

# Master 2 of Science in Computational Biology and Biomedicine

*Parcours du Master d'Informatique, spécialité Fondements et Ingénierie (Now), BIM (2012)*

Nice Sophia Antipolis University, France

Coordinators: E. De Maria, T. Papadopoulo

Teaching language: English  
Public: International attendance, applied mathematicians,  
Computer science students,  
Biologists (with good mathematical background)



## Computational biology and biomedicine (CBB)

An emerging **interdisciplinary** field that applies the techniques of **computer science, applied mathematics and statistics** to address biological/medicine problems.

- **Mathematical** modeling,
- **Computational** simulation techniques.
  
- Research Master 2 program CBB (3 years old).
- Hosted by the **Computer Science Department** of Nice Sophia Antipolis University (P. Lahire, J. Farré, C. Peyrat).
- Part of the BIM (Biology-CS-Maths) master program, starting in 2012 (F. Diener).

**Fellowships** available for foreign students (7) (800€/month).

# Organisation

**1 year** (september, 15<sup>th</sup> – end of august), applications up to june.

## **4 periods:**

- Basics in Mathematics and Biology (September).
- 2 periods of lectures: october-november, january-february.
- Training period: march-august (30 ECTS) + project (6 ECTS).

## **Lectures:**

- 2 ECTS per lecture.
- Half-day lectures (once per week).
- Mandatory courses (9).
- Elective courses (3).
- Total: 60 ECTS (10/20 on every course).

<http://cbb.unice.fr>  
[cbb@unice.fr](mailto:cbb@unice.fr)

November 12, 2030

## Dr House meets his patient Bill

Bill Krivitz suffers from knee arthritis...

Severe pain and reduced flexion



Dr House first looks for a medication?

- Sequences Bill's genome and tracks deficient genes
- Seeks drugs fixing the protein which malfunctions

Dr House and Bill agree on surgery

- Design of a patient specific prosthesis: pre-op simulations
- Computer monitored physiotherapy

Epilogue: Bill run the Boston Marathon

# Focus on the **human** being

From different **perspectives**:

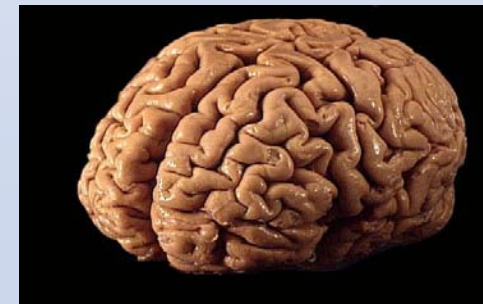
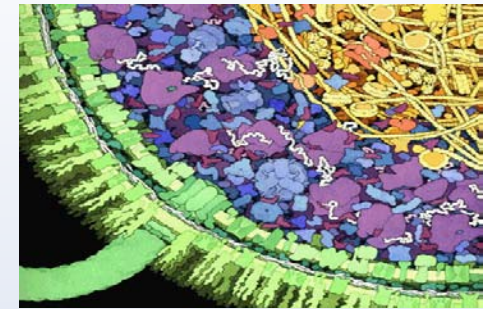
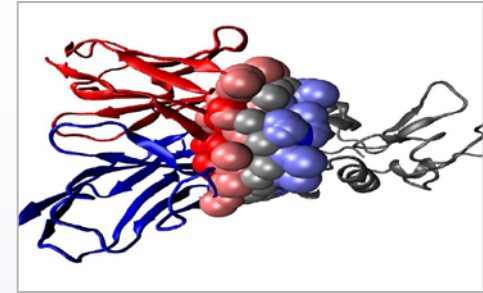
- Understanding and modeling functional aspects.
- Interpreting biomedical signals for various devices.

