Program NAV workshop 14 January 2016 at Park Inn, Engvej 171, 2300 Copenhagen

Program

Until 9.30	Coffee

9.30 - 9.35   Welcome (Lars-Inge Gunnarsson)

9.35 - 9.45   NAV homepage – starting point for breeding information (Terhi Vahlsten, Emma Carlén and Anders Fogh)

9.45 - 10.10   Future possibilities to use AMS data for improving genetic evaluation
- Use of udder coordinates to improve udder conformation (Elisenda Rius-Vilarrasa and Jan-Åke Eriksson
- Possibilities to improve temperament with AMS data (Anders Fogh and Rasmus Skovgaard Stephansen)

10.10 - 10.30   NTM – the improvement process
- Youngstock survival and Functional udder: follow up after NAV webinar, status and new results
- Introduction to group work: Questions to answer/discuss in relation to
  1) Youngstock survival, 2) Functional udder (Holstein only), 3) An efficient cow
  (Emma Carlén, Anders Fogh, Jan-Åke Eriksson and Gert Pedersen Aamand)

10.30 - 12.00   Group work per breed with focus on:
- Final recommendation in relation to Youngstock survival
- Final recommendation in relation to functional udder (Holstein only)
- Open discussion in smaller groups about an efficient cow

12.00 - 12.50   Lunch

12.50 - 13.10   Main conclusions from group work

13.10 - 13.40   Breeding goal (Morten Kargo, Jørn Pedersen, Jehan Ettema)
- Improved tools which can be used for making an update/revision of NTM
- What can be learned from a breeding goal survey?

13.40 - 13.50   Cross breeding –update on results (Anders Fogh and Morten Kargo)
- Dairy x dairy crosses
- Dairy x beef crosses

13.50 - 14.05   Coffee break

14.05 - 15.15   Status and plans - genomic prediction and traditional evaluation (Ulrik Sander Nielsen, Lars Peter Sørensen, Jukka Pösö and Gert Pedersen Aamand)
- Weekly genomic prediction
- GEBV reliability,
- GEBV for young stock survival,
- Do realised EBVs full fill the expectations based on GEBV?
- Improvements in pipeline
- Fee system

15.15 – 15.30   Evaluation and closing of the workshop