

Education

- ★ Graduate Student (Bioinformatics) April 2006 - Present
UCD Complex & Adaptive Systems Laboratory and UCD Conway Institute
University College Dublin, Ireland
- ★ Bachelor of Technology (Electrical Engineering) 2001
Indian Institute of Technology, Kanpur, India

Professional Experience

- ★ Bioinformatics Research Fellow Jul. 2003 - Mar. 2006
University of Miguel Hernández, Alicante, Spain
- ★ Bioinformatics Software Developer Jan. 2002 - Jun. 2003
Institute of Genomics & Integrative Biology, Delhi, India
- ★ Bioinformatics Software Consultant Dec. 2001 - Jan. 2002
Institute of Genomics & Integrative Biology, Delhi, India

Publications

- ★ Mira A., **Pushker R.** and Rodríguez-Valera F. (2006) The Neolithic revolution of bacterial genomes. *Trends in Microbiology*, 14(5):200-206.
- ★ D'Auria G., **Pushker R.** and Rodríguez-Valera F. (2006) IWoCS: analyzing ribosomal Intergenic Transcribed Spacer configuration and taxonomic relationships. *Bioinformatics*, 22(5):527-531.
- ★ Zaballos M., López-López A., Ovreas L., Bartual S. G., D'Auria G., Legault B., Alba J. C., **Pushker R.**, Dae F. L. and Rodríguez-Valera F. (2006) Comparison of prokaryotic diversity at two offshore oceanic locations reveals a different microbiota in the Mediterranean Sea. *FEMS Microbiology Ecology*, 56(3):389-405.
- ★ **Pushker R.**, D'Auria G., Alba-Casado J.C. and Rodríguez-Valera F. (2005) Micro-Mar: a database for dynamic representation of marine microbial biodiversity. *BMC Bioinformatics*, 6:22.
- ★ Mira A. and **Pushker R.** (2005) The silencing of pseudogenes. *Molecular Biology and Evolution*, 22(11): 2135-2138.
- ★ Mira A., **Pushker R.**, Legault B. A., Moreira D. and Rodríguez-Valera F. (2004) Evolutionary relationships of *Fusobacterium nucleatum* based on phylogenetic analysis and comparative genomics. *BMC Evolutionary Biology*, 4:50.
- ★ **Pushker R.**, Mira A. and Rodríguez-Valera F. (2004) Comparative genomics of gene-family size in closely related bacteria. *Genome Biology*, 5:R27.

Book Chapter

- ★ Mira A. and **Pushker R.** (2007) Genome architecture and evolution of bacterial pathogens. In: *Evolutionary Biology of Bacterial and Fungal Pathogens*. (Baquero F., Nombela C., Cassell G. H. and Gutiérrez J. A., Eds.). Chapter 12. ASM Press, Washington, D.C.

Invited Reviewer

- ★ *Bioinformatics*, Oxford University Press
- ★ 3rd ISCB Student Council Symposium (**SCS3**), Vienna, Austria

Research Talks

- ★ Micro-Mar, 2007 updates Jul. 2007
The 3rd Annual International Conference of Metagenomics, UCSD, San Diego, CA.
- ★ RNAGeneTracer: a tool to trace evolutionary history of the ribosomal genes Jun. 2007
Evolutionary Genomics Group, Universidad Miguel Hernández, Alicante, Spain
- ★ RNAGeneTracer: a tool to trace evolutionary history of the ribosomal genes Dec. 2006
International Symposium on Computational Biology & Bioinformatics, Bhubaneshwar, India
- ★ Evolution of genome size in Bacteria Nov. 2005
MRC Laboratory of Molecular Biology, University of Cambridge, UK
- ★ Micro-Mar: a database for dynamic representation of marine microbial biodiversity Nov. 2005
Max Planck Institute for Marine Microbiology, Bremen, Germany
- ★ Evolution of genome size in Bacteria Aug. 2005
CMBI, Radboud University, Nijmegen, Netherlands
- ★ Evolution of genome size in Bacteria Aug. 2005
Royal College of Surgeons, Dublin, Ireland
- ★ Micro-Mar: a prokaryotes database to correlate habitat with taxonomy Jun. 2004
University of East Anglia, Norwich, UK

Fellowships/Grants

- ★ Metagenomics 2007 Student Travel Grant Jul. 11 - 13, 2007
3rd Annual International Conference on Metagenomics, UCSD, CA
- ★ ECCB 2006 Travel Fellowship Jan. 21 - 24, 2007
5th European Conference on Computational Biology, Eilat, Israel
- ★ Science Foundation Ireland Fellowship Apr. 2006 - Mar. 2009
Conway Institute, University College Dublin, Ireland
- ★ European Commission Fellowship Jul. 2003 - Mar. 2006
University of Miguel Hernández, San Juan, Alicante, Spain
- ★ TCG Chembiotek Fellowship Jan. 2002 - Jun. 2003
Institute of Genomics & Integrative Biology, Delhi, India

