

AT-A-GLANCE PROGRAM

Oral presentations are in Grand Ballroom. Reception/Poster sessions are outside the Grand Ballroom. All meals are in Ramada Plaza Taian.

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| Monday April 9th | | |
| 8:30 - 9:00 | Registration | |
| 9:00 - 9:05 | Welcome | |
| 9:05 - 10:25 | Oomycete biology, populations, and evolution I | Chair: Audrey Ah-Fong and Kai Tao |
| 9:05 - 9:20 | Cellular signaling and cytoskeleton dynamics in <i>Phytophthora</i> | Francine Govers |
| 9:25 - 9:40 | Biochemical and genetic analyses of the oomycete <i>Pythium insidiosum</i> provide new insights into clinical identification, pathogenicity, and evolution of metabolism-related traits | Theerapong Krajaejun |
| 9:45 - 10:00 | Mating type change in <i>Phytophthora infestans</i> regulated by histone deacetylases | Li-Yun Guo |
| 10:05 - 10:20 | Ecology and evolution of oomycete communities in response to soybean seed treatments | Zachary Noel |
| 10:25 - 11:00 | Group photo and break | |
| 11:00-12:20 | Keynote Speech | Chair: Francine Govers |
| 11:00 - 11:35 | Exploring fungal DNA Virus to control crop diseases, challenge and opportunity | Daohong Jiang |
| 11:40 - 12:15 | Omics in Plant Breeding | Sanwen Huang |
| 12:20 - 2:00 | Lunch | |
| 2:00 - 3:40 | Effectors I | Chair: Petra Boevink and Kelsey Wood |
| 2:00 - 2:15 | Dissecting the transcriptional plasticity of <i>P. sojae</i> | Brett Tyler |
| 2:20 - 2:35 | Identification of <i>Plasmopara viticola</i> candidate RXLR effector repertoire in <i>Nicotiana benthamiana</i> | Jiang Lu |
| 2:40 - 2:55 | How are effectors secreted and delivered to the host? | Shumei Wang |
| 3:00 - 3:15 | Plant recognition of <i>Phytophthora</i> GH12 MAMPs | Yan Wang |
| 3:20 - 3:35 | Characteristics of RxLR effector structures in oomycete pathogens | Li Zhao |

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| 3:40 - 4:10 | Break | |
| 4:10 - 6:10 | Genomics I | Chair: Paul Birch and Diya Sen |
| 4:10 - 4:25 | Towards the identification of effectors and secondary metabolites involved in oomycete-oomycete interactions and mycoparasitism using comparative genomics | Laura Grenville-Briggs |
| 4:30 - 4:45 | Comparative genomics of four species of <i>Peronosclerospora</i> | Kyle Fletcher |
| 4:50 - 5:05 | What role does <i>Phytophthora kernoviae</i> play in New Zealand forests? | Rebecca McDougal |
| 5:10 - 5:25 | Land-use affects the growth and gene expression of <i>Phytophthora agathidicida</i> | Preeti Panda |
| 5:30 - 5:45 | Transcriptional regulation of <i>Phytophthora sojae</i> effectors for successful infection | Wenwu Ye |
| 5:50 - 6:05 | Genome sequence of <i>Plasmopara viticola</i> reveals effector repertoire and pathogenicity mechanisms | Ling Yin |
| 6:10 - 7:30 | Dinner | |
| 7:30 - 10:00 | Reception/Poster session | |

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| Tuesday April 10th | |
| 9:00 – 5:00 | Excursion/free time |
| 6:00 - 7:30 | Conference Dinner |

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| Wednesday April 11th | | |
| 8:30 - 9:50 | Host Interactions and Resistance Mechanisms I | Chair: Jun Zhao and Veronica Ancona |
| 8:30 - 8:45 | <i>Phytophthora palmivora</i> establishes tissue-specific intracellular infection structures in the earliest divergent land plant lineage | Sebastian Schornack |
| 8:50 - 9:05 | A pathogen suppressor of an NLR immune network | Lida Derevnina |
| 9:10 - 9:25 | A plant pathogen effector utilizes host susceptibility factor NRL1 to degrade the immune regulator SWAP70 | Qin He |
| 9:30 - 9:45 | A potato STRUBBELIG-RECEPTOR FAMILY gene StLRPK1 involves in plant immunity by association | Haixia Wang |

