
Suyog S. Kuwar

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EDUCATION

Max Planck Institute for Chemical Ecology, Jena, Germany

01/2010 - 06/2016

- PhD, Molecular Biology, Excellent (summa cum laude / with highest distinction)
- IMPRS International Max Planck Research School fellow

Institute of Bioinformatics and Biotechnology, University of Pune India

06/2004 - 03/2009

- Master of Science in Biotechnology (5 years integrated)

SKILLS

Data Analysis	Molecular Biology	Languages	Other
<ul style="list-style-type: none">• C-Programming-Basic• R-Studio• Transcriptome analysis• Statistics• Bioinformatics• Mathematical modeling (basic)• Python (intermediate)	<ul style="list-style-type: none">• Phage display, PCR/qPCR• Protein modifications/ engineering• Basic separation techniques• Western blot/pull down assay• Enzyme Kinetics• Microbiology, Animal and Plant cell culture technique• Fermentation technology	<ul style="list-style-type: none">• Marathi (Native)• Hindi (Native)• English (fluent)• German (Basic)	<ul style="list-style-type: none">• Project coordination• Report preparation• Public outreach• Scientific writing• Experimental design• Adobe package• Website development-Jekyll

RESEARCH EXPERIENCE

L.V. H. Arts, Science and Commerce College, Panchavati, Nashik, India

08/2019 - 04/2020

Department of Entomology and Nematology, University Florida, USA

02/2017 - 12/2018

Post Doc, Department of Entomology, Iowa State University, USA

10/2016 - 02/2017

- “Analysis of next generation DNA sequencing data using linux command line tool for the modification of toxins derived from *Bacillus thuringiensis* for enhanced toxicity against selected insect pests”

PhD, Max Planck Institute for Chemical Ecology, Jena, Germany

01/2010 - 06/2016

- “The adaptive response of the serine protease superfamily of the cotton bollworm *Helicoverpa armigera* to dietary protease inhibitors”
- Created a comprehensive resource of all the digestive proteases from *H. armigera*, *S. frugiperda* and *M. sexta*.
- Analyzed the transcriptome data using MetaboAnalyst2.0, MeV and Genespring Gx.
- Analyzed the proteomics data using PLGS software.

M. Tech., Indian Institute of Science Education and Research, Pune

04/2009 - 12/2009

- “Mass spectrometry data analysis for phloem proteome of potato in response to pathogen challenge”

M. Sc Project, National Chemical Laboratory, Pune, India

06/2008 - 03/2009

- “Identification and Characterization of *Helicoverpa armigera* gut amylases”
- Analyze enzyme assays using SPSS

M. Sc Biotechnology, Savitribai Phule Pune University, Pune

06/2006 - 03/2007

ACHIEVEMENTS

- ISCE Student Poster Award: for outstanding contribution as a poster in Melbourne, Australia, Aug 2013
- Selected to attend the Centre for Plant Integrative Biology (CPIB) summer school “Mathematical Modeling for Biologist 2010” at Sutton Bonington Campus, Nottingham, UK
- International Max Planck Research School Fellowship for PhD studies, Max Planck Institute for Chemical Ecology, Jena, Germany. January 2010
- Graduate Aptitude Test in Engineering 2009 Qualified with 84.45 percentile-All India Exam

PUBLICATIONS

- Published 6 papers in peer reviewed international journals, e.g. BMC Genomics, Scientific reports etc.

PROFESSIONAL SERVICES

Article reviewing for: PLOS ONE, Biology — Open Access Journal of Biochemistry & Molecular Biology, Life — Open Access Journal of Origins and Evolution of Life