Posters Presented

- ★ **Pushker R.**, Mira A. and Rodríguez-Valera F. (Jan. 2007) Genomic location and the evolution of paralogous genes. 5th European Conference on Computational Biology, Eilat, Israel.
 - ★ **Pushker R.** and Mira A. (Dec. 2006) Genome asymmetry : evaluating genome balance hypothesis in bacteria. International Conference in Bioinformatics (INCOB) 2006, New Delhi, India.
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Posters communicated

- ★ **Pushker R.**, Gautam R. and Mira A. (Jul. 2007) RNAGeneTracer: a tool to trace evolutionary history of the ribosomal genes. 15th Annual International Conference on Intelligent Systems for Molecular Biology (ISMB) & 6th European Conference on Computational Biology (ECCB), Vienna, Austria.
- ★ D'Auria G., **Pushker R.**, Alba-Casado J.C. and Rodríguez-Valera F. (Oct. 2006) Micro-Mar: a database for dynamic representation of marine microbial biodiversity. The First Annual International Conference of Metagenomics, UCSD, CA.
- ★ D'Auria G., **Pushker R.**, Alba-Casado J.C. and Rodríguez-Valera F. (Jun. 2006) Micro-Mar: a database for dynamic representation of marine microbial biodiversity. International Census of Marine Microbes 1st Annual Meeting, The Netherlands.
- ★ Mira A., **Pushker R.** and Rodríguez-Valera F. (Sep. 2005) Evolution of paralogous genes. CIAR Evolutionary Biology Program, Nanaimo, Canada.
- ★ Mira A. and **Pushker R.** (Sep. 2005) The silencing of pseudogenes. 4th European Conference on Computational Biology, Madrid, Spain.
- ★ **Pushker R.**, D'Auria G., Alba-Casado J.C. and Rodríguez-Valera F. (Sep. 2005) Micro-Mar: a database for dynamic representation of marine microbial biodiversity. XX Congreso Nacional de la SEM, Cáceres, Spain
- ★ Alba-Casado J. C., **Pushker R.** and Rodríguez-Valera F. (Nov. 2004) Micro-Mar : a marine prokaryotes database to correlate habitat with taxonomy. International Conference on Marine Biodiversity Data Management, Hamburg, Germany.
- ★ Zaballos M., López-López A., Ovreas L., Bartual S. G., Legault B., Alba J. C., **Pushker R.**, Dae F. L. and Rodríguez-Valera F. (Aug. 2004) Comparison of bacterial diversity at two offshore oceanic locations and depths. 10th International Symposium on Microbial Ecology ISME 10, Cancun, Mexico.
- ★ Alba-Casado J. C., **Pushker R.** and Rodríguez-Valera F. (Oct. 2003) Micro-Mar : A marine prokaryotes database to correlate habitat with taxonomy. XIX Congreso Nacional de Microbiología, Santiago de Compostela, Spain.

Meetings & Workshops

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| ★ Innovations in Drug Discovery and Development
IBM Life Sciences Virtual Conference and Expo | Apr. 16 - 17, 2008 |
| ★ MASAMB-2006, Mathematical and Statistical Aspects of Molecular Biology
Conway Institute, University College Dublin, Ireland | Apr. 10 - 11, 2006 |
| ★ GEMINI: Gene Discovery through Environmental Metagenomics, Annual Meeting
University of East Anglia, Norwich, UK | June 26, 2004 |
| ★ Parallel and Grid Computing
Institute of Physics, Bhubaneswar, India and INRIA, France | Mar. 11 - 16, 2002 |
| ★ C++ and Object Oriented Programming
Indian Institute of Technology, Delhi, India | Jul. 09 - 28, 2001 |
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Memberships

- ★ International Society for Computational Biology (ISCB), Student Membership 2006 - Present
- ★ Public Library of Science (PLoS), Individual Membership 2006
- ★ Institute of Electrical and Electronics Engineers (IEEE), Student Membership 2006

Courses done during PhD

- ★ Population Genetics & SNP Analysis Jun. 6 - 7, 2007
Institute of Molecular Medicine, St James's Hospital, Dublin, Ireland
- ★ Drug Design: Principles and Techniques Apr. 19 - 24, 2007
Royal College of Surgeons in Ireland, Dublin, Ireland

Technical Skills

- Programming Languages** : C, PERL, PHP, SQL, BioPERL
- Internet Technologies** : Javascript, CGI in C and PERL,
LAMP (The open source web platform)
- Markup Languages** : HTML, L^AT_EX
- Databases** : MySQL
- Software Tools** : CVS, GIMP, gnuplot, Adobe Photoshop, Illustrator and Flash
- Bioinformatics Tools** : BLAST, EMBOSS, PHYLIP, PAML, Clustalw,
FASTGroup, GLIMMER, OriLoc, Reputer, DNAMaster, ARB
KEstimator, BioEdit, TreeView, Artemis
- Operating Systems** : Linux (Fedora), Unix
MS Windows 9X/NT/2K/XP/Vista
- Parallel Computers** : Linux Cluster having 28 processors
Compaq Alpha Server having 32 processors

Personal Details

- Father's Name : Ram Swaroop
 - Date of Birth : 1978
 - Gender : Male
 - Marital Status : Single
 - Nationality : Indian
 - Correspondence : Clinical Bioinformatics Lab
 - Address : UCD Complex & Adaptive Systems Laboratory
University College Dublin
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