



Chairs - Dr. Robbie Waugh, James Hutton Institute, Dundee, *Scotland*, UK;

Dr. Mats Hansson, Lund University, SW

08:30 - 09:00	Prof. Dr. Nils Stein IPK Gatersleben, <i>Germany</i>	At the dark side of barley diversity - the gap between discovery and user- friendly access.
09:00 - 09:20	Asis Shrestha Heinrich-Heine-Universität Düsseldorf, <i>Germany</i>	The double round robin population unravels the genetic architecture of grain size in barley
09:20 - 09:40	Outmane Bouhlal ICARDA-CGIAR // Chouaib Doukkali University, <i>Morocco</i>	CGIAR BARLEY BREEDING TOOLBOX: A diversity panel to facilitate breeding and genomic research in the Developing World
09:40 - 10:00	Ronja Wonneberger The James Hutton Institute, <i>United Kingdom</i>	Comprehensive expression atlas of six tissues coupled with high density SNP data provides a resource for studies and characterization of European two-rowed spring barleys and identified a major haplotype switch on chromosome 5H