

Akshar Lohith

Employment History

- 10/2014 – 08/2016 Junior Specialist Genome Sequencing-Nader Pourmand Lab,
Santa Cruz, CA
- Perform sequencing activities including automated cDNA synthesis and Illumina sequencing library preparation of RNA samples from various projects. Processing and QC methods for sample preparation include quantification with qPCR, bead purification with AmPureXP and Caliper, and DNA size selection with Caliper LabChipXT. Operation of sample-handling robots, including programming sample-handling robots to automate Smart-Seq2 cDNA synthesis and KAPA Biosystems' Hyper-Prep Library Preparation protocols. Sample handling robot protocols include genomic DNA extraction from whole blood. Extractions are then processed for Illumina Sequencing. Maintain server file-share space, including Illumina Sequencing data sample demultiplexing and transfer of data to research collaborators. Perform research activities with single cell nanobiopsies from mammalian cells for several projects. Biopsied samples underwent cDNA synthesis for RNA sequencing in multiple projects. Optimize injection protocol and worked to automate nanoinjections of various materials into cells with BioStinger, single-cell interrogation instrument. For automating nanoinjections, TNF α :NF κ B pathway inhibitory drugs are injected into a mammalian cell culture modified for GFP-expression under TNF α exposure. Other materials injected include various toxins for Reactive Oxygen Species (ROS) detection with fluorescent probe DCFDA. Performed fluorescent imaging and subsequent analysis on cells exposed to intracellular pH probe BCECF. Projects include; ISPY 2 and SU2C RNA Sequencing cancer research projects in collaboration with groups at UCSF, RNA nanobiopsy from human sperm for genetic mutation analysis and sex identification in collaboration with Stanford School of Medicine, Intracellular ROS detection by nanoinjection of toxins, and Single-cell Automated Imaging and Nanoinjection.
- 6/2014 – 9/2014 Genome Sequencing Intern Genome Sequencing-Nader Pourmand Lab,
Santa Cruz, CA
- Learned and operated sample-handling robots to perform automated cDNA synthesis and Illumina Library Preparation for RNA Sequencing on Illumina Sequencing platforms. Performed laboratory sample QC methods such as qPCR library quantification and Bioanalyzer High-Sensitivity DNA Assay as well as sample purifications with AmPureXP and Caliper beads.
- 2/2012 – 6/2014 Undergraduate Research Nader Pourmand Lab, Santa Cruz, CA
Assistant
- Fabricated nanopipettes from glass capillaries, using Sutter P2000 laser puller, to use with the modified Scanning Ion Conductance Microscope setup developed for single cell analysis. This platform is capable of single cell nanobiopsies and injections, and was used for senior thesis project: Illuminating the Mitochondria and the Mitochondrial Genome's role in Neurodegeneration with nanopipettes. Using this platform, single mammalian cells cultured in compartmentalized Death Galaxy device were nanobiopsies for RNA sequencing, a new application of the single cell nanobiopsy technique. Maintained several mammalian cell cultures and extracted genomic DNA from cell populations using Invitrogen Genomic DNA mini kit.
- 10/2011 – 6/2014 Intramural Referee Office of Physical Education, Recreation and
Sports at UCSC, Santa Cruz, CA
- Refereed intramural soccer, futsal and dodgeball games on campus
- 6/2013 – 9/2013 Bioinformatics Intern Boston Heart Diagnostics, Framingham, MA
- Explored to expand image analysis of High Density Lipoprotein (HDL) size separation maps in order to study HDL particle formation and potential cardiovascular health benefits they may have. This was performed through analysis of over 3700 HDL maps. Microsoft Excel Macro-enabled Templates and workflows were developed for the Image Analysis Data Collection team with SOP addendums.

(408) 887-3335 • lohith.akshar@gmail.com

Akshar Lohith

6/2009 – 9/2013

QA Intern

Tricipher, Inc., Los Gatos, CA

Verified expected flow of links on web-based software product, MyOneLogin, for quality and new features. Searched for errors in both production and pre-production versions.

References

References are available on request.