# Philippe Desjardins-Proulx

CONTACT Graduate (Ph.D.) student

Department of Biology, Université du Québec à Montréal

Quebec Center for Biodiversity Science, McGill University, Canada.

Phone: +1-418-732-9877

E-mail: philippe.d.proulx@gmail.com WWW: http://phdp.github.com/ GitHub: https://github.com/PhDP

CITIZENSHIP Canada.

LANGUAGES French & English. Basic knowledge of Japanese.

Professional Interests Artificial Intelligence; Machine Learning; Artificial Neural Networks; Complexity; Bayesian Inference; Information theory; Biodiversity; Theoretical Population Genetics; Theoretical Ecosystem Ecology.

OTHER Interests Effective Technical Writing in English; Epistemology; Scientific Computing; Functional Programming; Open Source Software; Computational Finance.

Major Awards

From: Natural Sciences and Engineering Research Council of Canada

- Award: Alexander Graham Bell Graduate Scholarship, 09/2012–08/2015
- Value:  $105\ 000\$$  ( $\approx 105\ 000\ USD \approx 8\ 150\ 000\ JPY,\ 2012\ est.$ )

#### EDUCATION

### Department of Biology, Université du Québec à Montréal, Montréal, Canada.

Ph.D., September 2012 – August 2015 [expected]

- Thesis Proposal: Artificial Intelligence and the Puzzle of Genomic Diversity
- Adviser: Dr. Dominique Gravel
- Area of Study: Artificial Intelligence (machine learning: neural networks) and its applications to biodiversity.

### College of Engineering, University of Illinois at Chicago, Chicago, USA.

Graduate Certificate in Bioinformatics, 2012,

• Area of Study: Machine learning (Artificial Intelligence) and biostatistics.

## Université du Québec, Québec, Canada.

B.S., 2009,

- Major in Biology,
- Minor in Mathematics & Computer Science.

## REFEREED JOURNAL PUBLICATIONS

- [1] **P Desjardins-Proulx**, EP White, JJ Adamson, K Ram, T Poisot, and D Gravel. Developing a preprint culture in biology.

  Submitted to PLOS Biology (in review)
- [2] R Vergilino, TA Elliott, P Desjardins-Proulx, TJ Crease and F Dufresne. Evolution of a transposon in Daphnia hybrid genomes. Accepted in Mobile DNA

[3] D Ai, **P Desjardins-Proulx**, C Chu, and G Wang. The influence of immigration and dispersal limitation on the repeatability of niche and neutral communities. *PLOS ONE* 7(9): e46164, 2012.

DOI: 10.1371/journal.pone.0046164

- [4] P Desjardins-Proulx and D Gravel. A complex speciation-richness relationship in a simple neutral model. *Ecology and Evolution* 2(8): 1781–1790, 2012. DOI: 10.1002/ece3.292
- [5] P Desjardins-Proulx and D Gravel. How likely is speciation in neutral ecology? The American Naturalist 179(1):137-144, 2012.
   DOI: 10.1086/663196

# OTHER CONTRIBUTIONS

[6] P Desjardins-Proulx. The case for arXiv and a broader conception of peerreviews. Invited blog, International Network of Next-Generation Ecologists, 2012.

http://www.innge.net/?q=node/330.

- [7] **P Desjardins-Proulx**. A foot in the neutral trap. Invited comment for *Trends in Ecology & Evolution*, 2012.
- [8] P Desjardins-Proulx. L'origine de la Biodiversité. Le Mouton Noir, Mai-Juin. Cahier Spécial sur la Biodiversité p.2, 2010. Selected and republished by Gaia-Presse, a group sponsored by the Université Laval.

# Teaching & Training

### Université du Québec, Québec, Canada.

- 2013. I organized group studies of MacKay's Information Theory, Inference, and Learning Algorithms.
- 2012. CUDA training (intensive one-day course).
- 2012. Scientific computing with C (grad. students/post-docs).
- 2011. Scientific computing with C (grad. students/post-docs).

## Referee Service

Journal of Theoretical Biology, Theoretical Ecology, Acta Biotheoretica, Journal of Plant Ecology.

## COMPUTER SKILLS

- Programming languages:
  - Advanced: C++11, C.
  - Intermediate: JavaScript, Haskell, Java, Go, Python, R.
  - **Basic:** Ruby, MatLab/Octave.
- Tools & Frameworks
  - Operating systems: Linux (Debian/Ubuntu, Arch).
  - Writing: LaTeX  $2\varepsilon$ , LibreOffice.
  - Compilers: llvm/clang, gcc, intel, ghc.
  - Parallel computing: CUDA, OpenMP.
  - Linear algebra: Armadillo.
  - **Revision control:** git, mercurial.
  - **Database:** Redis, MongoDB.
  - Web: Node.js (Express).
- Web sites:
  - Personal page: http://phdp.github.com.
  - TEE's website: http://chaire-eec.ugar.ca.
- Primary working tools:
  - $\text{LAT}_{FX} 2_{\varepsilon}$ , vim, tmux.

Professional Memberships	<ul> <li>Institute of Electrical and Electronics Engineers</li> <li>International Society for Computational Biology</li> <li>Society for the Study of Evolution</li> <li>Quebec Center for Biodiversity Science</li> </ul>	2012 2010 2008 2012
GRADUATE COURSES	<ul> <li>2012. Datamining (machine learning) [A (4.0/4.0), 4 credits]</li> <li>2011. Biostatistics [A (4.0/4.0), 4 credits]</li> <li>2010. Intro. to bioinformatics [A (4.0/4.0), 4 credits]</li> <li>2010. Reading course on Ancestral Recombination Graphs [A, 3 credits]</li> </ul>	UIC UIC UIC UQAR

#### Referees

## Dr. Dominique Gravel

- Professor (Université du Québec à Rimouski)
- Canada Research Chair.
- ullet e-mail: dominique\_gravel@uqar.qc.ca
- phone: 1.418.723.1986 #1752
- \* I worked as a research profesionnal in Dr. Gravel's lab from September 2009 to August 2012, we also collaborated on many scientific projects. I am a Ph.D. student in his lab since September 2012.

## Dr. James Rosindell

- Post-doctoral researcher, Imperial College London, UK.
- e-mail: j.rosindell@imperial.ac.uk
- phone: +44 (0)2075 942263
- $\star$  I have collaborated with Dr. Rosindell on several occasions.