTransmembraneConditionalTest :=  
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  BFLigand(TransmembraneReceptor) IN extracell,  
  BFTransmembraneReceptor/Ligand(Substrate) IN transmembrane |  
  BFSubstrate(TransmembraneReceptor/Ligand) IN cytosol,  
  BFProduct IN cytosol
```

Biological Functionalities description

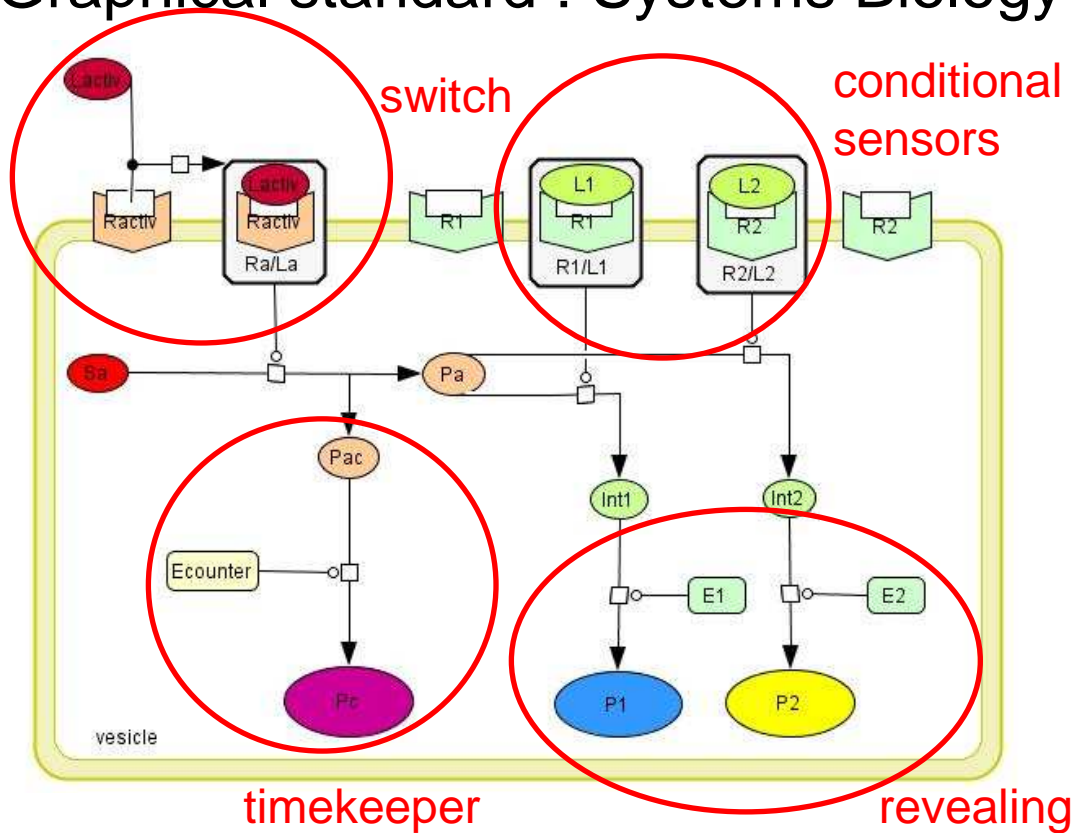
```
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  BA_Binding(Ligand)  
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  BA_EnzymaticActivity(Substrate) -> Product  
  IN cytosol  
  
BFLigand(TransmembraneReceptor) :=  
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BFSubstrate(TransmembraneReceptor/Ligand) :=  
  BA_ProteinBinding(TransmembraneReceptor)  
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BFProduct  
  BA_ProteinBinding-1(TransmembraneReceptor)  
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```



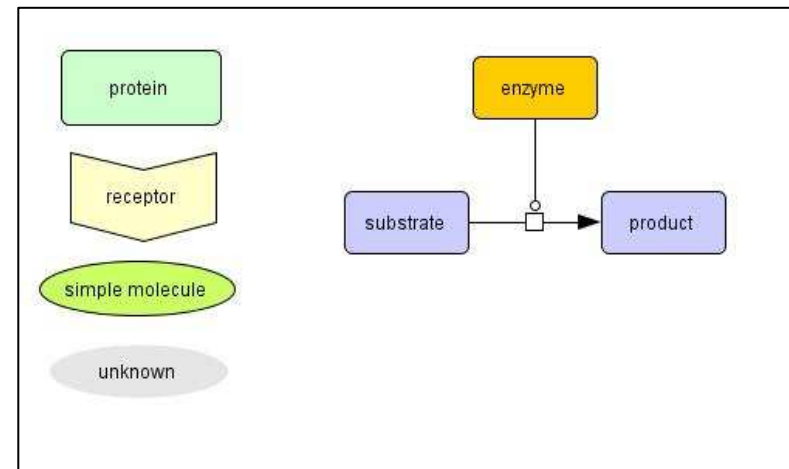
Nanobiosystem *in silico* design

Graphical representation tool : CellDesigner

Graphical standard : Systems Biology Graphical Notation (SBGN)



