Migun Shakya, PhD

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EDUCATION

Ph.D Genome Sciences \diamond University of Tenneessee \diamond Knoxville, TN 2013 **Dissertation** Validating Approaches for Studying Microbial Diversity to Characterize Communities from Roots of *Populus deltoides*

B.S Biology ◊ Southwerstern Oklahoma State University ◊ Weatherford, OK

2008

RESEARCH EXPERIENCE

Computational Biology Post doc

2016 - present

- Bioscience Division, Los Alamos National Laboratory, Los Alamos, NM
- · Developing pipeline for RNA-seq data analysis to DE genes in eukaryotes and prokaryotes.
- · Developing pipeline for evolutionary analysis based on SNPs and maximum likelihood.

Postdoctoral Associate 2013 - 2016

- · Department of Biology, Dartmouth College, Hanover, NH
- · Evolution of Gene Transfer Agents across Alphaproteobacteria
- · Developing SVM-based algorithm to distinguish phage genes from GTA homologs.

Graduate Research Assistant

2009 - 2013

- · Genome Science and Technology Program
- · University of Tennessee and Oak Ridge National Lab, Oak Ridge, TN

PUBLICATIONS

Hynes A.P*, **Shakya M.***, Mercer R.G, Grull M.P, Brown L., Davidson F, Steffen E., Matchem H., Peach M.E, Berger T, Grebe K., Zhaxybayeva O., Lang A.S, (2016). Functional and evolutionary characterization of a gene transfer agent's multi-locus "genome?", *Molecular Biology and Evolution* *These authors contributed equally

Hurt, R.A., Robeson II M.S., **Shakya M.**, Moberly, J.G., Vishnivetskaya, T.A., Gu B., and Elias, D.A. Improved Yield of High Molecular Weight DNA Coincides with Increased Microbial Diversity Access from Iron Oxide Cemented Sub-Surface Clay Environments. *Plos ONE* 2014.

Shakya, M., Gottel N., Castro H., Yang Z.K., Gunter L., Labbe J., Muchero W., Bonito G., Vilgalys R., Tuskan G., Podar M., Schadt C.W. A Multifactor Analysis of Fungal and Bacterial Community Structure in the Root Microbiome of Mature Populus deltoides Trees. *Plos ONE* 2013. doi:10.1371/journal.pone.0076382

Shakya, M., Quince, C., Campbell, J., Yang, Z., Schadt, C.W., Podar, M. Comparative metagenomic and rRNA microbial diversity characterization using Archaea and Bacteria synthetic communities *Environmental Microbiology* 2013.doi:10.1111/1462-2920.12086

Flores E.G., **Shakya**, **M.**, Meneghin J., Podar, M., Seewald., J.S., Reysenbach A-L. Inter-field Variability in the Microbial Communities of Hydrothermal Vent Deposits from a Back-Arc Basin. *Geobiology* 2012. doi:10.1111/j.1472-4669.2012.00325.x

BOOK CHAPTERS

Culley A.I., **Shakya M.**, Lang A.S. (2015) Viral evolution at the limits. In *Microbial Evolution under Extreme Conditions* (pp.209-222).Germany.degruyter.

HONORS AND AWARDS

University of Tennessee, Knoxville

2011 Genomic Sciences Meeting Student Travel Grant

Southwestern Oklahoma State University,

In-State Tuition Waiver 2003-2008

Ottis and Buena Ballard Memorial Scholarship 2007

Who is Who Among American Universities 2008

SELECTED SCIENTIFIC CONFERENCES

Li P.E., Lo C-C., Anderson J., Davenport K., Bishop-Lilly K., Xu Y., Voegtly L., Feng S., Senin P., **Shakya M.**, Philipson C., Gans J., Miller N., Challacombe J., Hamilton T., Chain P.S.G. Addressing the Analysis Bottleneck Created by Next Generation Sequencing (NGS) Using EDGE Bioinformatics. In:asm microbe, June 1-5, 2017. Poster and Selected talk.

