

Thursday, **November 28, 2013**

Max Delbrück Communications Center, **Axon 1**

MDC Berlin-Buch, Robert-Rössle-Str. 10, **13125 Berlin**

# SYMPOSIUM

## GROUP LEADER FOR A SCIENTIFIC BIOINFORMATICS PLATFORM

10:00 - 10:10	<b>Nikolaus Rajewsky</b> , MDC Berlin	Introduction
10:10 - 10:40	<b>Altuna Akalin</b> , FMI Basel	Genome-wide epigenetic dysregulation in subtypes of acute myeloid leukemia
10:40 - 11:10	<b>Juliane Dohm</b> , CRG Barcelona	Genome reconstruction and characterisation using high-throughput sequencing
11:10 - 11:30	Coffee Break	
11:30 - 12:00	<b>Denis Bauer</b> , CSIRO Australia	Assessing the target specificity of triplex forming oligonucleotides using DNA microarrays
12:00 - 12:30	<b>Peter Frommolt</b> , University of Cologne	Computational models for networks of molecular regulation in <i>C. elegans</i>
12:30 - 13:15	Lunch Break	
13:15 - 13:45	<b>Kristin Reiche</b> , Helmholtz Centre Leipzig	Long non-protein coding RNAs regulated by cell cycle, oncogenic and tumor suppressor pathways
13:45 - 14:15	<b>Konrad Förstner</b> , University of Würzburg	From RNA-Seq data towards an understanding of pathogenicity regulation