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## **Yannick Wurm**

5.03A Fogg Building Organismal Biology Queen Mary University of London London E1 4NS, UK

I aim to understand the genetic mechanisms involved in the evolution of animal societies. I use genomics and bioinformatics to characterize genes involved in behavioral decisions and caste-specific developmental fates in ants and create novel bioinformatics tools for genomics work with emerging model organisms.

## **HISTORY**

2014-		SENIOR LECTURER IN BIOINFORMATICS, Queen Mary University of London (QMUL)			
2012-2	2014	LECTURER IN BIOINFORMATICS, Queen Mary University of London (QMUL)			
2010-2	2011	1 Postdoctoc with L. Keller, Dept. Ecology & Evolution, U Lausanne &			
		EMBEDDED BIOINFORMATICIAN with I. Xenarios, Vital-IT, Swiss Institute of Bioinformatics (SIB)			
2006-2	2009	Member of SIB PhD Training Network			
2005-2	2009	PHD: SOCIOGENETICS OF FIRE ANTS with L. Keller, Dept. Ecology & Evolution, U Lausanne			
2004		RESEARCH TRAINEE (6 MONTHS) with R. Roy, Biology Dept., McGill, Montreal			
2003		JUNIOR SOFTWARE DEVELOPER (6 MONTHS) with N. Zinovieff, Krugazor Software, Paris			
1999-2	2004	BIOINFORMATICS & MODELING (MASTERS); Natl. Institute of Applied Sciences, INSA Lyon			
1999		SCIENTIFIC BACCALAUREATE (BIOLOGY & MATH) Cité Scolaire Internationale de Lyon			

France (10y; citizen); Germany (4y; citizen), USA (9y), Switzerland (7y), Indonesia (1 y), UK (since 2012).

## **DISTINCTIONS & GRANTS**

	ONS & GIVANTS				
2014	VP-research fund (Wurm et al (with 12 others) 45,000£)				
2015-2017					
2014-2017					
2014-2017	- 0 1				
2014-15	NERC Big Data Capital Call (local co-I with R. Nichols; 85,000£ to QMUL)				
2014	Queen Mary Faculty of Science & Engineering Research Performance Award.				
2013	Google Summer of Code (GSoC) with A. Priyam and MA. Dragan (\$5,500)				
2013-14	NEScent Working Group. Building non-model species genome curation communities				
	(with A. Papanicolaou & M. Muños-Torres) (approx. \$50,000 total).				
2012-14	BBSRC sTDRF. Genome Annotation for the Masses. (148,136£ Full economic cost)				
2011	NERC BioMolecular Analysis Facility proposal NBAF627 with RL. Hammond (5,656£)				
2011	Society for Molecular Biology & Evolution Postdoc travel award (\$2,500)				
2010	PhD award: Prix de la faculté de l'Université de Lausanne (1,000 CHF)				
2009	Infectigen: Ultra-high throughput sequencing for discovery & characterization of viral				
	genomes in fire ants. With E. Zdobnov, L. Keller, D. Shoemaker, S. Valles (100,000 CHF)				
2008	Swiss Zoological Society & Fondation du 450 grants to attend				
	International Congress of Entomology (3,500 CHF)				
2008	Ecogenics innovation award for oral communication at Biology08 (400 CHF)				
2004	Région Rhône-Alpes and Office Franco-Québécois de la Jeunesse				
	to visit Prof. Richard Roy's laboratory at McGill (2,800€)				
2002	Apple WorldWide Developer Conference Scholarship (San Jose; \$1,600)				
1999	Scientific baccalaureate honors: "Mention Bien"				
-555	Colonial Dacoular Care Honoror Mention Dien				

MEDIA COVERAGE Has been extensive - not least because social insects are particularly media-friendly. See http://yannick.poulet.org/pressCoverage/

**REVIEWER** for BBSRC, Research Foundation Flanders, Nature Communications, Proc B, Genome Biol, Mol Ecol, Mol Biol Evol, Bioinformatics, BMC Evol Biol, BMC Genomics, Insect Mol Biol, Nucl Acids Res, Bull Insectol, Insectes Sociaux, PLoS One, Curr Zool. iConcept Press.