At different **scales**:

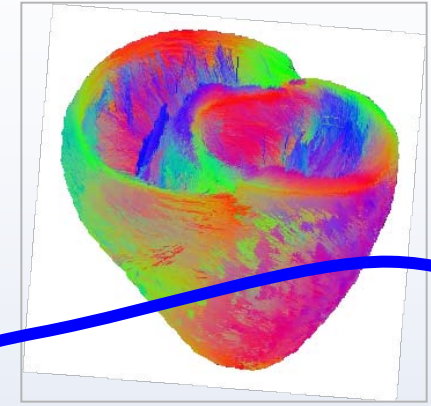
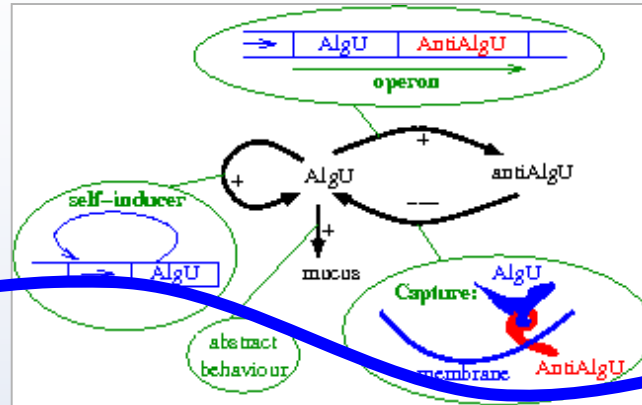
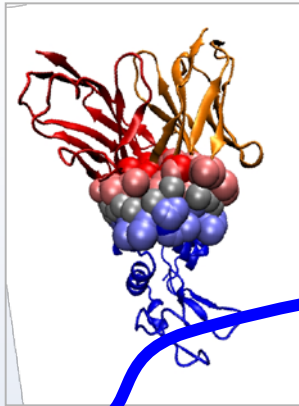
- From molecules to organs and the whole organism.

## Three **main topics**

- Bioinformatics
- Biomedical signal and image analysis
- Modeling in neuroscience



## Bioinformatics: Open problems



**Structure:** Protein complexes are ubiquitous:

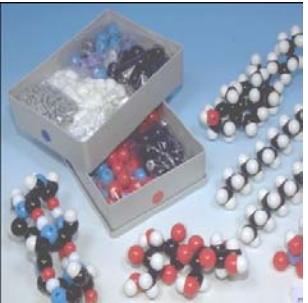
- Stability and specificity of macro-molecular complexes.
- Prediction ? (with little/no structural information).

**Networks,** systems biology:

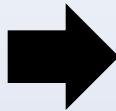

- Structure of interaction networks (topology).
- Associated dynamics (feedbacks and control loops).

# Bioinformatics: Methodology

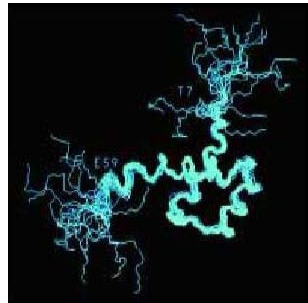
Biochemistry



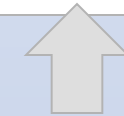
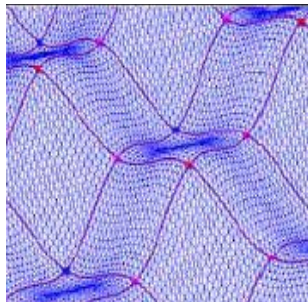
Biophysics



Geometry

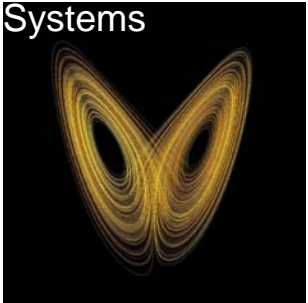


Topology

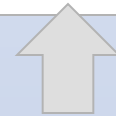
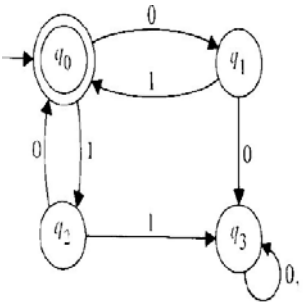


