



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)

Sebastian Will PR 

Philippe Chassignet MCF 

Jean-Marc Steyaert PR 
(Aemeritus)


Hélène Thibault ITA 

Postdoc: Sarah Berkemer – 2021 

PhD Students:

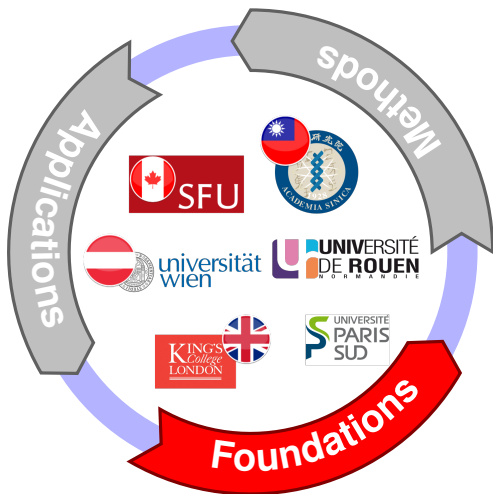
Hua-Ting Yao – 2018 

Bertrand Marchand – 2020 

Taher Yacoub – 2020 

Ha Nguyen Ngoc  – 2017 

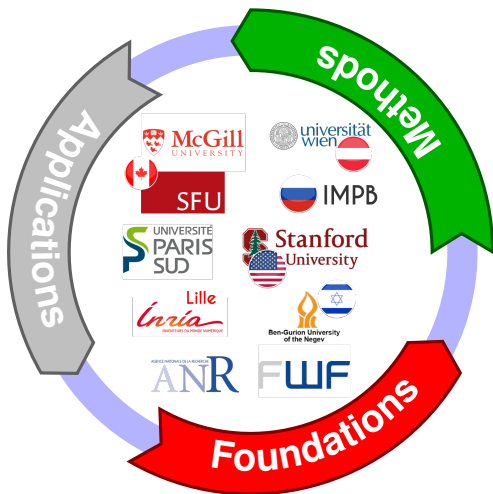
In a nutshell



- ▶ (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 myristoylation [Nature Chemistry'18]
- ▶ Chemical probing [NAR'17'20]

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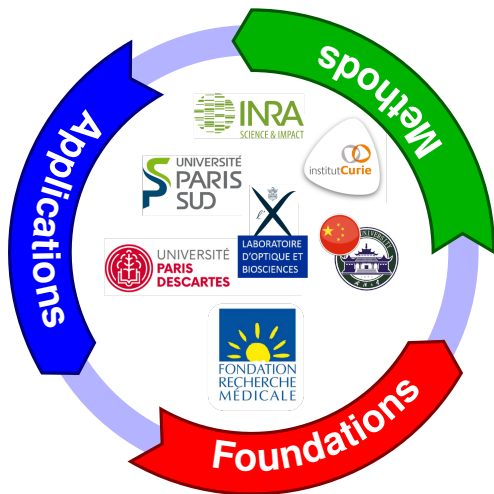


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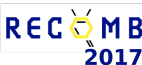
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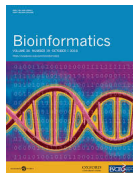
Community and visibility



Program Committees



Organizers



Editorial duties

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