Tuesday 23/8	
	WABI
08:30-09:00	Registration
09:00-10:15	Session 6CChair: ?
09:00	Sorina Maciuca, Carlos Del Ojo Elias, Gil McVean and Zamin Iqbal: A natural encoding of genetic variation in a Burrows-Wheeler Transform to enable mapping and genome inference
09:25	Adam Novak, Erik Garrison and Benedict Paten: A Graph Extension of the Positional Burrows-Wheeler Transform and its Applications
09:50	Svend V Nielsen, Simon Simonsen and Asger Hobolth: Inferring Population Genetic Parameters: Particle filtering, HMM, Ripley's K-function or Runs of Homozygosity?
10:15-10:35	Coffee Break
10:35-11:25	Session 7CChair: Jan Baumbach
10:35	Leena Salmela and Alexandru I. Tomescu: Safely filling gaps with partial solutions common to all solutions
11:00	Jay Ghurye and Mihai Pop: Better Identification of Repeats In Metagenomic Scaffolding
11:25-11:30	Break
11:30-12:30	WABI Invited Kiyoshi Asai Chair: ?
12:30-14:00	Lunch
14:00-15:15	Session 8CChair: ?
14:00	Yun Deng and David Fernández-Baca: Fast Compatibility Testing for Phylogenies with Nested Taxa
14:25	Arnon Benshahar, Vered Caspi, Danny Hermelin, and Michal Ziv-Ukelson: A Biclique Approach to Reference Anchored Gene Blocks and its Applications to Pathogenicity Islands
	Mohammed El-Kebir, Ben Raphael, Ron Shamir, Roded Sharan, Simone Zaccaria, Meirav Zehavi and Ron Zeira: Copy-Number Evolution Problems: Complexity and Algorithms
14:50	Copy-Number Evolution Froblems. Complexity and Algorithms
15:15-15:30	Break
15:30-16:45	Session 9CChair: ?
15:30	WABI Poster
15:55	2 6666
16:20-16:35	Break
16:35	Session 10Cchair: ?
16:50	

17:15	
17:40	WABI Poster
18:05	
18:30	Bus to conference dinner
19:00	Conference Dinner (Godsbanen, downtown)