



WHEAT WORLD CONGRESS

Workshop 10: Functional Genomics

Sunday, 22 September 2024

Perth Conference & Exhibition Centre (PCEC), Perth, Western Australia

Meeting Room 2, PCEC

Tentative Program

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Time	Co-Chairs: Prof Harvey Millar, The University of Western Australia, Australia Dr Phillipa Borrill, John Innes Centre, UK	
13:00 – 13:10	Introduction	
13:10 – 13:30	The discovery of tandem kinase r-genes: origin, function, and potential in resistance breeding	Prof Tzion Fahima University of Haifa, Israel
13:30 – 13:50	Unveiling the micronutrient architecture of wheat grains through spatial transcriptomics at single-cell resolution	Dr Chen Ji John Innes Centre, UK
13:50 – 14:10	Deciphering the role of small RNAs in regulation of restorer-of-fertility gene expression in wheat	Dao Anh Thien Tran The University of Western Australia, Australia
14:10 – 14:30	Developing cisgenic resistance gene stacks for improved resistance to wheat stem rust disease	Dr Ming Luo Commonwealth Scientific and Industrial Research Organisation (CSIRO), Australia
14:30 – 14:50	Tea/Coffee break	
14:50 – 15:10	Nicotianamine synthase gene discovery, characterisation, and overexpression in bread wheat	Dr Jesse Beasley The University of Melbourne, Australia
15:10 – 15:30	Does-response assays and transcriptomic analyses evaluate novel fungicide-supplement combinations on elevating wheat rust disease resistance	Dr Meng Li University of British Columbia, Canada

15:30 – 15:50	Identification of wheat rDNA unit variants and changes in their expression following partial chromosomal deletions	Mr Yohta Hyuga Prefectural University, Kyoto, Japan
15:50 – 15:55	Closing remarks by Co-Chairs	