

Poster presentations

Poster no.	Presenter	Affiliation	Country	Category	Title
P-1	Biwen Wang	U Amsterdam	NL	Evolution and mobile genetic elements	Transposon insertion sequencing of a minimal <i>Bacillus subtilis</i> genome
P-2	Biwen Wang	U Amsterdam	NL	Biotechnology	Induction of CtsR-regulated chaperones improves xylanase production in <i>Bacillus Gintool</i> : Transcriptome analysis using regulation directionality information
P-3	Biwen Wang	U Amsterdam	NL	Others	Lethal ROS production upon membrane depolarization of dormant <i>Bacillus subtilis</i>
P-4	Biwen Wang	U Amsterdam	NL	Antimicrobials and toxins	Development of probiotic agents containing <i>Bacillus natto</i> against <i>Campylobacter</i> food poisoning
P-5	Ryosuke Kadoya	Sugiyama Jogakuen	JP	Antimicrobials and toxins	Aberrant <i>Bacillus subtilis</i> cell's morphology emerges as consequence of active lysogeny
P-6	Valentina Andrea Floccari	U Ljubljana	SI	Microbial interaction	Tracking Bacilli swimmers in microalgae
P-7	Romain Briandet	INRAE Jouy	FR	Microbial interaction	SPbeta-like viruses trigger changes in <i>Bacillus subtilis</i> behaviour under defined environmental conditions
P-8	Virginie Grosboillot	U Ljubljana	SI	Microbial interaction	The potential of membrane vesicles as immunostimulant modulating <i>Clostridioides difficile</i> colonization
P-9	Yotaro Isamu	Tsukuba U	JP	Microbial interaction	Isolation and characterization of bacterial-derived extracellular membrane vesicles in Kin discrimination and cooperative behaviours in <i>Bacillus subtilis</i>
P-10	Miku Matsushita	Tsukuba U	JP	Microbial interaction	Secreted nuclease facilitates extracellular DNA (eDNA) repurposing during biofilm
P-11	Polonca Stefanic	U Ljubljana	SI	Microbial interaction	Construction and directed evolution of <i>B. subtilis</i> synthetic consortia
P-12	Stephen Landier	Northwestern U	USA	Microbial interaction	Luciferase luminescence of colonies to assess NADPH levels
P-13	Anne-Gaëlle Planson	INRAE Jouy	FR	Microbial interaction	in <i>Bacillus subtilis</i> cells
P-14	Yuzheng Wu	Kobe U	JP	Biotechnology	Metabolic Engineering Design Based on Flux Balance Analysis to Improve
P-15	Nunthaphan Vikromvarasiri	RIKEN	JP	Biotechnology	Bio-production from Glycerol in <i>Bacillus</i>
P-16	Koki Tanaka	Tokyo Agri U	JP	Biotechnology	Analysis of segmented filamentous bacteria genome function in <i>Bacillus subtilis</i>
P-17	Katsumi Amano	Tokyo Agri U	JP	Biotechnology	Introduction and heterologous expression of heliobacterial
P-18	Satoru Watanabe	Tokyo Agri U	JP	Biotechnology	photosynthetic gene cluster in <i>Bacillus subtilis</i>
P-19	Takahiro Morita	Tokyo Agri U	JP	Biotechnology	Gene expression profile of <i>CyanoBacillus</i> , carrying chimeric genome of <i>Bacillus subtilis</i> and cyanobacterium <i>Synechocystis</i> sp. PCC
P-20	Hiroataka Matsubara	Amano enzyme	JP	Biotechnology	Development of the genetic engineering methods of <i>Cyanobacillus</i>
P-21	Junko Yamamoto	Shinshu U	JP	Biotechnology	Genome analysis of <i>Bacillus amyloliquefaciens</i> industrial strains and its application to heterologous protein
P-22	Taiki Kanzaki	Shinshu U	JP	Biotechnology	establishment of gene introduction into alkaliphilic bacteria using conjugation system consisting with Type IV secretion system (T4SS) and conjugative factor RP4
P-23	Claudia Borgmeier	BRAIN Biotech	DE	Biotechnology	Determination of the oriT minimum region in conjugation
P-24	Wakana Suda	Tokyo Agri U	JP	Biotechnology	between <i>Escherichia coli</i> and <i>Bacillus subtilis</i> . Modulation of a built environment microbiome by a <i>B. subtilis</i> strain
P-25	Rina Nogami	Kobe U	JP	Biotechnology	Effect of restriction/modification on plasmid transfer in <i>Bacillus subtilis natto</i>
P-26	Thomas Konjetzko	FZ Jülich	DE	Biotechnology	Elucidation of the poly-γ-L-glutamic acid (γ-L-PGA) synthesis mechanism and its mutational impacts in <i>Bacillus subtilis</i>
P-27	Etienne Dervyn	INRAE Jouy	FR	Biotechnology	Unravelling the potential of thermophilic <i>Geobacillus</i> spp. as chassis organisms for bioplastic upcycling
					OSIRIS: Orthogonal Sigma for Internal Resources Implementation towards Synthesis

