H3AFRICA GENOME ANALYSIS WORKSHOP

Disease Gene Discovery in African Populations Spier Hotel and Conference, Stellenbosch, South Africa

June 02, 2014, 9:00 AM - 4:00 PM

TOPIC	SPEAKER	TIME
Introduction to the workshop	Genome Analysis WG	09:00 - 09:05
Overview of H3Africa	Jennifer Troyer (NHGRI, NIH)	09:05 - 09:15
Study Designs for Disease Gene Discovery:		09:15 - 10:00
Genotyping (GWAS) versus Sequencing	Kathleen Barnes (Johns Hopkins University)	
Finding disease genes using whole exome capture	Phillip Awadalla (CARTaGENE/University of	10:00 - 11:00
and sequencing (WES) and WGS approaches;	Montreal)	
WES versus exome arrays	Adebowale Adeyemo (NHGRI, NIH)	
	Tea/Coffee Break	11.00 - 11.20
African Genome Variation: Implications for finding and replicating disease genes	Manj Sandhu (Wellcome Trust Sanger Institute)	11.20 - 11.50
One Study, Multiple African populations: Strength or Weakness?	Charles Rotimi (tbc) (NHGRI, NIH)	11:50 – 12:20
Special populations (e.g. admixed populations, population isolates) in gene discovery studies in Africa	Marlo Moller (University of Stellenbosch), Carlos Bustamante (tbc) (Stanford University)	12:20 – 13:15
	Lunch	13:15 - 14:15
PRESENTATIONS & PANEL DISCUSSION	Chair: Charles Rotimi (NHGRI, NIH)	14:15 - 15:55
Searching for the "ideal" GWAS chip for African	Durantaki	
populations (panel discussion – custom chip	Presentations: • Doing GWAS in African populations using	
task group, other scientists, industry partners)	• Doing GWAS in African populations using Illumina arrays (<i>Illumina rep</i>)	
	 Doing GWAS in African populations using 	
	Affymetrix arrays (<i>Affymetrix rep</i>)	
	 Evaluating current GWAS arrays for studies 	
	of African populations - Nicki Mulder	
	(H3ABioNet, UCT)	
	 Designing a new GWAS array for African populations - Zane Lombard (WITS) 	
	Panel	
	Michele Ramsay (WITS), Guida Landoure	
	(University of Bamako), Ambroise Wonkam (UCT), Zane Lombard (WITS), Nicki Mulder	
	(UCT), Rep (Illumina), Rep (Affymetrix), Rep	
	(BGI), Rep (CPGR), Rep (Ingaba Biotec)	
	(= 2-), (or or-), rep (mqubu 210000)	
Closing remarks	Genome Analysis WG	15:55 -16.00

Registration deadline: April 28 For more information, visit www.h3africa.org