

Opening Symposium June 13-14 2013

Aarhus & Foulum, Denmark



# Thursday June 13<sup>th</sup>, Aarhus

### 9:30 Registration

- 10:00 Welcome & introduction to GenSAP Center Director Mogens Sandø Lund, Center for Quantitative Genetics & Genomics (QGG), Aarhus University (AU)
- 10:15 Session 1 (chaired by Peter Sørensen):

Understanding the genetic basis of complex traits Senior Researcher Peter Sørensen, QGG, AU

Charting the genotype-phenotype map: Lessons from Drosophila Professor Trudy Mackay, Dept. of Genetics, North Carolina State University, USA

## 11:15 Coffee break

Genome-wide association studies – lessons learned and future directions Senior Researcher Goutam Sahana, QGG, AU

GxE and epigenetics in plants Senior Researcher Torben Asp, Dept. of Molecular Biology & Genetics, AU

Connecting biological pathways and the genomics of complex traits PhD Student Stefan McKinnon Høj-Edwards, QGG, AU

- 12:45 Lunch
- 13:45 Session 2 (chaired by Luc Janss):

Genomic prediction models: towards use of full sequence and large populations *Senior Researcher Luc Janss, QGG, AU* 

The 1000 bull genomes project - reconstructing the past genetic history of cattle, and accelerating future gains

Research Leader Ben Hayes, Dept. of Primary Industries, Victoria, Australia

## 14:45 Coffee break

Genomic selection using QTL information Senior Scientist Didier Boichard, INRA, France



Visions and perspectives for genomic prediction in animals, plants and humans Professor Daniel Gianola, University of Wisconsin-Madison, USA

The benefits of haplotypes as predictors of genetic merit PhD Student Beatriz Castro Dias Cuyabano, QGG, AU

- 16:15 Poster session & reception
- 17:30 Bus to Viborg
- 19:00 Dinner at Golf Salonen, Viborg

## Friday June 14<sup>th</sup>, Foulum

- 9:00 Welcome Center Director Mogens Sandø Lund, QGG, AU
- 9:10 Session 3 (chaired by Christian Sørensen):

Assessment and optimization tools in genomic breeding programs Senior Researcher Christian Sørensen, QGG, AU

Genomic selection and the design of breeding schemes Professor Theo Meuwissen, University of Life Sciences, Norway

## 10:10 Coffee break

On the doorsteps of a new era in plant breeding Senior Scientist Christian Sig Jensen, DLF Trifolium

Making a world-class pig-breeding program even better with genomic selection Chief Geneticist Mark Henryon, Danish Pig Research Center

Medium to long-term effects of selection in finite locus models with non-additive effects EGS-ABG PhD Student Hadi Esfandyari, Wageningen University/QGG, AU

- 11:45 General assembly meeting lunch meeting for GenSAP partners
- 12:30 Lunch
- 13:30 PhD defense:

Genomic predictions using combined populations and SNP marker panels PhD Candidate Rasmus Froberg Brøndum, QGG, AU