

Poster Board Number	Title	Presenting author	Affiliation	Country
PS2.29.395	<i>Agricultural intensification factors reduce the diversity of rare soil bacteria and fungi: insights from a microbiome meta-analysis</i>	Purbasa Kole	North Dakota State University	United States of America
PS2.29.396	<i>Assessing the impacts of peatland restoration on microbial communities in Northumberland, UK</i>	Isis Cohen Nicholson	Northumbria University	United Kingdom
PS2.29.397	<i>Augmenting and stimulating microbiomes: seed and soil based strategies for native grass revegetation</i>	Sasha Tetu	Macquarie University	Australia
PS2.29.399	<i>Beyond genomic prediction: experimental validation of oxidative stress responses in coral probiotic bacteria</i>	Luanny Martins Fernandes		Australia
PS2.29.400	<i>Bin there, done that: Evaluating metagenomic binning pipelines to recover novel MAGs</i>	Ben Kilbrei		Canada
PS2.29.401	<i>Bioprospecting Endophytic Plant Growth-Promoting Bacteria Associated with Rhizophora mangle L. in a Brazilian Mangrove</i>	Douglas Monteiro		Brazil
PS2.29.402	<i>Characterising the ectomycorrhizal-microbial-soil nutrition axis supporting urban Dipterocarp species</i>	Sarah Geiger		Singapore
PS2.29.403	<i>Conservation preconditioning facilitates microbial rewilding of gut microbiome structure and function in endangered black-footed ferrets</i>	Sally Bornbusch	Smithsonian National Zoo and Conservation Biology Institute	United States of America
PS2.29.404	<i>Diet-microbiome covariation across three giraffe species in a close-contact zone</i>	Elin Videvall	Uppsala University	Sweden
PS2.29.405	<i>Flavobacteria mitigate heat-induced photoinhibition of coral endosymbiotic algae by restricting light exposure</i>	Toshiyuki Takagi	The University of Tokyo	Japan
PS2.29.406	<i>Hidden eukaryotic diversity and risk unveiled by high-accuracy long amplicons sequencing</i>	Xuemei Mao		Hong Kong
PS2.29.407	<i>Making your microbial biodiversity data reusable, citable, and visible</i>	Tobias Frøslev		Denmark
PS2.29.408	<i>Multi-Scale Evidence for Declining Microbial Carbon Fixation Along Forest Succession Gradients</i>	Shu-Yi-Dan Zhou	Chinese Academy of Sciences	China
PS2.29.409	<i>Reproductive microbiomes are associated with reproductive metrics in threatened felids and domestic cats</i>	Katherine Dami	University of Michigan	United States of America

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PS2.29.410	<i>Seasonal Variation of Phyllosphere Microbial Communities Under Warming</i>	Chaotang lei		China
PS2.29.411	<i>Separating direct thermal stress from microbiome-mediated effects on juvenile kelp growth</i>	Alice Simon	University of British Columbia	Canada
PS2.29.412	<i>Soil microbiomes reveal hidden trajectories of forest restoration beyond vegetation surveys</i>	Andrew Dopheide	Manaaki Whenua – Landcare Research group	New Zealand
PS2.29.413	<i>Strain-level differentiation of Escherichia coli, the dominant gut microbiome member in the critically endangered kākāpō</i>	Annie West	The University of Edinburgh	United Kingdom
PS2.29.414	<i>Symbiont Digestion under Nutrient Limitation</i>	Jade Shivak		New Zealand