

Program

10.30 Welcome by Lars-Inge Gunnarsson

10.30 Size of reference group, different SNP sets (3K, 50K, 700K) by Mogens Sandø Lund and Freddy Fikse

11.05 EBVs with medium reliability

Ranking at different evaluation runs

Use of breeding values in practice

by Gert Pedersen Aamand

11.30 Break

11.40 Challenge in new breeding schemes with use of effective Genomic Selection by Anders C. Sørensen

12.10 The effect of single genes – genes with large effect and recessive lethals by Nina Schulman

12.30 Closing remarks Lars-Inge Gunnarsson

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