

Poster Board Number	Title	Presenting author	Affiliation	Country
PS1.03.163	<i>120,000 new species from across the globe enrich the tree of life</i>	Samuel Aroney	Queensland University of Technology	Australia
PS1.03.164	<i>Adapting the GTDB workflow for Fungi</i>	Maria Chuvochina	Aalborg University	Denmark
PS1.03.165	<i>Comparative genomics and experimental validation of hurdles along the path to foreign gene domestication</i>	Tess Brewer	Eawag	Switzerland
PS1.03.166	<i>Discovery of novel Rhodothermota lineages in marine sponges: Phylogenomic diversification and functional adaptation to a host-associated lifestyle</i>	André C. Pereira	Institute for Bioengineering and Biosciences	Portugal
PS1.03.167	<i>Distinctive Evolutionary Patterns in Archaeal NRPS C-Domain Superfamily Revealed by Large-Scale Genome Mining</i>	Rei C. Abe	Institute of Science Tokyo	Japan
PS1.03.169	<i>Integrative taxonomic reassessment of Penicillium in marine environments</i>	Ji Seon Kim	Seoul National University	Korea, Republic of
PS1.03.170	<i>Molecular Detection and Phylogenetic Analysis of Macrorhabdus ornithogaster in Companion and Zoo Birds in Japan</i>	Naoki Kobayashi		Japan
PS1.03.171	<i>Potential immune mechanisms exhibited by oyster Magallana bilineata in preventing viral-mediated horizontal gene transfer</i>	Juri Nazareth Ochotorena	University of the Philippines Diliman	Philippines
PS1.03.172	<i>Quartet Ensemble Discordance</i>	Russell Neches	Kyoto University	Japan
PS1.03.173	<i>Spatiotemporal genomic catalog of the Han River: reconstruction of 2,229 metagenome-assembled genomes</i>	Do-Hoon Lee	Chung-Ang University	Korea, Republic of
PS1.03.174	<i>Stop codon reassignment to tryptophan in members of the bacterial phylum Actinomycetota</i>	Donovan Parks		Canada
PS1.03.175	<i>Uncovering the Phylogenetic Structure and Ecological Traits of the Bifidobacterium catenulatum-Related Lineages</i>	Kento Orihara		Japan