

Department of Mathematics and Statistics

COLLOQUIUM

Tuesday, October 28th, 2014

4:00 – 5:00 pm, Adel Mathematics Bldg., Room 164 (refreshments at 3:45)

Dr. Viacheslav Fofanov

Informatics and Computing Program
Department of Mathematics and Statistics
Northern Arizona University

Computational Challenges in Microbial Forensics

Abstract

Microbial Forensics seeks to use biochemical properties of bacterial samples to track a germ used as a weapon (as part of criminal activity or bioterrorism event) to its source. In this context, the presence of rare variants in otherwise homogeneous bacterial populations, could act as a sample's fingerprint. Recently developed High Throughput Sequencing technologies (HTS) can, in a cost acceptable manner, produce sufficient amount of sequence data for sensitive detection of ultra-rare variants (present in 0.1% of the sample). However, the specificity of ultra-rare variant SNP detection (at levels <1%) still remains a major limiting factor of this technology.

This talk will focus on the statistical, computational, and biological challenges (and potential solutions) associated Microbial Forensics, the HTS technology and the ultra-rare variant detection.

ACGT Seminar: Tuesdays 12:45 – 1:45 pm, AMB 164.

Applied Math Seminar (AMS): Thursdays 12:45 – 1:45 pm, AMB 164.

Friday Afternoon Undergraduate Mathematics Seminar (FAMUS) meets Fridays at 3pm, AMB 164