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| | with StSERK3/BAK1 | |
| 9:50 – 10:20 | Break | |
| 10:20-12:00 | Effectors II | Chair: Suomeng Dong and Lida Derevnina |
| 10:20 - 10:35 | <i>Phytophthora infestans</i> RXLR Effector Pi22926 suppresses plant immunity by targeting the host MAP3 β to manipulate MAP3K signalling pathway | Zhendong Tian |
| 10:40 - 10:55 | Biochemical basis of an RXLR-WY effector suppression of an NLR network | Jessica Upson |
| 11:00 - 11:15 | Characterization of host-recognized WY domain effectors from <i>Bremia lactucae</i> that lack the canonical RxLR motif | Kelsey Wood |
| 11:20 - 11:35 | Functional analysis of an effector PcRxLR101 in <i>Phytophthora capsici</i> | Jing Li |
| 11:40 - 11:55 | A potato chloroplast kinase is required for Rpi-vnt1 mediated immunity against late blight pathogen | Chuyun Gao |
| 12:00 - 2:00 | Lunch | |
| 2:00-3:20 | Oomycete biology, populations, and evolution II | Chair: Daolong Dou and Theerapong Krajaeun |
| 2:00 - 2:15 | Molecular genetics of the oomycete pathogen of lettuce, <i>Bremia lactucae</i> . | Lin Zhang |
| 2:20 - 2:35 | Structure determination of three enzymes from <i>Phytophthora capsici</i> | Weiwei Song |
| 2:40 - 2:55 | Profiling of the <i>Phytophthora</i> epigenomes | Han Chen |
| 3:00 - 3:15 | Immunity triggered by the small cysteine rich effector PC2 from <i>Phytophthora infestans</i> requires proteolytic cleavage by the host protease P69B | Shuaishuai Wang |
| 3:20 - 3:50 | Break | |

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| 3:50-5:30 | Genomics II | Chair: Sucheta Tripathy and Wenwu Ye |
| 3:50 - 4:05 | New insights into pathogenicity of the emerging tropical pathogen: <i>Phytophthora colocasiae</i> on taro | Diya Sen |
| 4:10 - 4:25 | Mutations in oxysterol binding protein-related protein conferring oxathiapiprolin resistance confirmed using CRISPR/Cas9 in <i>Phytophthora capsici</i> and <i>P. sojae</i> | Jianqiang Miao |
| 4:30 - 4:45 | Detection of multiple oomycetes in metagenomic data by using E-probe Detection of Nucleic Analysis (EDNA) | Maria Fernanda Proano |
| 4:50 - 5:05 | EumicrobeDBLiteV11: a lightweight genomic resource and analytic platform for draft oomycete genomes | Arijit Panda |
| 5:10 - 5:25 | Diploid genome assembly works better when Pacbio long reads are supplemented with Illumina reads: <i>P. ramorum</i> assembly a case study | Mathu Malar C |
| 5:30 - 6:20 | Committee meeting | |
| 6:30 - 7:30 | Dinner | |
| 7:30 - 10:00 | Reception/Poster session | |

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| Thursday April 12th | | |
| 8:30 - 9:50 | Host Interactions and Resistance Mechanisms II | Chair: Rebecca McDougal and Qin He |
| 8:30 - 8:45 | Synergistic anti-oomycete effect of melatonin with a biofungicide against oomycetic blank shank disease | Maozhi Ren |
| 8:50 - 9:05 | <i>Phytophthora</i> utilizes an RxLR effector to hijack the host ER stress as part of their infection strategy | Maofeng Jing |
| 9:10 - 9:25 | A conserved host target of oomycete Avr3a family effectors positively regulates plant resistance to <i>Phytophthora</i> | Ruirui Feng |
| 9:30 - 9:45 | NbMORF8 encoding protein localized in mitochondria and chloroplasts negatively regulates plant immunity to <i>Phytophthora</i> pathogens | Yang Yang |
| 9:50 - 10:20 | Break | |

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| 10:20 - 11:40 | Effectors III | Chair: Jiang Lu and Biao Gu |
| 10:20 - 10:35 | The evolutionarily conserved Phytophthora effector RxLR24 interferes with the secretion of antimicrobial compounds | Iga Tomczynska |
| 10:40 - 10:55 | Structure and function analysis of effector PcRxLR145 of Phytophthora capsici | Liyang Wang |
| 11:00 - 11:15 | The Phytophthora sojae essential effector Avh238 targets soybean GmACSs to suppress ethylene biosynthesis and promote infection | Bo Yang |
| 11:20 - 11:35 | Host mRNA methylation pathway, a new target for Phytophthora effector proteins? | Jie Huang |
| 11:40 -12:00 | Closing remarks | |
| 12:00 -2:00 | Lunch/Adjourn | |