F. Cazals

Dynamical Systems

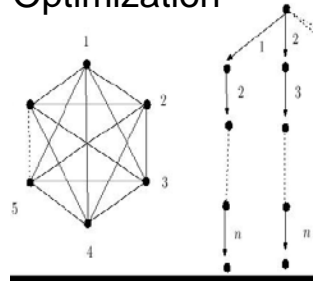


CS

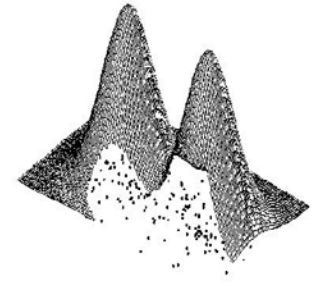


G. Bernot

Combinatorics  
Optimization

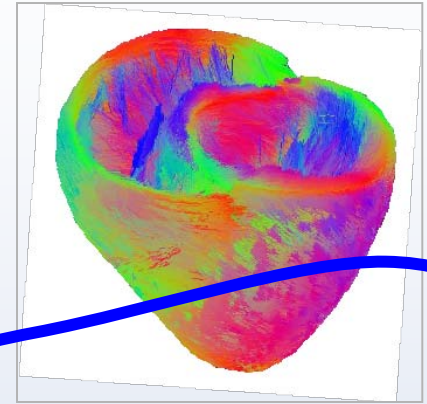
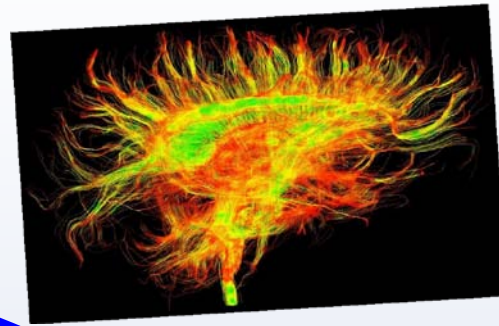
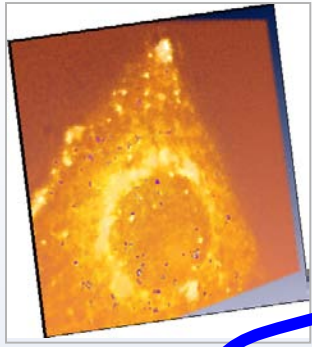


Statistics



Biophysical models  
and experimental  
data

## Biomedical signal and image analysis : Open problems



### Signal processing and inverse problems:

- Image denoising and enhancement.
- Inverse problems.
- Coupling to physical properties of sensors and tissues.

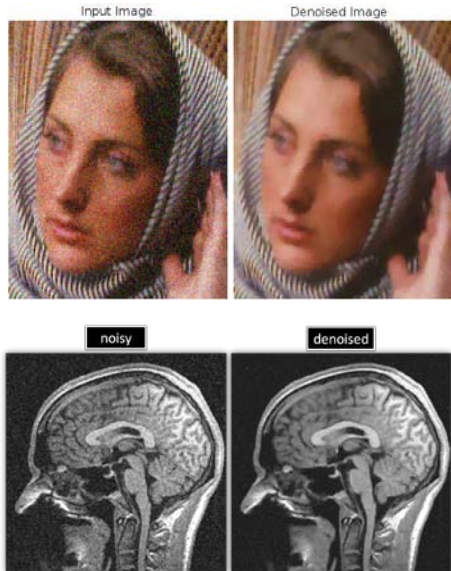
### Virtual human and patient specific modeling:

- Parameter identification.
- Statistical analysis (in shape spaces).
- Simulations.



