

	July 20 Wednesday	July 21 Thursday	July 22 Friday	July 23 Saturday
9:00-9:50		Introduction of the participants	Novel Computing Systems in Biology Llc: <i>A Java library for binding site search algorithms</i>	Peter Laczko (Technical University Budapest): <i>Finding rearrangements from NGS data</i>
9:50-10:40		Istvan Miklos: <i>The TransFoot software package</i>	Philip Stegmaier (BIOBASE Gmbh): <i>Tools and databases for NGS analysis</i>	István Nagy (BayGen): <u>Single experimental setup = multiple evaluation possibilities</u>
10:40-11:10		Coffee break	Coffee break	Coffee break
11:10-12:00		Miklos Csuros (University of Montreal): <u>Algorithm (re-)engineering for next-generation sequencing</u>	Adam Novak (Oxford): <i>Using spannoids in comparative sequence analysis</i>	Balint L. Balint (University of Debrecen): <u>Understanding variability in ChIP Seq experiments and the consequences for data interpretation</u>
12:00-12:50		Attila Berczes (Omixon Ltd.): <u>Accurate Genome Variant Detection using Omixon Gapped SOLiD Alignment tools</u>	Endre Sebestyén (Agricultural Research Institute of the Hungarian Academy of Sciences): <u>The DoOP database</u>	Endre Barta (University of Debrecen): <i>Meta-analysis of ChIP-seq data</i>
12:50-14:00		Lunch	Lunch	Lunch
14:00-15:00	Registration	<i>Tutorial: Plugin Workshop for the CLC bio Platform. Workbench as well as Server. Tutor: Mikael Flensburg, Clc Bio</i>	<i>Tutorial on the DoOP database</i>	Discussion: <i>Towards personalized medicine</i> (Moderator: István Peták, KPS Molecular Diagnostics)
15:00-15:30	Coffee break	Coffee break	Coffee break	Coffee break
15:30-17:00	Introduction of the COGANS consortium Press conference	Tutorial continued.	Poster session	Informal discussions
17:00-19:00	Wine & cheese party	Informal discussions	Sightseeing & social event	
19:00-				