

iGRAD-Plant Retreat 2021 – Program

Day 1 – Wednesday, Oct 6, 2021

10:15	10:40	Registration - Heinrich-Heine Saal (2nd floor)		
10:40	11:00	Welcome – Andreas Weber		
11:00	11:20	Vanessa Reichel-Deland	Unraveling drought signal transduction of the facultative CAM plant <i>Talinum fruticosum</i> in orthogonal systems	Chair: Francesco
11:20	11:40	Mara Schultz	The role of UPF0016 members in manganese homeostasis of photoautotrophs	
11:40	11:50	Katarzyna Krawczyk	Peroxisomes of photosynthetic organs - modulate resource allocation during abiotic stress	
11:50	12:00	Ghazala Butt	Transcriptional regulation of systemic acquired resistance	
12:00	12:10	Jonas Schön	Quantitative understanding of the ethylene signaling network through synthetic reconstruction and optogenetic control in orthogonal mammalian cell systems	
12:10	13:25	Lunch Break		
13:30	13:50	Sajjad Ghaffarinasabsharabiani	The effect of metabolic network size in adaptability of unicellular organisms	Chair: Tim
13:50	14:10	Allegra Corelli Grappadelli	Mass spectrometrical imaging method development for studying cell wall polymers	
14:10	14:30	Isaia Vardanega	Identification and characterization of CLV receptors and CLE peptides regulating barley shoot meristem maintenance and development	
14:30	14:55	Coffee Break		
15:00	15:20	Welcome – Susanne Hoffmann-Benning & Andreas Weber		
15:20	15:50	Shin Han Shiu	IMPACTS - Integrated training Model in Plant and Computational Sciences	Chair: Ksenia
15:50	16:15	Michael Feldbrügge	Linking transport and translation of mRNAs with endosomes and mitochondria	
16:20	16:40	Coffee Break		
16:40		Vera Göhre	Living inside the host for years: the novel smut fungus <i>Thecaphora thlaspeos</i> as a model pathogen in model plants	Chair: Natascha
	17:20	Lesley Plücker	Mg ²⁺ transport and its role for virulence in the smut fungus <i>Ustilago maydis</i>	
17:20		Petra Bauer	Iron deficiency signaling networks and integration during the life of a plant	
	18:00	Dibin Baby	Can IRONMAN vanquish BRUTUS' acts? A saga of plant transcription factors in Fe acquisition pathway	
18:00	18:15	Closing Day 1		

Day 2 – Thursday, Oct. 7, 2021

11:00	11:10	Welcome	
11:10	11:30	Ksenia Trofimov	Analysis of subcellular organization within the iron uptake regulation in plants
11:30	11:50	Natascha Heßler	Sugar partitioning between the Brassicaceae smut fungus <i>Thecaphora thlaspeos</i> and its host plants
11:50	12:10	Chen Deng	Identification and characterization of a sugar allocation system in <i>Arabidopsis-B. Cinerea</i> interaction
12:10	13:25	Lunch Break	
13:30	13:50	Gesa Helmsorig	<i>HvLWD1</i> is a candidate gene underlying the <i>early maturity 7 (eam7)</i> locus in barley
13:50	14:10	Tianyu Lan	Effects of High Ambient Temperature on Inflorescence Development in Barley (<i>Hordeum vulgare</i>)
14:10	14:30	Francesco Cosenza	Combined single population and multi-parent population QTL analysis to detect loci involved in control of flowering time and plant height in barley
14:30	14:55	Coffee Break	
15:00	15:30	Danny Ducat	PhotoSynthetic: Using synthetic biology approaches to gain insight into natural metabolic processes in cyanobacteria
15:30		Shizue Matsubara	Thylakoid maintenance and remodeling — from a point of view of pigments
	16:10	Ana Carolina dos Santos Sá	Natural variation in sink-source interaction and carbon allocation in <i>Arabidopsis thaliana</i> under fluctuating environment
16:10		Oliver Ebenhöf	Quantitative models of photosynthetic acclimation
	16:50	Tim Nies	Investigating Photoinhibition by Computational Modelling
16:50	17:10	Coffee Break	
17:10	17:35	Matias Zurbruggen	Engineering photoreceptors into optogenetic tools for the control and understanding of cellular processes in microbial, animal and plant systems
17:35	18:05	Amanda Koenig & Martin Kulke	Examples of national and international collaborations: Use of molecular simulation and optogenetics to understand protein-lipid interactions in long-distance signaling
18:05	18:15	Closing Day 2	
18:45		BBQ	

Day 3 – Friday, Oct 8, 2021

09:45	13:00	Social Event: City Rallye – Team Challenge
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