Shakya M., Senin P., Feng S., Lo C., Hu. B, Chain P.S.G. PiReT: Pipeline for Reference Based Transcriptomics. In: *Sequencing*, *Finishing*, *analysis In the Future Meeting*, May 16-18, 2017. Poster.

Shakya M. and Zhaxybayeva O. Do bacteria domesticate viruses?. In: *Microbiology and Molecular Pathogenesis Program Retreat*, February 11-12, 2016. Poster and Talk.

Shakya M. and Zhaxybayeva O. Origin and Evolution of gene transfer agents in α -proteobacteria. In: *Mobile Genetic Elements: In Silico, In Vitro, In Vivo*, September 3-5, 2015. Selected talk.

Shakya M. and Zhaxybayeva O. Origin and Evolution of gene transfer agents in α -proteobacteria. In: *Gordon Research Conference: Microbial Population Biology*, July 19-24, 2015. Poster.

Shakya M., Drivers of microbial community structure in rhizosphere and endosphere of *Populus deltoides*.In: *American Society for Microbiology KY-TN Branch Meeting*, October 26-27, 2012.Talk.

Shakya M., Gottel, N.R., Castro, H.F., Yang Z.K., Kerley M., Podar M., Doktcyz, M.J., Schadt, C.W. Archaea associated with Populus and its surrounding soil and trees. In: *International Society of Microbial Ecology 14, Copenhagen, Denmark*, August 19-24, 2012. Poster.

Shakya M., Gottel, N.R., Castro, H.F., Yang Z., Kerley M., Podar M., Doktcyz, M.J., Schadt, C.W. Plant-Microbe Interface: Dynamics of Bacterial Microbiome of Populus del-toides In: 2012 Department of Energy Genomic Science Awardee Meeting X, February 26-29, 2012. Poster

Shakya M., Gottel, N.R., Castro, H.F., Yang Z., Kerley M., Podar M., Doktcyz, M.J., Schadt, C.W. Plant-Microbe Interface: Dynamics of Bacterial Microbiome of *Populus deltoides* In: *2012 Department of Energy Genomic Science Awardee Meeting X*, February 26-29, 2012. Poster.

Shakya M., Campbell, J.C., Schadt, C.W., Podar M. Experimental and Computational Approaches for Microbial Diversity Characterization Using Artificial Communities. In: *2011 Genomic Science Awardee Meeting IX*, April 10-13, 2011. Poster and Talk.

Shakya M., Campbell, J.C., Yang, Z.K., Podar M. Experimental and Computational Approaches for Microbial Diversity Characterization Using Artificial Communities In: *International Society of Microbial Ecology 13*, August 22-27, 2010. Poster.

TEACHING EXPERIENCE

Los Alamos National Laboratory, Los Alamos, NM

Taught a class on phylogenetic analysis for annual workshop.

Taught a class on Transcriptomics analysis for annual workshop.

University of Tennessee, Knoxville

Teaching Assistant for BCMB 230: Human Physiology Fall 2009 (2 sections)

Teaching Assistant for BIO 140: Organization and Function of the Cell Spring 2010 (2 sections)

Southwestern Oklahoma State University

Teaching Assistant for Biological Sciences Fall 2007-Spring 2008

Grader for College Algebra and Trigonometry Spring 2007-Spring 2008

PEER REVIEW ACTIVITIES

Reviewer for: FEMS Microbiology Ecology, Canadian Journal of Microbiology, Frontiers in Microbiology, Molecular Biology and Evolution, BMC Microbiology, MDPI - *Toxins*

RELEVANT SKILLS

Programming/Scripting: Python, Perl, R, Bioconductor

Bioinformatics: Genome, Microbial ecology, Metagenome, Metagenetics, Methylome, SNP analysis, phylo-

genetics, etc.

Other: Linux, shell scripting, high performance computing