## **ADDITIONAL RESPONSIBILTIES**

Current: **Supervision/line management**: postdoc (E Stolle); full PhD students (R Pracana, B Vieira, L Santiago), shared PhD student (N Arajo); Sysadmin (S Moss), software developer (A Priyam), Five 3rd year project students; **PhD panel** of: S Dodsworth, S Bailey, L Sollars, J Zohren, D Radcliffe, S Saha. **Advisor** to ~30 undergraduate students. Chair of QMUL School of Biology web committee; **Member**, QMUL High Performance Computing steering group; **Teaching** (Year 1 Evolution, Year 3 Ecological & Evolutionary Genomics, Year 3 Research Methods, MSc Bioinformatics for genomics). **Leading** http://antgenomes.org and http://sequenceserver.com.

Previously: 3 Bioinformaticians: (O Riba-Grognuz, F Ricci, M Bota, B Vieira), 3 Lab technicians (C LaMendola, C Stoffel, N Tzaud), 3 Masters students (S Buechel, P Siguier, H Darras).

INVITED O	RAL PRESENTATIONS	2012/01/20	European Course on Comparative
2014/12/11	U Helsinki, Finland	001110-101	Genomics, ENS Lyon
2014/12/8	Oulu, Finland	2011/07/31	Laboratory of Insect Ecology, Okayama
2014/05/28	Balti & Bioinformatics @ Birmingham U	2011/06/29	International Social Insect Genomics Research Conference, Shenzhen
2013/11/25	Sheffield University	2011/05/06	
2013/09/05	Cream Teas & Bioinformatics, @ Exeter	2011/05/06	Communication in Social Insects, National Taiwan University, Taipei
2013/04/11	Sussex University	2011/04/11	Laboratoire de Biométrie et
2013/03/19	Bristol University	2011/01/11	Biologie Évolutive, Lyon
2013/03/15	Institut Jacques Monod, Paris	2011/03/31	Les Doctoriales, Lausanne
2013/03/13	UCL Department of Genetics,	2011/03/29	UCL Department of Genetics
	Evolution and Environment		Ecology & Environment, London
2013/01/17	Zoologisches Kolloquium, U Regensburg	2010/10/31	Jakarta Ruby Brigade
2012/11/05	UCL MRes (Biodiversity,		meeting, Indonesia
	Evolution and Conservation)	2010/08/31	UK RAD Sequencingmeeting, Edinburgh
2012/09/06	European Conference of Apidologie	2009/01/15	The Past, Present and Future of Ant
	(EurBee 5), Halle an der Saale		Genomics, Arizona State University
2012/06/18	SIB workshop on Evolutionary &	2008/12/01	Interaction Seminar, Zurich
	Ecological Genomics, Adelboden	2008/10/23	Symposium of Ecology and Evolution
<b>PUBLICATI</b>	ONS ("corresponding authorship: "equal cor	Doctoral Students, Lausanne	

PUBLICATIONS (°corresponding authorship; \*equal contributions)

2014 Nipitwattanaphon, Wang, Ross, Riba-Grognuz, Wurm, Khurewathanakul, Keller. Effects of ploidy and sex-locus genotype on gene expression patterns in the fire ant Solenopsis invicta. Proc B 281: 20141776

