

UPDATED SCHEDULE FOR TUESDAY MORNING, SEPTEMBER 28

Tuesday, September 28		
8:30 AM	Registration	
9:00 AM	Keynote 4—Rainer Spang. From large datasets to small networks	
	PROCEEDINGS TRACK (Auditorium)	TECH TRACK (Vd Goes)
9:50 AM	PT18 - Prototypes of elementary functional loops unravel evolutionary connections between protein functions. <i>Alexander Goncarenko and Igor Berezovsky</i>	
10:10 AM	PT19 - Improved sequence-based prediction of disordered regions with multilayer fusion of multiple information sources. <i>Marcin J. Mizianty, Wojciech Stach, Ke Chen, Kanaka Durga Kedarisetti, Fatemeh Miri Disfani and Lukasz Kurgan</i>	TT7 - CLC bio, a comprehensive platform for NGS data analysis. <i>Roald Forsberg</i>
10:30 AM	PT20 - A predictor for toxin-like proteins exposes cell modulator candidates within viral genomes. <i>Guy Naamati, Manor Askenazi and Michal Linial</i>	
10:50 AM	Coffee break	

From large data to small networks

Rainer Spang

University of Regensburg, Germany

Rainer Spang is professor and chair of statistical bioinformatics at the University of Regensburg, Germany. His research focuses on modeling gene expression and signal disruption in human cancers. More recently, he addressed problems of reconstructing cellular information flow from molecular perturbation experiments.

Corrections

The speaker for *TT8—DNA Sequencing with Illumina Instruments and Chemistry: Current and Future Applications* will be Klaus Maisinger.

Poster A-67 was withdrawn at the request of the authors.

Poster B-35: the last author is Posada D.

Poster I-38: the last author is Leys N.