Webinar

The Hunt for Missing Heritability

Challenges and Opportunities for Novel Locus Discovery in Non-European Populations

TUESDAY, JANUARY 31, 2012

12 noon Eastern • 9 a.m. Pacific • 5 p.m. UK

Characterizing the genetics of complex diseases has, to date, focused on common variants and predominantly on populations of European descent. GWAS methodologies have been successful in uncovering novel susceptibility loci for common disorders but the heritability of many disorders remains to be explained. Now, a growing number of investigators are looking beyond European cohorts to study common and rare variants in populations around the world, including African, Asian, and other ancestries in the hunt for novel susceptibility genes.

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Bethesda, MD

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THE THOUGHT-LEADERS ON OUR WEBINAR PANEL WILL:

- Discuss how population genetics integrates with the genetics of complex disease to reveal novel disease genes
- Describe how the discovery of population-specific rare variants expands our understanding of complex diseases
- Explain the importance of population-optimized strategies and tools that account for differences in genetic diversity and population admixture
- Answer your questions live!

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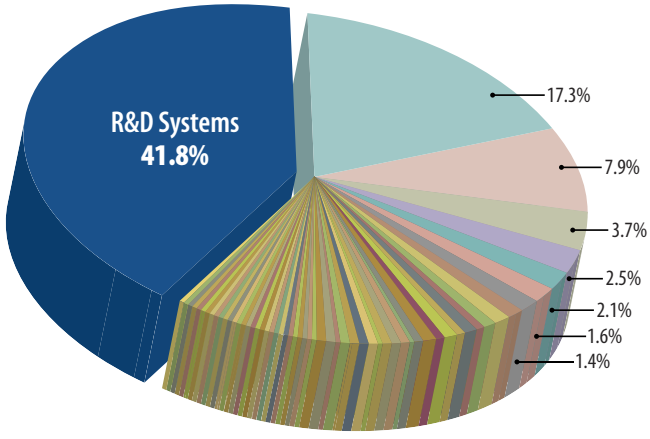

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