- 2014 Purcell, Brelsford, <u>Wurm</u>, Perrin, Chapuisat. **Convergent genetic architecture underlies social organization in ants** Current Biology.
- 2014Buechel\*, <u>Wurm</u>\*, Keller L. Social chromosome variants differentially affect queen determination and the survival of workers in the fire ant *Solenopsis invicta*. Molecular Ecology 23:5117-27
- 2013 Valles°, Shoemaker, <u>Wurm</u>, Strong, Varone, Becnel, Shirk. **Discovery & molecular characterization of an ambisense densovirus from South American populations of** *Solenopsis invicta***. Biological Control 67:431-39.**
- 2013 Corona, Libbrecht, <u>Wurm</u>, Riba-Grognuz, Studer, Keller°. **Vitellogenin underwent subfunctionalization to acquire caste and behavioral specific expression in the harvester ant** *Pogonomyrmex barbatus***. PLoS Genetics 9(8):e1003730.**
- 2013 Manfredini, Riba-Grognuz, <u>Wurm</u>, Keller, Shoemaker, Grozinger°. **Sociogenomics of cooperation and conflict during colony foundation in the fire ant** *Solenopsis invicta*. PLoS Genetics 9:e1003633.
- 2013 38 authors (incl. Wurm Y). Social insect genomes exhibit dramatic evolution in gene composition and regulation while preserving regulatory features linked to sociality. Genome Research gr.155408.113.
- 2013 Wang\*°, <u>Wurm\*</u>°, Nipitwattanaphon, Riba-Grognuz, Huang, Shoemaker, and Keller°. **A Y-like social chromosome causes alternative social organization in fire ants.** Nature 493: 664-68.
- 2013 Privman, <u>Wurm</u>°, Keller° **Duplication and concerted evolution in a master sex determiner under balancing selection.** Proceedings of the Royal Society B.
- 2013 Kulmuni°, <u>Wurm</u>, Pamilo. Comparative genomics of chemosensory protein genes reveals rapid evolution and positive selection in ant-specific duplicates. Heredity.
- 2012 Ingram°, Kutowoi, <u>Wurm,</u> Shoemaker, Meier, Bloch. **The molecular clockwork of the fire ant** *Solenopsis invicta*. PLoS One 7(11):e45715.
- 2012 Chittka, Wurm, Chittka. Epigenetics: The Making of Ant Castes. Current Biology 22(19):R835-8.
- 2012 Gadau°, Helmkampf, Nygaard, Roux, Simola, Smith, Suen, <u>Wurm</u>, Smith. **The genomic impact of 100** million years of social evolution in seven ant species. Trends in Genetics 28:14-21 [cover photo].
- 2011 Riba-Grognuz°, Keller, Falquet, Xenarios, <u>Wurm°</u>. **Visualization and quality assessment of** *de novo* **genome assemblies.** Bioinformatics, 27(24): 3425-26.
- 2011 Hunt, Ometto, <u>Wurm</u>, Shoemaker, Yi, Keller, Goodisman°. **Relaxed selection is a precursor to the evoluton of phenotypic plasticity.** PNAS 108(38):15936-41
- 2011 Nygaard°, Zhang, Schiøtt, Li, <u>Wurm</u>, and 11 additional authors including Boomsma). **The genome of the leaf-cutting ant** *Acromyrmex echinatior* **suggests key adaptations to advanced social life and fungus farming.** Genome Research 21:1339-48.
- 2011 Wurm Y°. Behind the scenes of an ant genome project. Formosan Entomologist, 31(2):149-56.
- 2011 Wurm Y° et al (38 authors). The Genome of the fire ant Solenopsis invicta. PNAS, 108(14):5679-84.
- 2011 Gotzek, Robertson, <u>Wurm</u>, Shoemaker. **Odorant binding proteins of the Red Imported Fire Ant**, *Solenopsis invicta*: **An example of the problems facing the analysis of widely divergent proteins.** PLoS ONE 6: e16289.
- 2010 Wurm°, Keller. Parasitoid Wasps: From natural history to genomic studies. Current Biology 20:R242-4.
- 2010 Wurm°, Wang, Keller. Changes in reproductive roles are associated with changes in gene expression in fire ant queens. Molecular Ecology 19(6).
- 2009 <u>Wurm</u>°, Uva, Ricci, Wang, Jemielity, Iseli, Falquet, Keller. **Fourmidable: a database for ant genomics.** BMC Genomics 10(5):5.
- 2007 Wang, Jemielity, Uva, <u>Wurm</u>, Gräff, Keller. **An annotated cDNA library and microarray for large-scale gene-expression studies in the ant** *Solenopsis invicta***. Genome Biology 8:R9.**
- 2007 Wurm, Wang, Keller. Behavioral genomics: A, bee, C, G, T. Current Biology 17:R51-3.
- 2005 Zerlin M, Greiffenberg, <u>Wurm Y</u>, Winkler I, Kreusel D, Toemoe S, Ballabas L, Martin J. **Cryostock, Software for Cell Culture Management**. ESACT Proceedings: Animal Cell Technology Meets Genomics. 397-400.