



BARCELONA NGS'16: GENOME ANNOTATION

APRIL 4-6, 2016 BARCELONA, SPAIN

Mark Your Calendars for the NGS 2016!

This year's conference will focus on cutting-edge methods for genome annotation. It will bring together researchers interested in sequencing and annotation of new reference genomes using NGS technologies. The meeting will include emerging and future trends in genomics including eco-system analysis and non-model genomes.

CONFERENCE TOPICS:

- Novel software and algorithms.
- New visualisation techniques.
- New applications of NGS technologies in genomics and biology.
- Post genomic sequencing and analysis including RNASeq, ribosome profiling, histone modifications, nucleosome positioning and proteogenomics.
- Clinical and Translational applications.
- Applications in Agriculture and biotechnology.
- Annotation of non-reference genomes.
- Genome Annotation impact on Health.

KEYNOTE SPEAKERS

Dave Burt, Roslin Institute, Edinburgh, UK

Rachel Karchin, John Hopkins University, Baltimore, US

Eric Karsenti, European Molecular Biology Laboratory, Heidelberg DE

Suzi Lewis, Lawrence Berkeley National Laboratory, Berkeley US

Peter F. Stadler, University of Leipzig, DE

KEY DATES

- Paper/ oral and poster submissions open

16 November 2015

- Submission closes

18 January 2016

- Early registration closes

29 February 2016

CONFERENCE CHAIRS

Jennifer Harrow, Wellcome Trust Sanger Institute, Hinxton, Cambridgeshire UK

Cedric Notredame, Centre for Genomic Regulation (CRG), Barcelona ES

website: www.iscb.org/ngs2016