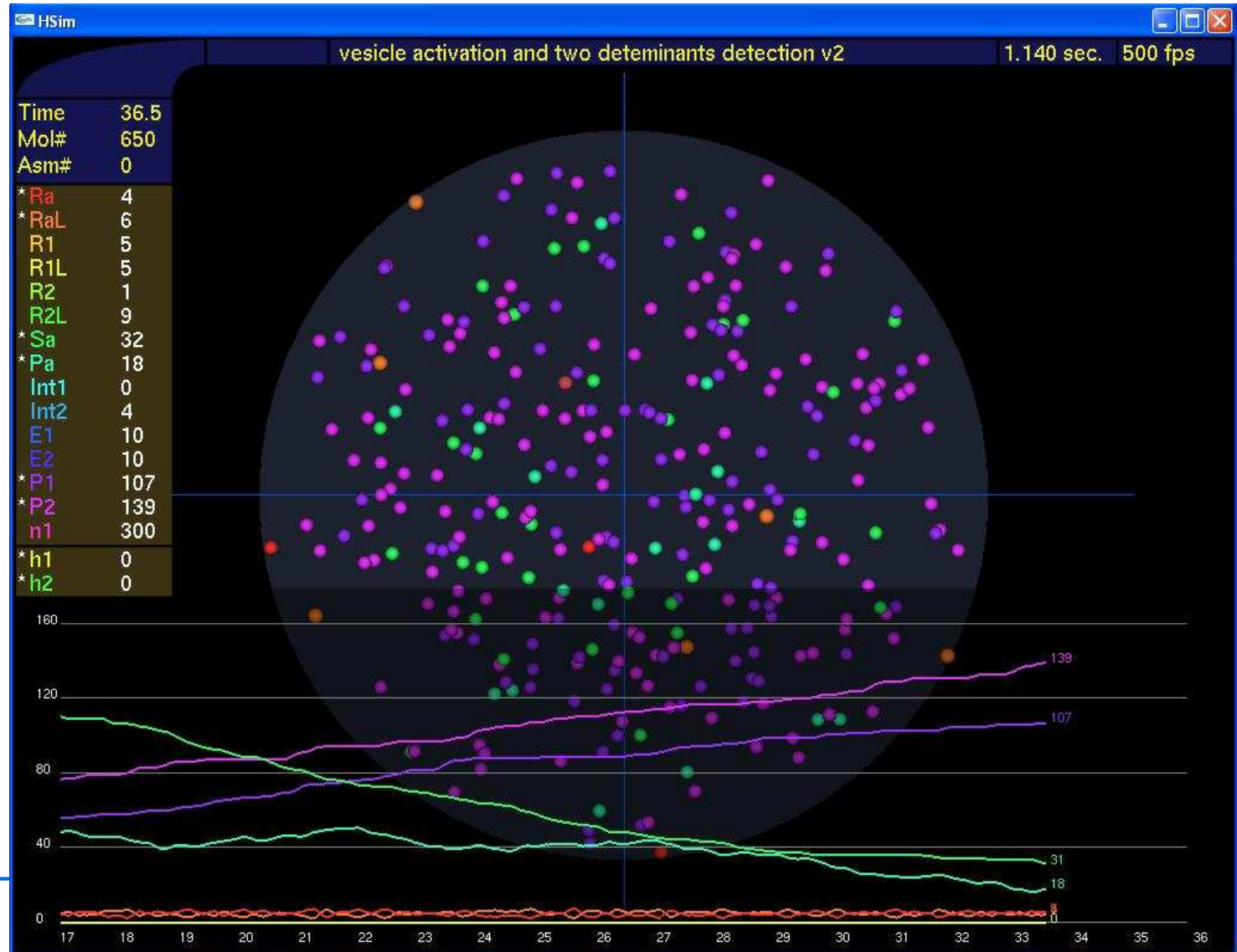
Légende



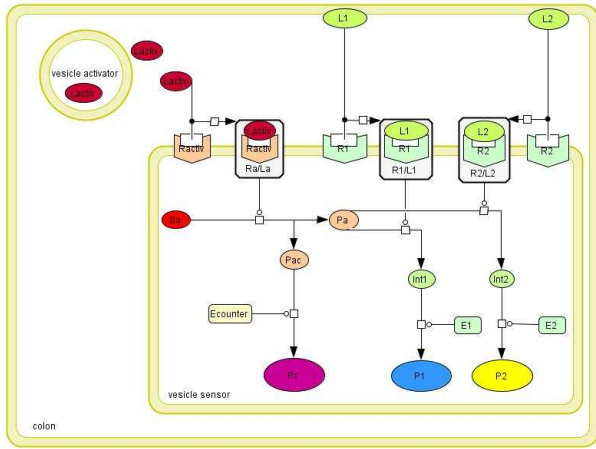


Modelling, simulations, analyses

HSIM
(Amar, 2004)