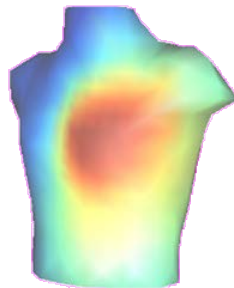
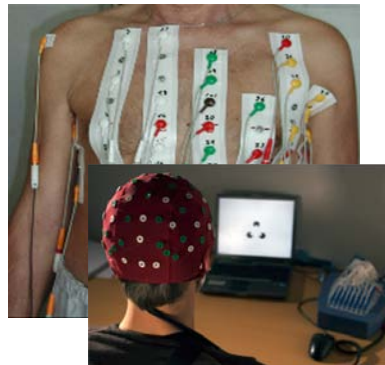
# Biomedical signal and image analysis : Methodology

Variational or Markovian models.  
Wavelets and



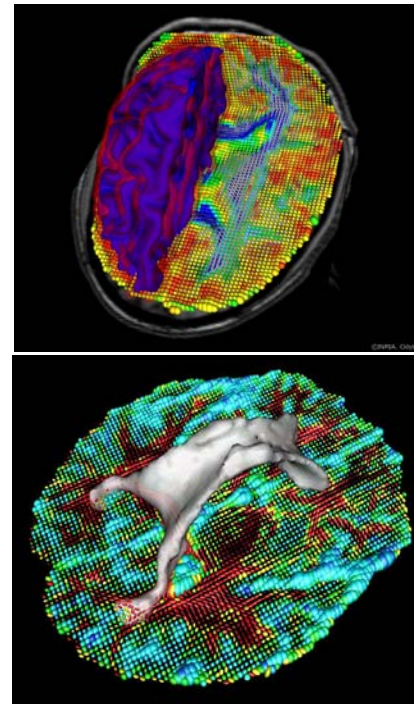
J. Zerubia

Inverse problems



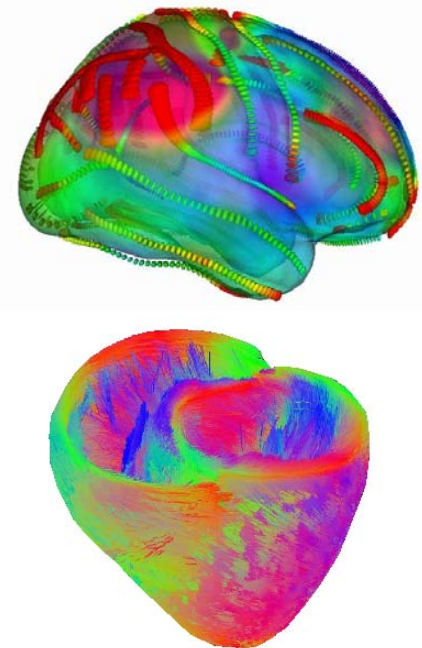
O. Meste, M. Clerc

Tensor calculus



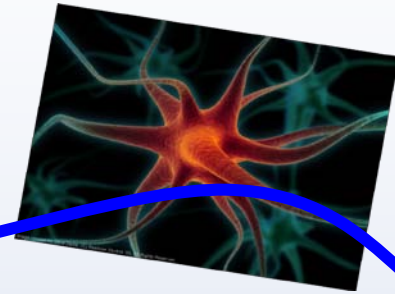
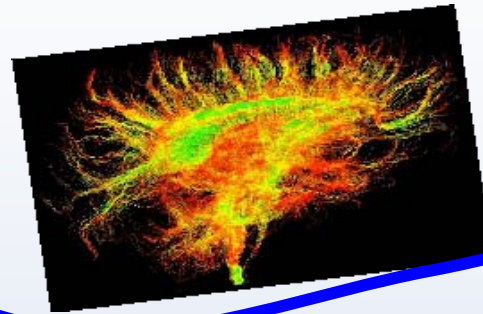
R. Deriche

