

Time	Speaker	Topic
Tuesday		Moderator: Katrin Heer
09:00	Katharina Budde	Fourteen years of Alpine Forest Genomics Network (AForGeN): Advances in the research field and perspectives ahead
09:20	Thibaut Capblancq	Assessing maladaptation risk in Alpine conifers
09:40	Thomas Francisco	Demographic history shapes forest tree vulnerability to climate change
10:00	Mark Riedi	National representativeness of Swiss gene conservation units for Swiss stone pine (<i>Pinus cembra</i>)
10:20	Break	
		Moderator: Katharina Budde
10:50	Jonathan Feichtner	Expansion load compromises local adaptation in a long-lived tree species
11:10	Samuel Christe	Genetic structure of bog pine, <i>Pinus mugo</i> ssp. <i>rotundata</i> , in the Black Forest.
11:30	Devrim Semizer-Cuming	Investigating the adaptive potential of <i>Abies alba</i> in southern Germany using a multi-method approach
11:50	Maria Höhn	Climate change hazards and Carpathian conifer forests
12:30	Lunch	
		Moderator: Jill Sekely
14:00	Lucian Curtu	Contrasting adaptation strategies of European beech provenances. A case study at the Eastern margin of its natural distribution range
14:20	Christian Lampei	Selection in European beech in seedlings and adult trees
14:40	Ivan Scotti	Gradients, gradients! Latest news about the genomics of microgeographic and regional adaptation in European beech
15:00	Niels Müller	Towards understanding the pangenomic basis of natural variation in European beech
15:20	Break	
		Moderator: Charalambos Neophytou
16:00	Felix Rentschler	Resistance rising? Early insights from newly established common ash progeny trials
16:20	Aglai Szukala	The role of introgression in shaping taxonomic, genetic, and adaptive diversity in Southeast European White Oaks
16:40	Agatha Hurel	Future potential of <i>Quercus cerris</i> in Austria under climate change: insights from phenotypes and genotype–environment associations
17:00	Christian Rellstab	Contrasting genetic and plastic effects on trait variability in sessile oak and European beech across networks of common gardens
Thursday		Moderator: Christian Rellstab
09:00	Feedback from Group discussions	
09:40	Jian Feng Mao	GraphMana and GraphPop: new computational platforms for population genomics
10:00	Felix Gugerli	A brief update on <i>Abies alba</i> reference genome progress
10:45	Break	
11:00	AForGenMeeting	