Finding genes for human disease

from genetics to genomics

Oporto

7 - 11 July 2014

Faculty

Marianna Bevova, Michel Georges, Yurii Aulchenko

Program

Monday 7 th July			
9:30-9:45	Course introduction (Marianna Bevova)		
9:45-10:30	Lecture:	Variation in human genome (Marianna Bevova)	
10:45-11:30	Lecture:	Monogenic disorders. Linkage analysis (Marianna Bevova)	
11:45-13:30	Exercise:	Lod score exercise (Marianna Bevova)	
13:30-14:30	Lunch		
14:30-15:30	Lecture:	Monogenic disorders. Next generation sequencing (Marianna Bevova)	
15:45- 18:00	Exercise:	Virtual Cloning (haplotype analysis)	
Tuesday 8 th July			
9:30-10:00	Discussion Exercises (Marianna Bevova)		
10:00-13:30	Exercise:	Analysis of the next generation sequence data (Marianna Bevova)	
13:30-14:30	Lunch		
14:30-15:30	Lecture:	Multifactorial disease and association studies (Yurii Aulchenko)	
15:30-18:00	Exercise:	Introduction to R and association analysis (Yurii Aulchenko)	
Wednesday 9 th July	<i>I</i>		
9:30-10:00	Lecture:	Genome wide association studies (Yurii Aulchenko)	
10:15-11:00	Lecture:	Power and coverage of genome wide association studies	
11.30-13.30	Exercise:	(Yurii Aulchenko) Power estimation; GWAS QC	
13:30-14:30	Lunch		
14:30-15:30	Lecture:	Confounding in Genome wide association studies (Yurii Aulchenko)	

15:30-16:30	Exercise:	GWAS in GenABEL
		(Yurii Aulchenko)
17:00-18:00	Exercise:	Developing a study design

Thursday 10th July

9:30-10:15	Lecture:	Positional cloning for complex diseases (Michel Georges)
10:30-11:15	Lecture:	Functional annotation of the human genome (Michel Georges)
11:30-13:30	Exercise:	Developing a study design
13:30-14:30	Lunch	
14:30-18:00	Exercise:	Developing a study design

Friday 11th July

9:30-10:15	Lecture:	Applications of the medical genetics findings (diagnostic, pharmacogenomics)
10:30-11:15	Lecture:	(Marianna Bevova, Yurii Aulchenko) Beyond exome sequencing and GWAs: Functional Genomics
1:15-13:30	Exercise:	Developing a study design
13:30-14:30	Lunch	
14:30-17:00		Presentation and discussion of the study design. Conclusions