Association Mapping: GWAS and Sequencing Data

Instructors: Timothy Thornton and Michael Wu

Summer Institute in Statistical Genetics (SISG) July 2021

Introduction: Course Aims

This is a course on statistical methods and software for genetic association studies of complex traits. We aim to cover:

- Case-Control Association Testing
- Genetic Association Testing with Quantitative Traits
- Gene and Pathway Level Analysis
- Population Structure/Ancestry Inference
- Genetic Association Testing in Samples with Structure
- Interaction Analysis
- Collapsing Tests for Rare Variant Analysis
- Kernel Tests and Omnibus Tests for Rare Variants
- Power and Sample Size, Design Considerations, Emerging Issues

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Introduction: Resources

Importantly, the class site is:

http://faculty.washington.edu/tathornt/SISG2021.html

Contains (or will contain);

- PDF copies of slides (in color)
- Datasets needed for exercises
- Exercises for you to try
- Our solutions to exercises (later!)
- Links to software packages

Introduction: About Mike



- Associate Member, FHCRC Public Health Sciences Division
- Affiliate Associate Professor, UW Boistatistics
- Research in:

High dimensional data Kernel machine methods Variable selection and regularization Pathway and network based analysis Translational research Statistical genomics/genetics

Introduction: About Tim



- Professor,
 UW Biostatistics
- Affiliate Investigator, Fred Hutchinson
 Cancer Research Center
- Research in: Genetic Association Studies Correlated Genetic Data Inferring Genetic Ancestry Relatedness Estimation Pharmacogenomics

Our department's finest - here to help you:

Andrea Horimoto



Amarise Little

Anya Mikhaylova



Slack Channel

Expect to 'see' them on Zoom chat, and our Slack channel: https: //uwbiostatisticssisg.slack.com/archives/C0239BJBG5P

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Introduction: Course Structure

10 sessions, 80 minutes each, over 2.5 days

- What to expect in a typical session;
 - 40 mins teaching/lecture
 - 25 mins hands-on exercises
 - 15 mins summary, discussion