

Patrik D'haeseleer

Department of Genetics, Harvard Medical School
The New Research Building, 77 Avenue Louis Pasteur, Boston, MA 02115
Phone: 617-432-6511; Fax: 617-432-7266
E-mail: patrik@genetics.med.harvard.edu
<http://genetics.med.harvard.edu/~patrik>

Education

- 2000 **Ph.D. in Computer Science.**
University of New Mexico, Albuquerque, NM.
- 1993 **M.S. in Computer Science** (Artificial Intelligence), GPA: 4.0.
Stanford University, Stanford, CA
- 1991 **M.S. (“Burgerlijk Ingenieur”) in Electrical Engineering** (electronics), with distinction.
Ghent University, Ghent, Belgium.
- 1988 **M.E. (“Industrieel Ingenieur”) in Electrical Engineering**, with distinction.
Katholieke Industriële Hogeschool Oost-Vlaanderen, Ghent, Belgium.

Research and Professional Experience

- 2001-present Postdoctoral Fellow, Harvard Medical School. Study of complex networks in biology: regulatory motifs and expression patterns; reconstruction of regulatory and metabolic networks; tracing signaling pathways through the protein interaction network; protein interaction error rates and analysis of weighted interaction networks. Advisor: Prof. George Church.
- 1996-2000 Research Assistant, University of New Mexico. Ph.D. thesis work on inference of genetic regulatory networks from large-scale gene expression data. Advisor: Prof. Stephanie Forrest.
- 1995-1996 Research Assistant, Santa Fe Institute. Member of the EvCA (Evolving Cellular Automata) group. Investigated evolution of emergent computation in 2D cellular automata, in collaboration with Melanie Mitchell and Jim Crutchfield.
- 1995 Complex Systems Summer School. Santa Fe Institute, NM.
- 1994-1995 Research Assistant, University of New Mexico. Investigated the mathematical and algorithmic foundations of an artificial immune system.
- 1993-1994 Research Scientist, LSI Logic, Mountain View, CA. Development of novel methods for cell placement in integrated circuit design (based on evolutionary algorithms), resulting in several patents.
- 1990-1991 Thesis research, Ghent University. Implemented an artificial neural network for optical character recognition.
- 1987-1988 Thesis research, Katholieke Industriële Hogeschool Oost-Vlaanderen. Designed and implemented a system for capturing and digitizing video frames on a PC.

Teaching Experience

- 2001-2002 Organized and ran a bi-weekly Networks Discussion Group, Harvard Medical School.
- 2001 & 2002 Instructor, Instituto Gulbenkian de Ciência, Portugal. Co-taught a one-week intensive Introduction to Bioinformatics for the PhD program in Biomedicine.
- 1992-1993 Teaching Assistant, Stanford University. CS140: Concurrent Programming (three times) and CS221: Introduction to Artificial Intelligence (twice).

Awards

- 1996 Outstanding graduate student award, UNM Computer Science Department.
1995 First student to pass UNM CS comprehensive exams with distinction in 5 years.
1991 Honorary fellowship, Belgian-American Education Foundation.
1991 Graduated Ghent University with distinction.
1988 Graduated Katholieke Industriële Hogeschool Oost-Vlaanderen with distinction.

Invited Lectures

- June 2003 Southern Regional Council on Statistics, Summer Research Conference in Statistics.
March 2003 Harvard Center for Genomics Research.
June 2001 Boston University, Center for BioDynamics.
April 2000 University of New Mexico, Department of Biology.
February 2000 University of New Mexico, Department of Neurosciences.
December 1999 Los Alamos National Labs, Theoretical Biology and Biophysics group.
October 1999 Short course on microarrays, Society for Neuroscience Annual Meeting.
June 1999 National Center For Genome Resources, Santa Fe, NM.
May 1999 Fred Hutchinson Cancer Research Center/Santa Fe Institute Exchange Meeting: Melding Theoretical and Experimental Biology.
January 1999 Tutorial on Gene Expression and Genetic Networks, Pac. Symp. Biocomputing, Hawaii, HI.

