

SEQUENCE MAPPING AND ASSEMBLY ASSESSMENT PROJECT (SMAAP) RGASP3 / dnGASP

April, 5th-7th, 2011

PROGRAM

	Tuesday, April 5 th		Wednesday, April 6 th		Thursday, April 7 th
8:45	Registration				
9:00	Presentation			9:00	Greg Grant
9:30-11:00	Introduction to dnGASP/RGASP Ivo Gut/Roderic Guigo Report on GAW/Assemblathon Tyler Alioto/Paolo Ribeca Evaluation methods and overview of results Tyler Alioto/Andre Corvelo	9:00-11:00	Results of Evaluation Tyler Alioto/Andre Corvelo	9:40-11:00	Evaluation of predictions
11:00	Coffee break	11:00	Coffee break	11:00	Coffee break
11:30-1:00	11:30am-12:10pm Daniel Zerbino 12:10pm-12:35pm Shaun Jackman 12:35pm-1:00pm Zemin Ning	11:30-1:00	11:30am Discussion 12:30pm pmThe way ahead Ivo Gut (CNAG)	11:30-1:00	Evaluation of predictions
1:00	Lunch	1:00	Lunch	1:00	Lunch
2:00-2:40	Li Yingrui	2:00-3:00	Tom Gingeras	2:00-3:30	Validation Discussion on: a) Plans for Experimental Work b) Possible Publication c) New Editions
2:40-4:00	2:40 Bryan Downie(Mathias Platzer) 3:05 Victor Solovyev (Softberry) 3:30 Rayan Chikhi (Symbiose)	3:00-4:30	3:00pm Welcome by Roderic Guigó (CRG) 3:05pm Tim Hubbard "Gencode" 3:30pm Jen Harrow RGASP 1 and RGASP 2 4:00pm Paul Bertone "RGASP 3"		
4:00	Coffee break	4:30	Coffee break		
4:30-6:00	4:30 Alexey Sergushichev (SPbSU ITMO) 4:55 Ekaterina Khmrameeva (Genoalytica) 5:20 Jared Simpson (Richard Durbin)	5:00-7:30	5:00 Zening Ning (Seq informatics) 5:15 Paolo Ribeca (GEM) 5:30 Nicola Vitulo (CRIBI) 5:45 Victor Solovyev (Softberry) 6:00Xi Wang (Tsinghua University) 6:15 Jan Prins (MapSplice) 6:30 Sheng Li (Mason Lab) 6:45 Alexander Dobin (Gingeras lab) 7:00 Gunnar Räscht (Räscht lab) 7:15 Thomas Wu (Genentech)		