

Poster Board Number	Title	Presenting author	Affiliation	Country
PS3.17.158	<i>Age-associated remodelling of the gut microbiome is linked to a decline in host filtering</i>	Nivedhitha Ulaganathan	Indian Institute of Science	India
PS3.17.159	<i>Bio-enhanced seed pellets improve early establishment of Acacia confusa under simulated landslide soil conditions</i>	Wen-Ching Chen	National Chung Hsing University	Taiwan, Province of China
PS3.17.160	<i>Can seagrasses drive bacterial succession within a meadow?</i>	Rebecca Case		Singapore
PS3.17.161	<i>Capacity for luxury uptake and storage of nitrogen in cyanobacteria</i>	Nagasaijanani Rajpirathap	University of Auckland	New Zealand
PS3.17.162	<i>Co-resident plasmid interactions reshape extracellular vesicle-mediated gene transfer across bacterial hosts</i>	Uli Klümper	TU Dresden	Germany
PS3.17.163	<i>Emergent coexistence is common in mathematical models of microbial consumer-resource systems</i>	James Richardson		Australia
PS3.17.164	<i>How thermodynamic limitations shape metabolic strategies in anaerobes</i>	Christoff Odendaal	Delft University of Technology	Netherlands
PS3.17.165	<i>Just keep swimming - uncovering the principles governing regulatory network rewiring</i>	Mitchell Reynolds		United Kingdom
PS3.17.166	<i>Matrix-driven architecture and resilience in Salmonella biofilms</i>	Thibault Debord		Belgium
PS3.17.167	<i>Multiplicity Quantifies Intra-Cluster Diversity in Metagenomic Analysis</i>	Felipe González-Casabianca		Estonia
PS3.17.168	<i>Potential Contribution of Microbial Interaction to Stochasticity of Microbiota Assembly</i>	Zhong Yu	Sun Yat-Sen University	China
PS3.17.169	<i>Predicting spatial structures of microbial communities by explicitly accounting for facilitative and inhibitory interactions</i>	Bharti Yadav		India
PS3.17.170	<i>Predicting the worst-case timing of multiple pulse perturbations</i>	James Orr		Australia
PS3.17.171	<i>Quantifying stochasticity among replicated communities in experimental microbial community assembly</i>	Ibuki Hayashi	Kyoto University	Japan
PS3.17.172	<i>Reconstructing the infection dynamics of R. solanacearum by tracing bacterial lineages with neutral genetic markers</i>	Sylvain Vicente	Laboratoire des Interactions Plante-Microbes-Environnement (LIPME, INRAE, France)	France
PS3.17.173	<i>Shaking Regimes Influence the Growth of Human Gut Microorganisms</i>	Adedamola Godwin Daodu	KU Leuven	Belgium
PS3.17.174	<i>The Power-Law of Species-Time Relationships in Microbial Ecology</i>	Kai Feng	Research Center for Eco-Environmental Sciences, Chinese Academy of Sciences	China