Seminars for Second Year Ph.D. Students

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Impact of Coverage of Causal Variation in GWAS

WEDNESDAY, April 14, 2010, at 5:15 PM 110 Eckhart Hall, 5734 S. University Avenue

ABSTRACT

In this presentation I will present a method for measuring how well a particular platform or set of SNPs covers the causal variation in the human genome. I will discuss possible implications coverage has on finding individual causal variants in a genome-wide association study (GWAS) and, time permitting, implications on finding gene-gene interactions.

Information about building access for persons with disabilities may be obtained in advance by calling Kelly Macias at 773.834.5169 or by email (kmacias@galton.uchicago.edu).