Selected Publications

D'haeseleer, P. and Church, G.M. (2003) Estimating and improving protein interaction error rates. *Submitted to RECOMB.*

Segrè, D., Zucker, J., Katz, J., Lin, X., **D'haeseleer, P.**, Rindone, W., Karchenko, P., Nguyen, D., Wright, M., and Church, G.M. (2003) From annotated genomes to metabolic flux models and kinetic parameter fitting. *Omics* 7:301-16

Steffen, M., Petti, A., **D'haeseleer, P.**, Aach, J., and Church, G.M. (2002) Automated Modeling of Signal Transduction Networks. *BMC Bioinformatics* 3:34-44.

Fuhrman, S., **D'haeseleer, P.** and Somogyi, R. (2002) Tracing Genetic Information Flow from Gene Expression to Pathways and Regulatory Networks. In *Microarrays for the Neurosciences: An Essential Guide*, eds. Geschwind D.H. and Gregg J. (The MIT Press).

Chao, D.L. and **D'haeseleer P.** (2001) The Distribution of Variable-length Phatic Interjectives on the World Wide Web. Technical Report TR-CS-2001-23, The University of New Mexico.

D'haeseleer, P. (2000) Reconstructing Gene Networks from Large Scale Gene Expression Data. Ph.D. dissertation, University of New Mexico.

D'haeseleer, P., Liang, S., and Somogyi, R. (2000) Genetic Network Inference: From Co-Expression Clustering to Reverse Engineering. *Bioinformatics* 16(8):707-26.

D'haeseleer, P., Wen, X., Fuhrman, S., and Somogyi, R. (1999) Linear modeling of mRNA expression levels during CNS development and injury. *Pacific Symposium on Biocomputing* (World Scientific Publishing), pp. 41-52.

D'haeseleer, P., Wen, X., Fuhrman, S., and Somogyi, R. (1998) Mining the gene expression matrix: Inferring gene relationships from large scale gene expression data. In *Information Processing in Cells and Tissues*, eds. Paton, R.C. and Holcombe, M. (Plenum Publishing), pp. 203-212.

D'haeseleer, P. (1996) An Immunological Approach to Change Detection: Theoretical Results. *Proceedings of the 9th IEEE Computer Security Foundations Workshop* (IEEE Computer Society Press), pp. 18-26.

D'haeseleer, P., Forrest, S. and Helman, P. (1996) An Immunological Approach to Change Detection: Algorithms, Analysis and Implications. *Proceedings of the IEEE Symposium on Security and Privacy* (IEEE Computer Society Press), pp. 110-119.

D'haeseleer, P. and Bluming, J. (1994) Effects of Locality in Individual and Population Evolution. In *Advances in Genetic Programming*, K.E. Kinnear Jr. Ed. (MIT Press), pp.177-198.

D'haeseleer, P. (1994) Context Preserving Crossover in Genetic Programming. *Proceedings of 1st IEEE Conf. on Evolutionary Computation* (IEEE Press), pp. 256-261.

D'haeseleer, P. and De Backer, W. (1991) Neural Network for Optical Character Recognition (“*Neuraal Netwerk voor Schriftherkenning*”). Thesis work Ghent University, Ghent Belgium.

D'haeseleer, P. and Moreau, P. (1988) Digitizing and Processing of Video Images (“*Digitaliseren en Verwerken van Video Beelden*”). Thesis work K.I.H.O., Ghent Belgium.

Patents

D'haeseleer, P. and Scepanovic, R. (1999) Method and apparatus for computing minimum wirelength position (MWP) for cell in cell placement for integrated circuit chip. United States Patent 5,859,781.

D'haeseleer, P. and Boyle, D. B. (1997) Method of cell placement for an integrated circuit chip comprising integrated placement and cell overlap removal. United States Patent 5,619,419.

Scepanovic, R. and **D'haeseleer, P.** (1997) Optimal pad location method for microelectronic circuit cell placement. United States Patent 5,638,293.

References available upon request