



IP PARIS



# AMIB(io): Algorithms and Models for Integrative Biology

Discrete Bioinformatics for Molecular Biology  
RNA Structure, Design and Evolution

AMIBio team  
Ecole Polytechnique  
CNRS UMR 7161 LIX

Yann Ponty (DR2 CNRS)

# Members

Permanent staff:

Yann Ponty DR2  
(Head of AMIBio)



Philippe Chassagnet MCF



Hélène Thibault ITA



Sebastian Will PR



Jean-Marc Steyaert PR  
(Aemeritus)



Postdoc: Sarah Berkemer – 2021



PhD Students:

Hua-Ting Yao – 2018



Taher Yacoub – 2020



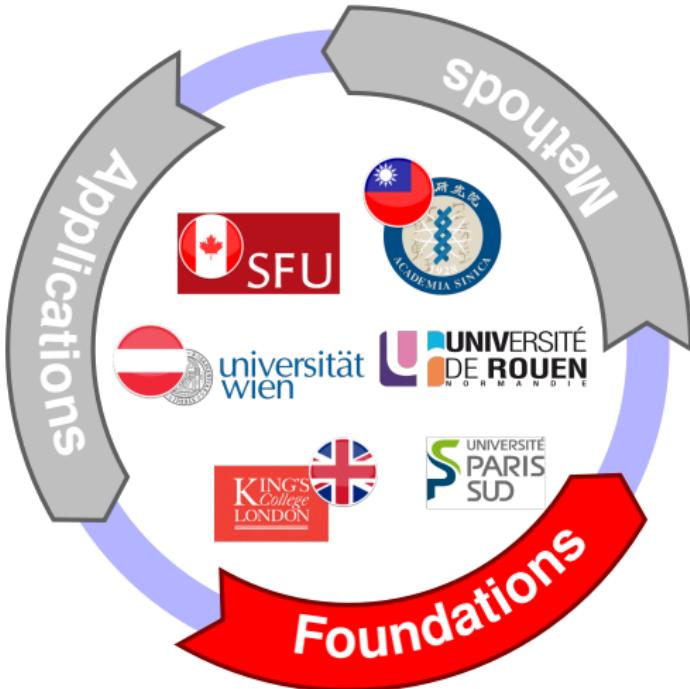
Bertrand Marchand – 2020



Ha Nguyen Ngoc – 2017

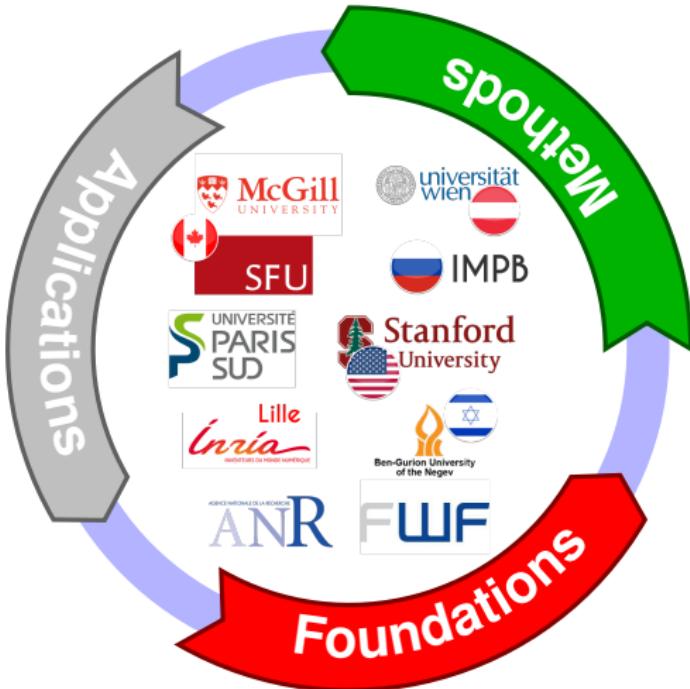


# In a nutshell



- ▶ (Algebraic) Dynamic Programming
- ▶ Enumerative/analytic combinatorics
- ▶ Stringology
- ▶ Random generation
- ▶ Machine learning
- ▶ Discrete optim./FPT algorithms
  
- ▶ Comparative genomics  
[J Math Biol'19]
- ▶ Statistical genomics
- ▶ Structural Bioinformatics  
[NAR'20, RECOMB'20, Bioinfo'21]
- ▶ RNA Design  
[RECOMB'18'21, BMC Bioinf'19]
  
- ▶ Circular RNAs [RNA Biology'17]
- ▶ HIV structure modeling [NAR'17]
- ▶ Structural basis of myristylation  
[Nature Chemistry'18]
- ▶ Chemical probing [NAR'17'20]

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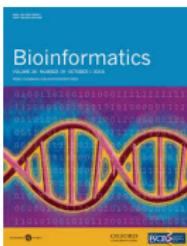
# Community and visibility



Program Committees



Organizers



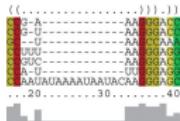
Editorial duties

# Main software

► LocaRNA/SPARSE

→ Univ. Freiburg

*De facto* standard for structural alignment of homologous RNAs



► VARNA

→ Univ. Paris Saclay

Visualization/modeling of RNA secondary structure(s)



► GenRGenS

→ Univ. Paris Saclay

Random generation toolkit for genomic sequences



► IPANEMAP

→ Univ. Paris Descartes

RNA structure prediction from chemical probing experiment(s)



► RNARedPrint/InfraRed

Declarative FPT framework for constrained sequence generation

# Current projects

- ▶ ANR/FWF PARNASSUS France/Austria → Univ. Descartes, Vienna
- ▶ ANR Decrypted → ENS Paris/Univ. Paris Descartes
- ▶ RNA design (PRC CNRS) → Univ Ben Gurion, Israel
- ▶ RNA modeling/geometric deep learning (CNRS?) → Univ McGill
- ▶ ArboRNA (DIM RFSI) → LIGM & Univ. Paris Descartes
- ▶ Reference-free cancer transcriptomics → Univ Paris Saclay
- ▶ RNA drug design for Alzheimer → Univ Paris Saclay
- ▶ SARS CoV-2 modeling → Univ. Paris Descartes & Univ. Wuhan

# Teaching activities

- ▶ Engineering/Bachelor@X
- ▶ Bioinformatics Modal/project@X
- ▶ AMI2B@Paris Saclay
- ▶ BIM@Sorbonne université

MERCI