

STØTTET AF

Promilleafgiftsfonden for landbrug

We would like to welcome you to the combined Robusta status- and RadiMax concluding meeting. In extension of the status meeting there will be a hands-on Robusta workshop the day after, which will host all persons who wish to have a demonstration on the in-silico tools available for marker- and genomic selection based breeding at genome.dk. We kindly ask you to email Lars Bonde Eriksen (lbe@seges.dk), what part of the program you would like to join:

- Status meeting 05/12-2019
- Dinner 05/12-2019
- Accommodation 05/12-2019
- Robusta Hands-on workshop 06/12-2019 morning

Program 5/12 Thursday

- 09:30 Welcome and overview of Robusta and RadiMax (Christian)
- 10:00 What did we learn so far? (Kristian)
- 10:30 Wheat 2018 phenotype data (RadiMax + field)
- 11:00 Wheat 2019 phenotype data
- 11:30 Wheat 2018/2019 trait heritability and genomic prediction
- 12:00 Potato 2018 data
- 12:30 Lunch
- 13:30 Barley phenotype data (Pernille Bjarup Hansen)
- 14:00 Barley multiomics (Anja Karine Ruud)
- 14:30 Grass 2017 phenotype and multiomics data (Marta Malinowska)
- 15:00 Grass 2019 phenotype data
- 15:30 Discussion and conclusions
- 16:00 End
- Guided Tour at DLF facilities
- 18:00 Workshop Dinner

Program 6/12 Thursday - Robusta genome.dk workshop

- 09:00 Introduction and overview of functions at genome.dk (Torben)
- 09:30 Different pipelines (Istvan Nagy and Anja Karine Ruud, all)
- 10:30 Different type of analyses (all)
- 12:00 Lunch