

4th Annual Meeting of NGFN-Plus and NGFN-Transfer

„Next Generation Sequencing Satellite Symposium“ (org. Philip Rosenstiel) September 26, 2011 Urania, Berlin

04:30-04:40 pm	Welcome and Introduction Philip Rosenstiel
Session I	
04:40-05:00 pm	Joris Veltmann (UMC St Radboud) Title: Identifying de novo mutations from complex NGS data sets
05:00-05:20 pm	Christoph Bock (Max-Planck-Institut für Informatik) Title: The methylome and beyond: using NGS to identify complete epigenomes
05:20-05:40 pm	James Hicks (Cold Spring Harbor Laboratory) Title: Molecular evolution of single tumor cells
05:40-06:00 pm	<i>Coffee Break</i>
Session II	
06:00-06:20 pm	Marie-Laure Yaspo (Max Planck Institute for Molecular Genetics) Title: To be announced
06:20-06:40 pm	Stefan Haas (Max-Planck Institute for Molecular Genetics) Title: Computational challenges of large scale NGS data sets
06:40-07:00 pm	Marc Zapatka (German Cancer Research Center) Title: After the first personal cancer genomes – where do we go ?
07:00-07:20 pm	Peer Bork (European Molecular Biology Laboratory EMBL) Title: Identification of functional and phylogenetic signals in gut metagenomes
07:20-07:30 pm	Concluding remarks
07:30 pm	<i>Supper</i>