

16th CRG Symposium: Seventh International Workshop on Genomic Epidemiology

Wednesday, Sept 20th	
14:00 – 14:15	Welcome
	Session 1 – Technology - Chair: Ivo Gut
14:15 - 14:45	Ivo Gut , CNAG-CRG, ES
14:45 - 15:15	
15:15 – 15:30	Selected Abstract talk
15:30 - 16:00	
16:00 - 16:30	Coffee break
16:30 – 17:00	Marc Marti-Renom , CNAG-CRG, ES
17:00 - 17:15	Selected Abstract talk
17:15 - 17:45	Manolis Kogevinas , CREAL, ES Contributions from twin studies to gene discovery
17:45 - 19:45	Welcome Reception + Poster Session (Terrasse)

Thursday, Sept 21st

	Session 2 - Epigenetics
09:00 - 09:30	Peter Jones , Van Andel Institute, USA Targeting Human Endogenous Retroviruses for Epigenetic Therapy
09:30 - 10:00	Caroline Relton , University of Bristol, UK
10:00 - 10:15	Selected Abstract talk
10:15 - 10:30	Selected Abstract talk
10:30 - 11:00	Iñaki Martin-Subero , IDIBAPS, Spain
11:00 - 11:30	Coffee break
	Session 3 – Resources / Big Initiatives
11:30 - 12:00	Hanns Lochmüller , Newcastle University, UK
12:00 - 12:15	Selected Abstract talk
12:15 - 12:30	Selected Abstract talk
12:30 - 13:00	
13:00 - 14:00	Lunch
14:00 - 14:30	Roderic Guigó , CRG, ES The human transcriptome across tissues and individuals
14:30 - 14:45	Selected Abstract talk
14:45 - 15:15	
	Session 4 – Diseases - Cancer
15:15 - 15:45	Nuria Malats , CNIO, ES
15:45 - 16:15	Richard Houlston , Institute Cancer Research, UK
16:15 – 16:45	Coffee break
16:45 – 17:15	Ben Lehner , CRG, ES

Friday, Sept 22nd	
09:00 - 09:30	Nuria López-Bigas , IRB, ES
09:30 – 09:45	Selected Abstract talk
09:45 – 10:00	Selected Abstract talk
	Session 4 – Diseases – Common Diseases
10:00 – 10:30	Momoko Horikoshi , RIKEN, JP
10:30 – 11:00	Patricia Munroe , Barts Institute, UK
11:00 - 11:30	Coffee break
11:30 - 12:00	George Thanassoulis , McGill, CA
12:00 – 12:15	Selected Abstract talk
12:15 - 12:30	Selected Abstract talk
12:30 - 13:00	Stephanie Debette , University Bordeaux, FR
13:00 - 14:00	Lunch
	Session 5 – Analytical methods and resources
14:00 - 14:30	George Davey-Smith , University of Bristol, UK Mendelian randomization: what does the future hold?
14:30 - 15:00	Dorret Boomsma , Vrije Universiteit Amsterdam, The Netherlands
15:00 - 15:30	Gonçalo Abecasis , University of Michigan, USA Sequencing and Analysis of 10,000s of Human Genomes: Challenges and Opportunities
15:30 - 16:00	Mark Caulfield , Genomics England, UK
16:00 – 16:30	Closing Remarks