

15 November 2010

Mini Seminar Genomic Selection

Date: Wednesday 24 November 2010 from 10.00 - 13.00

Location: Hotel Park Inn
Engvej 171
2300 Copenhagen S

Agenda: 10.00 Welcome by Lars-Inge Gunnarsson
10.05 Genomic predictions using 3K, 50K, 800K or
sequence data within and across populations –
results and speculations by Mogens Sandø Lund
10.45 Break
10.55 Genomic information in routine evaluation –
Genomic prediction, blending and publication
criteria by Gert Pedersen Aamand and Esa
Mäntysaari
11.35 Break
11.50 Genomic breeding plans
- Genotyping strategies using 3K and 50K
(Christian Sørensen)
- Reference population for 'new' and 'old'
traits (Line Buch & Morten Kargo)
- Practical recommendations for Sire
selection (Morten Kargo)
12.45 General discussion
12.55 Closing Meeting by Lars-Inge Gunnarsson

Language: English, chaired by Lars-Inge Gunnarsson

Questions: Please feel free to send questions about Genomic
Selection within breeding plans and breeding
evaluation or related questions to the head of your
breed, **before Monday 22 November 2010:** Lars
Nielsen, Lan@vikinggenetics.com; Auli Hiimanen,
auli.hiimanen@vikinggenetics.com; Lars-Olof
Bårström, lars-olof.barstrom@vikinggenetics.com;
Hans Stålhammar,
hans.stalhammar@vikinggenetics.com; Peter
Larsson, PGL@vikinggenetics.com.

Registration and costs

Registration for the meeting no later than 12 p.m. Monday 22
November 2010 to Ann Raaby, anr@vikinggenetics.com.

Sincerely yours

On behalf of the Steering Committee

Claus Fertin, VG / Mogens Sandø Lund, AU/Foulum /
Gert Pedersen Aamand, NAV / Søren Borchersen, SC