Statistics



X. Pennec

## Modeling in neuroscience : Open problems

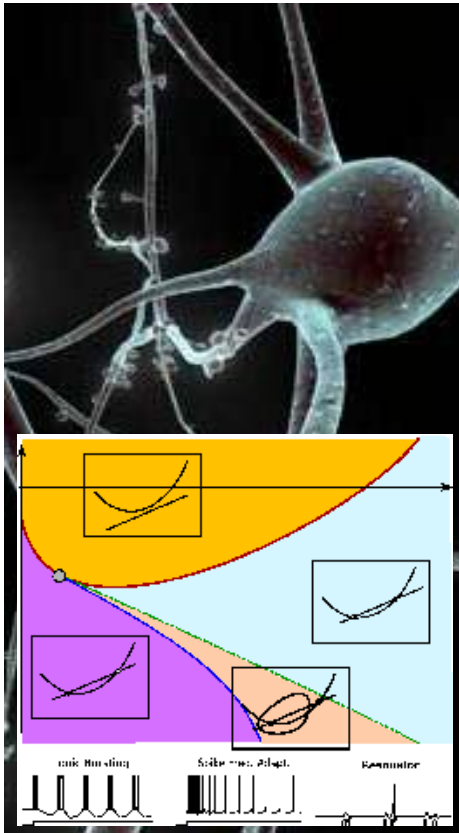


### A **multiscale** multidisciplinary problem

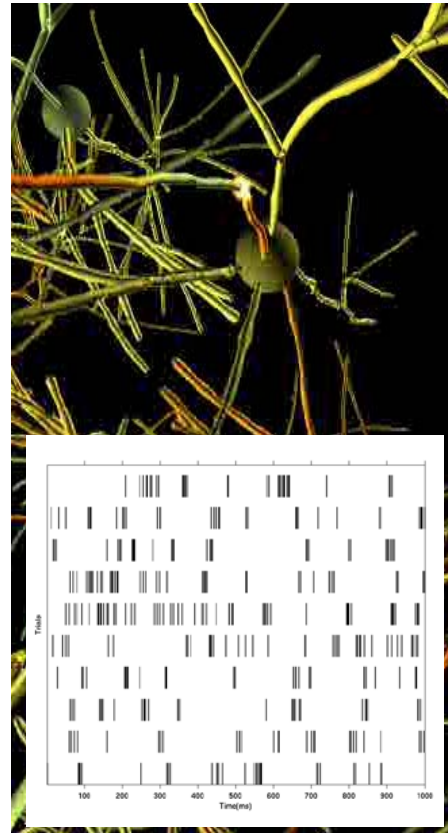
- Neurons and synapses: analysis of neuron dynamics
- Neuronal networks: Dynamical evolution. What about the statistics of spike trains?
- Neural masses: At a mesoscopic scale the neuronal substrate can be represented by a continuum where points represent neuronal populations.

# Modeling in neuroscience : Methodology

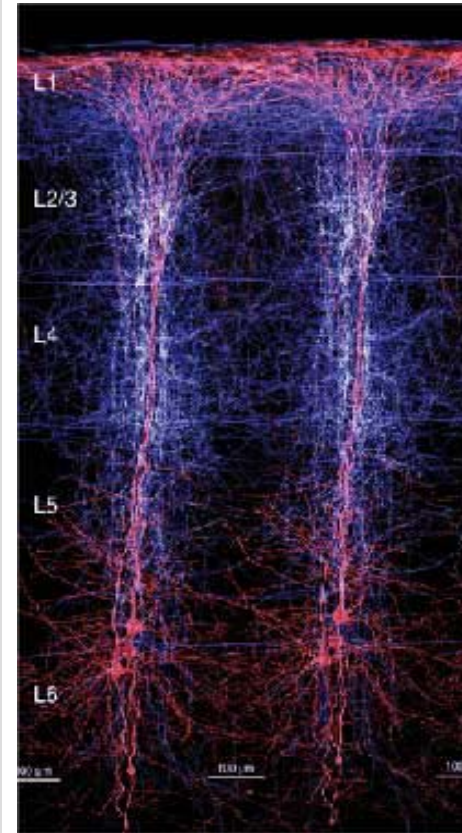
Dynamical systems  
(stability, bifurcations,  
asymptotic dynamics).



statistics



Integrodifferential  
equations



B. Cessac