

**August 16, 2018 - Thursday**

08:45-09:00	<b>COLLOQUIUM 2</b>
09:00-09:15	
09:15-09:30	
09:30-09:45	
09:45-10:00	
10:00-10:15	
10:15-10:30	
10:30-11:00	<b>Coffee Break / Poster Presentations</b>
<b>Session Chair: Dr. Elisabeth Dirlewanger and Dr. Vivian Panes</b>	
11:00-11:15	<b>Dr. Jun Song</b> An integrated `omics` approach to improve and maintain quality of apples
11:15-11:30	<b>Dr. Vivian Panes</b> Discovering Genes Involved in the Synthesis of Secondary Metabolites from the Seeds of Moringa oleifera Lam. through Transcriptome Analysis
11:30-11:45	<b>Dr. Elisabeth Dirlewanger</b> New sweet cherry genomic tools and their use in marker-assisted breeding
11:45-12:00	<b>Dr. SINEM ÖZTÜRK ERDEM</b> Genetic Relationship Between Some Strawberry Varieties and F1 Population Obtained from These Varieties
12:00-12:15	<b>Mr. William Nak</b> Temporary silencing targeted plant genes with exogenous application of dsRNA
12:15-12:30	<b>Madeleine Ms. Gleeson</b> From the Outside-In: topical application of microRNAs to avocado
12:30-14:00	<b>Lunch / Poster Presentations</b>
<b>Session Chair: Prof. Shenchun Q u and Prof. Dr. Bayram Cevik</b>	
14:00-14:15	<b>Ryutaro Tao</b> <b>Keynote speaker</b>
14:15-14:30	
14:30-14:45	<b>Prof. Shenchun Q u,</b> MicroRNA0137 Targeting to a AP2/ERF Family Transcription Factor ERF14 Mediates the Resistance against Botryosphaeria dothidea Infection in Malus
14:45-15:00	<b>Dr. Andres Cruz-Hernandez</b> miRNAs participation in prickly pear (Opuntia sp) fruit development



15:00-15:15	<b>Evelyne Costes</b> Deciphering the roles of dormancy-associated and flowering-time related MADS-box transcription factors during bud dormancy in apple
15:15-15:30	<b>Prof. Dr. Bayram Cevik</b> Transcriptional analysis of tomato-interactions with RNA viruses
15:30-16:00	<b>Coffee Break / Poster Presentations</b>
<b>Session Chair: Prof. Hongqiang Yang</b>	
16:00-16:15	<b>Dr. Panagiotis Kalaitzis</b> A prolyl 4 hydroxylase plays a role in the regulation of the tomato ( <i>Solanum lycopersicum</i> ) fruit growth program
16:15-16:30	<b>Mr. Kentaro Ono</b> Characterization of the pollen-part modifier gene involved in self-incompatibility reaction in sweet cherry
16:30-16:45	<b>Assist. Prof. Sherif Sherif</b> Abscisic acid-mediated transcriptional regulation of dormancy- associated MADS-box genes during peach bud dormancy
16:45-17:00	<b>Minori Sonoda</b> Mechanism of male organ differentiation mechanism in the sex determination system of kiwifruit
17:00-17:15	<b>David Lewis</b> Flower colour patterns in Cymbidium orchid linked with expression of a Myb transcription factor during flower development
17:15-17:30	<b>Prof. Hongqiang Yang</b> Transcriptome comparison among three types of annual roots of <i>Malus hupehensis</i> Rehd
17:30-17:45	<b>Prof. Dr. Kexuan Tang</b> Overexpression of blue light receptor AaCRY1 improves artemisinin content in <i>Artemisia annua</i> L
17:45-18:00	<b>Prof. Dr. Rosario Muleo</b> White colour is generated by a down-regulation of genes and metabolic pathway of flavonoids