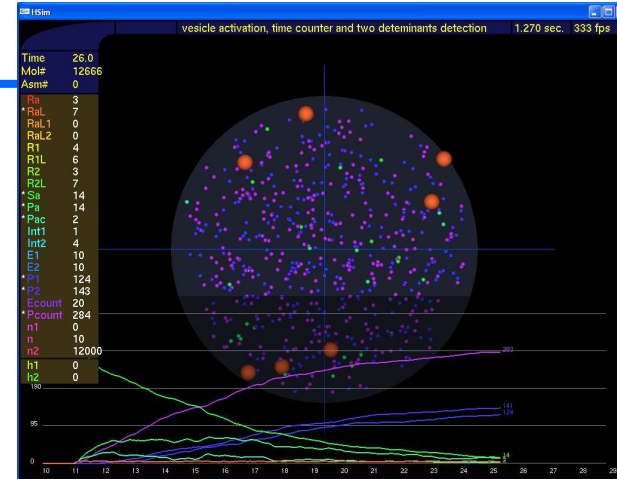


Simulation



```

Biological State description
Biological Role description
Biological Functionalities description
    
```



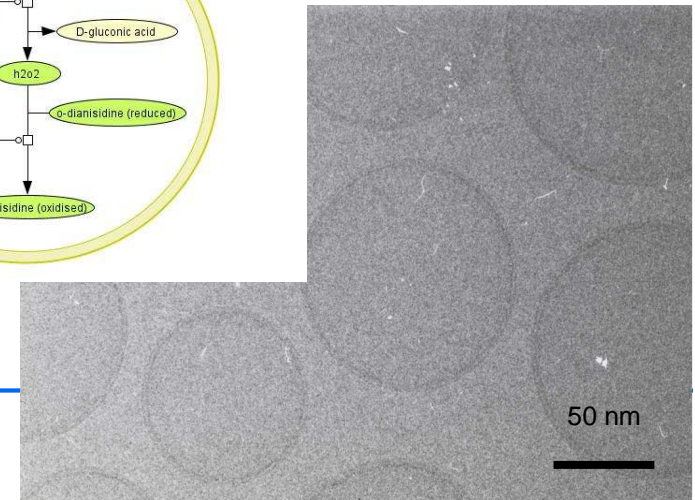
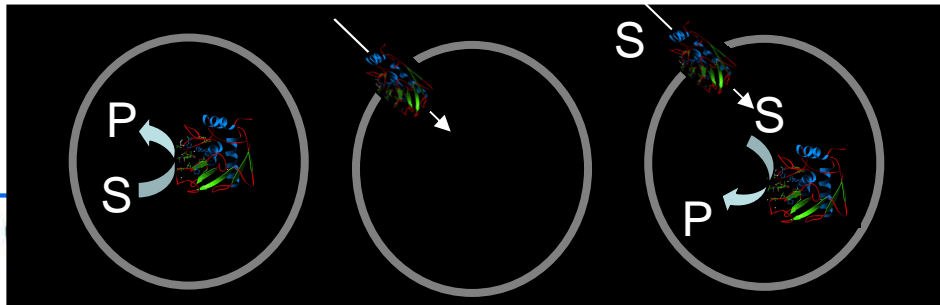
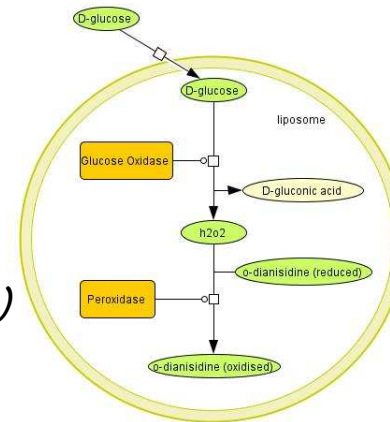
System design using our compound catalog
SBGN and *CellDesigner*

BioΨ modelling
Formal description

Simulation : Stochastic Cell automaton
and multi-agent

Experimental validation

- ☒ Stable Vesicles construction (liposomes) ~100nm
- ☑ Introduction of chosen functional compounds
- ☐ Opérational assays of full synthetic system (*in vitro*)





Expectations

- **Principles** and **method** for synthetic biological systems design from protein parts
- Set of **characterized "compounds"** re-usable in various synthetic systems
- **Formal description** of biological processes, including a description at the **protein domains level**
- A synthetic "bio-machine" dedicated to colorectal cancer **diagnosis**

CompuBioTic DB + CellDesigner + HSim Integrated tools for Synthetic Biology

Project team:

Franck Molina, SysDiag director
Stéphanie Rialle Doctorante BDI
Sabine Peres Postdoc
Alain Thierry Inserm
Liza Felicori Postdoc

External collaborators :

Patrick Amar, LRI, Orsay, Simulation
Marc Ychou, CRLCC, Montpellier Oncology colo-rectale
Andrew Griffiths, CNRS Strasbourg, nanodroplets
European BaSysBio partners UE FP6 syst. modelling



Thank you for your attention

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Modélisation et Ingénierie des **S**ystèmes Complexes Biologiques pour le **D**iagnostic

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