1st French-German Maize Breeders School

Genetic diversity in maize breeding programs







May 23 and 24, 2017 | Stuttgart-Hohenheim, Germany





A.S.F.
Association des Selectionneurs Français





Scientific organizers |

Albrecht E. Melchinger, University of Hohenheim, Stuttgart Alain Charcosset, INRA, Gif sur Yvette

Hosts |

Association des Sélectionneurs Français (A.S.F.), Paris Deutsches Maiskomitee e.V. (DMK), Bonn Gesellschaft für Pflanzenzüchtung e.V. (GPZ), Göttingen Institut National de la Recherche Agronomique (INRA), Paris University of Hohenheim, Stuttgart

Date and location |

May 23 and 24, 2017

Christkönigshaus Stuttgart, Paracelsusstr. 89, 70599 Stuttgart

Participation fee |

A conference fee of EUR 150,00 per person will be charged, including lunch, reception with supper and refreshments during break (students and postgraduates EUR 15,00, without reception and supper). Participation is limited to a total number of 100 persons and registration will be accepted on a first order first serve basis.

Room reservation |

A limited number of rooms are available in the Christkönigshaus — Stuttgart for about 77 EUR per person/night. Please make your reservation directly at the Christkönigshaus under the heading "Deutsches Maiskomitee".

Tel.: +49 (0)711-4582823

email: christkoenigshaus@tagungshaus.net

Registration |

with the enclosed registration form by Fax: +49(0)228 92658-20 — email: dmk@maiskomitee.de

Terms of payment |

After receipt of your registration you will receive a confirmation and the invoice. If it is not possible for you to join the event despite a registration, you are entitled to name an alternative participant or cancel the registration (free of charge until May 8). For cancellations after May 8 we charge 50 % of the participant fee. Please make your cancellation in written form.

Closing date: May 15, 2017

Comparison of different "omics"

pressure in temperate maize

technologies for hybrid prediction

Matthias Westhues, University of Hohenheim, Germany

Sandra Unterseer, Technical University of Munich, Germany

Identification of genes under differential selective

May 23, 2017				
11:00 12:15	Registration Welcome addresses	Session 2 Session chair: Chris-Carolin Schön Tools and approaches for broadening the genetic diversity with genetic resources		
Carain	Helmut Meßner, Deutsches Maiskomitee e. V. (DMK), Germany Albrecht E. Melchinger, University of Hohenheim, Germany	15:40	Use of Doubled Haploids in the Context of Genetic Resources Exploitation Thomas Lübberstedt, Iowa State University, USA	
Session 1 Session chair: Alain Charcosset Management and broadening of genetic diversity in breeding programs		16:10	Libraries of doubled-haploid lines: a new tool for harnessing landraces	
12:30	Developing and deploying climate-resilient maize in the tropics: progress and prospects Bodupalli Prasanna, CIMMYT, Kenya	16:30	Albrecht E. Melchinger, University of Hohenheim, Germany Association mapping of agronomic and metabolic traits in DH libraries of landraces	
13:00	Diversity in a global breeding program: status and possibilities for broadening Rainer Leipert, KWS, Germany	16:50	Juliane Böhm, University of Hohenheim, Germany Finding the optimum level of diversity for germplasm improvement	
13:30	Lessons learned and expectations for broadening the diversity in elite breeding programs Tobias Eschholz, Maisadour, Germany	17:10	Manfred Mayer, Technical University of Munich, Germany GBS based comparison of USDA and INRA	
13:50	Breeding technologies help to describe and utilize genetic diversity in maize breeding		inbred lines collections Brigitte Gouesnard, INRA, France	
14:10	Carsten Knaak, KWS, Germany Managing genetic diversity in an elite breeding program Alain Murigneux, Limagrain, France	17:30 18:20	DNA pooling for evaluating the originality of landraces Stéphane Nicolas, INRA, France Departure evening program	
14:40	Practical aspects of introgression of landraces into elite pools: The Piemont-Pyreneen case study		Point of departure: Christkönigshaus, Paracelsusstr. 89, 70599 Stuttgart	
15:00	Cyril Bauland, INRA, France Coffee break	19:00– 22.00	Reception and supper with wine tasting "Collegium Wirtemberg", Uhlbacher Kelter Uhlbacher Str. 221, 70329 Stuttgart	
	7 24, 2017 n 3 Session chair: Antoine Gaillard	11:45	Assessment of genetic heterogeneity using	
genomi	ing the genetic diversity in mapping populations and c prediction		whole-genome regression models Christina Lehermeier, Technical University of Munich, Germany	
08:30	Multiparental QTL mapping of biomass related traits Adama Seye, INRA, France	12:10	Lunch	
09:00	Choice of individuals for the training set in genomic prediction Laurence Moreau, INRA, France	Miscel	n 4 Session chair: Susanne Groh laneous topics II	
09:30	Genomic selection in synthetics — results from a simulation study Pascal Schopp, University of Hohenheim, Germany	13.10	New developments in the doubled-haploid technology in maize Willem Molenaar, University of Hohenheim, Germany	
09:55	Recurrent genomic selection in synthetics Dominik Müller, University of Hohenheim, Germany	13:30	Evaluation of quality traits for biogas production in maize Wolfgang Schipprack, University of Hohenheim, Germany	
10:20 Session	Coffee break 1 4 Session chair: Milena Ouzunova	13:50	Transcriptomic dissection of heterosis manifestation in young maize roots Frank Hochholdinger, University of Bonn, Germany	
Miscellaneous topics I		14:20	Variation for and inheritance of BYDV	

resistance in maize

Alain Charcosset, INRA, France

14:50 Concluding remarks

15:00 End of meeting

Benjamin Stich, University of Düsseldorf, Germany

Registration form

Fax: +49(0)228 92658-20

email: dmk@maiskomitee.de

Deutsches Maiskomitee e. V. (DMK) Brühler Straße 9 53119 Bonn | Germany

Current topics in maize breeding

May 23 and 24, 2017 at Christkönigshaus Stuttgart

I will participate at the event.

Please return until May 15, 2017

Name	
Organization	
Billing address	
Email address	
☐ I am student or postgraduate (15,00 € without reception and supper. Ple	ase add relevant certificate).
Signature	

Send Email