This subject requires common and basic knowledge.

The internship will be supervised by François Fages and Sylvain Soliman.

Assignment

The chemical reaction network formalism (CRN) is used to model biological processes at the cellular level. They explain the complex phenotypes as resulting from elementary molecular interactions. A CRN has a hypergraph structure (that is, a bipartite species / reaction graph labeled by a rate function) and can be interpreted at different levels of abstraction with different dynamic behaviors: differential, stochastic, discrete Petri net or Boolean.

Visualization of the structure of CRNs is not a well-solved problem today, and the automatic drawing tools are far from being able to compete with the manual drawings that modelers make in practice.

The aim of the internship is to try to use this static information (calculated by BIOCHAM) on the network structure so as to define drawing strategies and to set up the visualization tools of the interaction graph (Graphviz) so that he produces automatic drawings close to the manual drawings found in the publications.

Main activities

Our previous work on some BIOCHAM static analyzers is described in


The work will involve using this information to design and experiment with different species placement strategies and reactions in the GraphViz visualization tool. The evaluation can be done on the BioModels model warehouse by comparing the results with the drawings of the corresponding publications.

Results may be published in the Systems Biology Graphical Notation (SBGN) community.

Skills

This subject requires common and basic knowledge.
in algorithmics, programming, graphs and graphics.
Specific knowledge of the Prolog programming language or of Computational Systems Biology will be a plus.

**Benefits package**
- Subsidised catering service
- Partially-reimbursed public transport
- Social security
- Paid leave
- Flexible working hours
- Sports facilities

**Remuneration**
500 euros/month