



## SCHEDULE

09:20-09:30 Introduction

09:30-10:10 **Rob KLOSE**

*Department of Biochemistry, Oxford University, Oxford UK*

"Epigenetic conservation in vertebrates revealed by non-methylated DNA profiling"

10:10-10:50 **Gregor GILFILLAN**

*Norwegian Sequencing Centre, Oslo NO*

"ChIP-seq from low cell numbers: challenges and opportunities"

10:50-11:20 Coffee break

11:20-11:40 **Shankar PATTABHIRAMAN**

*Functional genomics & cancer, Institut de Génétique et de Biologie Moléculaire et Cellulaire (IGBMC), Illkirch FR*

"Linear DNA Amplification: recent developments"

11:40-12:00 **Marco-Antonio MENDOZA-PARRA**

*Functional genomics & cancer, Institut de Génétique et de Biologie Moléculaire et Cellulaire (IGBMC), Illkirch FR*

"A quality control system for the certification of ChIP-seq and enrichment-related massive parallel sequencing generated profiles"

12:00-12:40 **Michael KERTESZ**

*Technology Development, Illumina Inc, San Diego (CA) US*

"Molecule Sequencing for Human Genome Phasing and De-Novo Assembly of Complex Genomes"

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ACTIVE MOTIF

Enabling Epigenetics Research

CONDA

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