

3rd Annual Meeting
NGFN-Plus and NGFN-Transfer in the
Program of Medical Genome Research

Satellite Symposium
Next-Generation Sequencing

November 25, 2010

Scientific Organization: Bernhard Korn, Institute for Molecular Biology (IMB), Mainz, Germany

12.00 - 12.05 pm Introduction and Chair: **Bernhard Korn**

Session 1

12.05 – 12.25 pm Microdroplet-based PCR Amplification for Large Scale Targeted Sequencing
Steve Picton, RainDance Technologies, Inc., Lexington MA, USA

12.25 – 12.45 pm Fully automated library preparation for the Illumina Genome Analyzer – a comparison with the manual method
Kathryn Stemshorn, Cologne Center for Genomics, Germany

12.45 – 1.05 pm ‘Private variants’ – a common cause of disease
Alexander Hoischen, Nijmegen Centre for Molecular Life Sciences, The Netherlands

1.05 – 1.25 pm “NGS to Examine the Transcriptome: A Focus on ncRNAs and Viruses”
John Castle, TrOn GmbH, Mainz, Germany

1.25 – 1.45 pm Sequencing using Semiconductor Technology
Armin Winands, Ion Torrent Inc., San Francisco, USA

1.45 – 2.00 pm Coffee Break

Session 2

2.00 – 2.20 pm Complete Human Genome Sequencing for Large-Scale Disease Studies
Rick Tearle, Complete Genomics, Inc., Mountain View, CA, USA

2.20 – 2.40 pm Cool Runnings with the Roche 454 GS Junior
Ralph Oehlmann, IMG M Laboratories GmbH, Martinsried, Germany

- 2.40 – 3.00 pm High performance genomics with Galaxy on the ground and in the clouds
Nate Coroar, PennState University, Pennsylvania, USA
- 3.00 – 3.20 pm NGS Data Analysis: Genomic and Transcriptomic Variants Identification – Characterization – Visualization
Martin Seifert, Genomatix Software GmbH
- 3.20 – 3.40 pm A comprehensive platform for full-genome data analysis
Roald Forsberg, CLC bio, Aarhus, Denmark
- 3.40 – 4.00 pm From third generation sequencing to a multi-scale biology approach to understanding and treating human disease
Tobias W.B. Ost, Pacific Biosciences, Menlo Park, CA, USA
- 4.00 – 4.30 pm Coffee Break**