P-28	Ryosuke Fukuda	Tsukuba U	JP	Regulation	Temperature-dependent gene regulation for environmental adaptation in <i>Clostridium perfringens</i>
P-29	Mitsuo Ogura	Tokai U	JP	Regulation	Regulatory mechanism of the operon containing genes encoding 5-oxoprolinase and manganese importer
P-30	Ahmad Altoun	Marburg U	DE	Regulation	Regulation of bacterial transcription by 6S
P-31	Veronika Kočárková	Czech Academy of Science	CZ	Regulation	Novel transcription factors in <i>Bacillus subtilis</i>
P-32	Yuzuki Shimada	Saitama U	JP	Regulation	Analysis of sigI regulation via lipoteichoic acid synthase in <i>Bacillus subtilis</i>
P-33	Ling Juan Wu	Newcastle U	UK	Regulation	Global regulatory role of ParB through parS-mediated autoregulation and gene silencing
P-34	Richard Daniel	Newcastle U	UK	Regulation	Differentiation between old and new peptidoglycan is required for coordinated cell growth in Bacteria
P-35	Matthieu Jules	INRAE Jouy	FR	Regulation	(p)ppGpp sets the level of tRNA charging through continuous regulation of translation
P-36	Anne-Gaëlle Planson	INRAE Jouy	FR	Regulation	Greedy reduction of <i>Bacillus subtilis</i> genome yields emergent phenotypes of high resistance to a DNA
P-37	Naoko Tsuji	Kyoto Sangyo U	JP	Regulation	Isolation of novel translation arrest peptides with RAPP and RGPP sequence motifs
P-38	Koichiro Masuda	Tokyo Agri U	JP	Regulation	Analysis of the suppressor strain, which recovered the survivability of <i>Bacillus subtilis</i> sigma factor minimizing strain
P-39	Sari Ikawa	Tokyo Agri U	JP	Regulation	In vivo comparative analysis of SigA /RpoD family in <i>Bacillus subtilis</i>
P-40	Miho Omote	Tokyo Agri U	JP	Regulation	Analysis of regulatory mechanism of RNA polymerase expression under nutrient starvation conditions in <i>Bacillus subtilis</i>
P-41	Natsumi Kimura	Tokyo Agri U	JP	Sporulation and development	Effect of the acetylation state of 2-oxoglutarate dehydrogenase complex on sporulation in <i>Bacillus subtilis</i> .
P-42	Teppei Kawakami	Tokyo Agri U	JP	Sporulation and development	Analysis of Arg phosphorylation site of SigA in <i>Bacillus subtilis</i> sporulation initiation.
P-43	Ritsuko Kuwana	Setsunan U	JP	Sporulation and development	Comparative Analysis of Thioflavin T and other Fluorescent Dyes for Fluorescent Staining of <i>Bacillus subtilis</i> Vegetative Cell, Sporulating Cell, and Mature Spore
P-44	Nozomu Obana	Tsukuba U	JP	Sporulation and development	A novel conserved protein complex controls sporulation in <i>Clostridium</i>
P-45	Nobuki Kuwabara	Hosei U	JP	Sporulation and development	Non-secreted intercellular signal transduction during sporulation in <i>Bacillus subtilis</i>
P-46	Tsutomu Sato	Hosei U	JP	Sporulation and development	Identification of CgeA as a glycoprotein that anchors polysaccharides to the spore surface in <i>Bacillus subtilis</i>
P-47	Maja Popović	U Ljubljana	SI	Sporulation and development	Disruption of phage integration gene leads to unexpected changes in <i>Bacillus subtilis</i>
P-48	Alexandre D'Halluin	Institut de Biologie Physico-Chimique	FR	Sporulation and development	The sporulation specific 3' exoribonuclease KapD is involved in the spore crust and outer coat formation in <i>B. subtilis</i>
P-49	Daisuke Seo	Kanazawa U	JP	Stress response	Investigation of a physiological role of the ferredoxin-NADP+ oxidoreductase paralog found in <i>Bacillus subtilis</i>
P-50	Hiroko Fukuda	Shinshu U	JP	Stress response	Application of a High-Throughput Colony Growth Measurement System to <i>Bacillus subtilis</i> under Different pH Circumstances
P-51	Yuma Okubo	Chiba U	JP	Sporulation and development	<i>B. subtilis</i> tRNA ^{Arg} is indispensable for the competence development
P-52	Koichiro Akiyama	NIG	JP	Cell division	Divergent cell shapes by a mutation in the IIA domain of MreB
P-53	Shota Suzuki	Rikkyo U	JP	Cell division	Adaptive evolution of oriC through in vitro propagation in RCR
P-54	Tomoaki Okado	Saitama U	JP	Others	Expression analysis of genes involved in propidium iodine in <i>Bacillus subtilis</i>
P-55	Kimihiro Abe	NIID	JP	Others	Spo0A-dependent membrane vesicle production in <i>Bacillus subtilis</i>
P-56	Ryuji Yamazawa	Setsunan U	JP	Others	YabG is a novel arginine-specific cysteine protease

P-57	Shigeki Kada	Meg Milk	JP	Others	<p>Exploration of D-amino acid producing lactic acid bacteria by bioassay using <i>Bacillus subtilis</i> mutants.</p> <p>Isolation of the suppressor mutations that restore the growth of <i>zwf</i> mutant in <i>B. subtilis</i></p> <p>Dual-wield NTPases: An uncharacterized protein family mined from AlphaFold Protein Database conserved among Bacilli and Identification of a minimum set of tRNA repertoire in <i>B. subtilis</i></p> <p>Comparative Analysis of Growth Retardation Effects on <i>Bacillus subtilis</i> Spores via Diverse Microbial Control Methods: Heat, Gamma-Rays, UV, and Essential Oils from Spices for the Development of Optimal Combined Novel approach to the genome (giant DNA) synthesis</p>
P-58	Takashi Inaoka	NARO	JP	Others	
P-59	Koya Sakuma	Nagoya U	JP	Others	
P-60	Akiko Soma	Chiba U	JP	Others	
P-61	Masakazu Furuta	Osaka Metro U	JP	Stress response	
P-62	Shinya Kaneko	Titech	JP	Biotechnology	