

Genetics and Genomics in Clinical Research

An Immersion Course for Investigators at UAB

October 10th-14th, 2016 from 8:30-12:30pm

\$50- Faculty, \$25-Post Doc and Fellows, \$10 – Students/Staff Continental Breakfast served daily

> Location: Finley Conference Center 720 20th St. S, Birmingham, AL 35233

REGISTER

This course will review the principles, major technologies, and experimental approaches in genetics and genomics through both lectures and hands-on activities.

Learning Objectives:

- 1. Design an approach to identify a gene responsible for a phenotype in a family that segregates in a Mendelian manner.
- 2. Devise an appropriately powered case-control or transmission disequilibrium study to identify single nucleotide polymorphisms in linkage disequilibrium with a multifactorial disorder.
- 3. Develop a study comparing patterns of gene expression or methylation levels in normal vs. pathological tissue.
- 4. Formulate a protocol involved human research subjects for a genetic or genomic study to be submitted for IRB review.
- 5. Choose between alternative genotyping or next generation sequencing platforms appropriate for specific applications.
- 6. Utilize major bioinformatics databases to analyze genomic data.





| Schedule: | | | | | |
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| | Monday, Oct 10 | Tuesday, Oct 11 | Wednesday, Oct 12 | Thursday, Oct 13 | Friday, Oct 14 |
| 8:30- 9:00 | Breakfast | Breakfast | Breakfast | Breakfast | Breakfast |
| 9:00- 10:00 | Review of Genetic Principles Dr. Bruce Korf | Human Population Genetics Dr. Greg Barsh | Analytics of next-generation sequencing data Dr. Greg Cooper | Copy number variants (CNVs) and chromosomal microarray technologies Dr. Fady Mikhail | Approaches to Bioinformati cs Data Analysis Dr. David Crossman |
| 10:15- 11:15 | Overview of Genetics and Genomics in Clinical Research Dr. Bruce Korf | Genome Wide Linkage and Association Analysis Dr. Greg Barsh | Genomic counseling Ms. Whitley Kelley | Genotyping Technologies Next-Generation sequencing Dr. Mike Crowley | Transcripto me and Pathway Analysis Dr. David Crossman |
| 11:30- 12:30 | Informed Consent and Ethical Issues Dr. Ashley Cannon | Genome Wide Association Analysis Dr. Hemant Tiwari | Variant Review Dr. Greg Cooper | Use of Public Databases and Genome Browser to Interpret CNV Data Dr. Fady Mikhail | Genetics in Medicine Dr. Bruce Korf |

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For further information please contact Shaila Handattu at hande@uab.edu