



INTERNATIONAL GENETIC EPIDEMIOLOGY SOCIETY

The past decade has seen a virtual explosion in the generation and analysis of genomic data. Due in part to data sharing policies, much of this information is available through controlled access databases, such as the database of Genotypes and Phenotypes ([dbGaP](#)) and the European Genome-phenome Archive ([EGA](#)). Although there are a growing number of examples of key scientific findings stemming from this shared data, we are only at the tip of the iceberg in terms of fully leveraging these rich resources.

Is there potential for scientific advances as a result of broad data sharing?
How can you capitalize on these resources?

WORKSHOP: Advancing Scientific Progress through Genomic Data Sharing: Access and Analysis

Sign up now!
\$75 fee

Guest Speakers:

Steve Sherry, PhD—
NCBI/dbGaP

Nancy Cox, PhD—University
of Chicago

Brandon Pierce, PhD—
University of Chicago

Marylyn Ritchie, PhD—
Penn State

Laura Bierut, PhD—
Washington University in St. Louis
School of Medicine

Justin Paschall, MS—Team
Leader- Variation/EGA/DGVA EBI

Martin Tobin, PhD—on
behalf of UK Biobank

**International Genetic Epidemiology Society
Conference**

Sunday, September 15, 2013, 1:00-5:00pm
Intercontinental Hotel, Magnificent Mile, Chicago, Illinois,
USA

**Limited
Seating!!**

What will be discussed?

- **Overview of access policies and mechanisms**
- **Review of currently available datasets, providing information on format and content**
- **Research Panel will share expertise and first-hand experiences using shared genomic data (from both dbGaP and EGA) in their own research programs and stimulate discussion on current scientific issues arising when using existing data for analysis and methods development and provide practical advice on topics including: facilitating data sharing and navigating common pitfalls and obstacles in accessing controlled access data**