

# Philippe Desjardins-Proulx

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CURRENT POSITION	Ph.D. candidate, Université du Québec à Montréal, Canada, Quebec Center for Biodiversity Science, McGill University, Canada.
CONTACT	<i>email:</i> <a href="mailto:phdp@outlook.com">phdp@outlook.com</a> <i>email (alt.):</i> <a href="mailto:philippe.d.proulx@gmail.com">philippe.d.proulx@gmail.com</a> <i>phone:</i> +1-418-732-9877 <i>skype:</i> philippe.desjardins-proulx <i>www:</i> <a href="http://phdp.github.io/">http://phdp.github.io/</a> <i>github:</i> <a href="https://github.com/phdp/">https://github.com/phdp/</a>
CITIZENSHIP	Canada
LANGUAGES	FRENCH: Native language ENGLISH: Full professional proficiency JAPANESE: Elementary proficiency
EXPERTISE	<ul style="list-style-type: none"><li>• <b>Machine Learning &amp; Natural Language Processing:</b> My thesis focuses on the problem of deep transfer learning, which could be described as the capacity to automatically transfer knowledge between a source (the knowledge we want to transfer) and a target (the model we want to build). It comes naturally to us, e.g.: we use what we learned for walking when learning to run, but almost all machine learning algorithms build models from nothing. The current deep transfer learning algorithms based on Markov logic achieve poor performance in part because they don't know enough about the context to judge what knowledge is likely to be transferable. My plan is to use tools from natural language processing to build semantic networks capable of providing enough context to perform smarter mappings between the source and the target.</li><li>• <b>Scientific computing:</b> Almost all my undergraduate projects were done on scientific computing in biology and I worked 4 years (2009-2012) as a research professional, focusing on C/C++/CUDA simulations on the Canada Research Chair on Terrestrial Ecosystems' scientific cluster. I have implemented several complex algorithms, including high quality random number generators and evolutionary algorithms. I'm particularly interested in the design and deployment of machine learning and artificial intelligence algorithms on distributed systems.</li><li>• <b>Ecology:</b> I'm interested in complexity and nothing is more complex than ecosystems. I have contributed to theoretical ecology, in particular on the problem of speciation (the origin of species) and how it relates to the origin of ecological communities.</li></ul>
EDUCATION	<b>Department of Biology, Université du Québec à Montréal, Montréal, Canada.</b> Ph.D., September 2012 – <ul style="list-style-type: none"><li>• <b>Thesis Proposal:</b> <i>Achieving deep transfer learning with semantic networks, with applications to species distribution modeling</i></li><li>• <b>Adviser:</b> <a href="#">Dr. Dominique Gravel</a></li><li>• <b>Area of Study:</b> Machine learning; Artificial Intelligence; Natural Language Processing; Probabilistic Graphical Models; Statistical learning; Learning transfer; Ecology.</li></ul>

- **Courses:** Advanced Distributed Computing (A13).

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

Graduate Certificate in Bioinformatics, 2012,

- **Area of Study:** Data Mining & biostatistics.

**Université du Québec, Québec, Canada.**

B.S., 2009,

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

**AWARDS**

Alexander Graham Bell Graduate Scholarship (2012)

- **From:** Natural Sciences and Engineering Research Council of Canada
- **Description:** Most competitive Canadian scholarship in science.
- **Value:** 105 000\$ (equivalent to 105 000 USD or 8 150 000 JPY, 2012 est.)

Windows Azure Research Award (2013)

- **From:** Microsoft Research
- **Description:** Access to the Windows Azure cloud for selected science proposals.
- **Proposal:** *Growing Intelligence with Cloud Markov Logic.*
- **Value:** >40 000 USD.

**REFEREED  
JOURNAL  
PUBLICATIONS**

- [1] **P Desjardins-Proulx**, EP White, JJ Adamson, K Ram, T Poisot, and D Gravel. The case for open preprints in biology. *PLoS Biology* 11(5): e1001563
- [2] R Vergilino, TA Elliott, **P Desjardins-Proulx**, TJ Crease and F Dufresne. Evolution of a transposon in *Daphnia* hybrid genomes. *Mobile DNA* 4-7, 2013. DOI: 10.1186/1759-8753-4-7
- [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 peer-reviews. Invited blog, International Network of Next-Generation Ecologists, 2012. <http://www.innge.net/?q=node/330>.
- [7] **P Desjardins-Proulx**, JL Rosindell, T Poisot, and D Gravel. A simple model to study phylogeographies and speciation patterns in space, 2012. arXiv: 1203.1790.
- [8] **P Desjardins-Proulx**. A foot in the neutral trap. Invited comment for *Trends in Ecology & Evolution*, 2012.

- [9] **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.*

JOB EXPERIENCES	<b>Research Professional, Canada Research Chair on Terrestrial Ecosystem</b> <ul style="list-style-type: none"> <li>• From 2009 to 2012.</li> <li>• <b>Supervisor:</b> <a href="#">Dr. Dominique Gravel</a></li> <li>• <b>Responsabilities:</b> Programming high-performance simulations in C, C++, and CUDA on a distributed cluster; Design of ecological models to understand biodiversity; Teaching scientific computing to graduate students.</li> </ul>	
TEACHING EXPERIENCES	<b>Université du Québec, Québec, Canada.</b> <ul style="list-style-type: none"> <li>• 2013. I organized a series of meetings on information theory and inference.</li> <li>• 2012. CUDA training (intensive one-day course).</li> <li>• 2012. Scientific computing with C and C++ (grad. students/post-docs).</li> <li>• 2011. Scientific computing with C and C++ (grad. students/post-docs).</li> </ul>	
REFeree SERVICE	<i>Molecular Ecology Ressources; Journal of Theoretical Biology, Theoretical Ecology, Acta Biotheoretica, Journal of Plant Ecology.</i>	
COMPUTER SKILLS	A few of my favorite tools: <ul style="list-style-type: none"> <li>• <b>Programming languages:</b> <ul style="list-style-type: none"> <li>– <b>Advanced:</b> C, C++11, Java.</li> <li>– <b>Intermediate:</b> Clojure, F#, Haskell, Python.</li> </ul> </li> <li>• <b>Other skills:</b> <ul style="list-style-type: none"> <li>– <b>Operating systems:</b> Linux (primary), Windows.</li> <li>– <b>Database:</b> SQL, Neo4J, PostgreSQL, MongoDB, Redis.</li> <li>– <b>Cloud services:</b> Windows Azure, Heroku, Google App Engine.</li> </ul> </li> </ul>	
PROFESSIONAL MEMBERSHIPS	<ul style="list-style-type: none"> <li>• Institute of Electrical and Electronics Engineers</li> <li>• Quebec Center for Biodiversity Science</li> </ul>	2012–... 2012–...
GRADUATE COURSES	<ul style="list-style-type: none"> <li>• 2013. Advanced Distributed Computing [3 credits]</li> <li>• 2012. Datamining (machine learning) [4 credits]</li> <li>• 2011. Biostatistics [4 credits]</li> <li>• 2010. Intro. to bioinformatics [4 credits]</li> <li>• 2010. Reading course on Ancestral Recombination Graphs [3 credits]</li> </ul>	Athabasca UIC UIC UIC UQAR