

Faculty of Agricultural Sciences Department of Animal Sciences

Prof. Dr. Henner Simianer Animal Breeding and Genetics

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Göttingen, 01.02.2011

Course Announcement

Methods for Assessing Molecular Diversity in Farm Animals April 4-8, 2011

- In the scope of PAG Module 0060 'Advanced methods in animal breeding and statistical genetics' -

Lecturers:

- Dr. Steffen Weigend Institute of Farm Animal Genetics, Friedrich-Loeffler-Institut; Neustadt-Mariensee
- Dr. Georg Haberer MIPS/IBIS Institute of Bioinformatics and Systems Biology, Helmholtz Zentrum München
- Prof. Henner Simianer Animal Breeding and Genetics Group, Department of Animal Sciences, Georg-August-University Göttingen

Course outline:

F	\pr∠	1	Introd	duction	. molecu	lar mar	kers.	Sanger	seauen	cina. a	lianment	(Weiaend)

- Apr 5 Next Generation Sequencing, Blast, de novo assembly, SNP detection (Haberer)
- Apr 6 Measures of genetic diversity, F statistics, distance measures, marker-based kinship (Weigend)
- Apr 7 Phylogenetic structure, tree building, model based clustering (Weigend)
- Apr 8 Linkage disequilibrium, estimation of effective population size, haplotype reconstruction and imputing, selection signatures and selective sweeps (Simianer)

All lectures will be accompanied by practical exercises with relevant software tools.

Place: Department of Animal Sciences, Albrecht-Thaer-Weg 3, Göttingen

Registration: Participation is free for students enrolled at Georg-August-University, external participants will be charged a small course fee to cover expenses. The number of participants is limited to 30, applications will be accepted in the order of receipt.

Please register before 21. 3. 2011 by email to tierzucht@agr.uni-goettingen.de

For further information please contact hsimian @gwdg.de