

“RNA-seq in all flavors” Mini-symposium
Lausanne Genopode Auditorium C
Friday June 10, 2011

Program

14:00 Welcome address, **Ioannis Xenarios**, SIB/Swiss-Prot

14:10 **Keith Harshman**, UNIL
«Different flavors of RNA-seq library preparation»

14:35 **Jacques Rougemont**, EPFL
«Quantifying the expression of retro-elements with RNA-seq»

15:00 **Simon Anders**, EMBL
«Comparative analysis of RNA-Seq data»

15:45 *coffee break*

16:15 **Anamaria Necșulea**, UNIL
«Rapid evolution of alternative splicing in mammals revealed with RNA sequencing»

16:40 **Yvan Wenger**, UNIGE
«Transcriptome analysis of early hydra regeneration»

17:05 **Alexandre Reymond**, UNIL
«HTS to understand expression changes and the mechanisms beyond»

17:30 *end of meeting*



Organizers: Sylvain Pradervand, Keith Harshman, Christian Iseli, Jacques Rougemont, Ioannis Xenarios

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