MBSJ-EMBO Poster Award
List of Awardees for Day 1 (November 27)

No.	Title	Presenter	Affiliation
1P-131	Development of novel transcriptional control tools using type I-D CRISPR-Cas system	Ryuya Watanabe	Sch. of Life Sci and Tech., Science Tokyo.
1P-213	TIMM17A, a subunit of mitochondrial inner membrane translocator, is degraded by YME1L1 when being unoccupied by precursors.	Hiroki Kinefuchi	Medical Research Institute, Tokyo Medical and Dental Univ.
1P-341	Effects of morphological and mechanical properties of nuclei on neuronal function	Yu Nagata	Grad. Sch. of Pharma. Sci., Univ. of Tokyo
1P-363	The recruitment of 14-3-3 proteins to GPCRs is driven by different protein kinases.	CARLOMARION CODOG CARINO	Grad Sch of Pharm Sci, Tohoku Univ
1P-364	GDP-bound Rab27a regulates endocytosis in pancreatic beta-cells	Soshiro Kodera	university of shizuoka
1P-432	CENP-C is involved in maintenance of centromeric chromatin via direct interaction with HJURP	Yutaka Mahana	Osaka Univ.
1P-471	Artificial genetic circuit converted by serine recombinase function	Toshiki Saito	Dept. Life Sci. Eng., Grad. Sch. Sci. Eng., Tokyo Denki Univ.
1P-668	A novel mechanism of dendritic refinement in postnatal mouse barrel cortex unveiled by rapid protein knockdown	Ayane Nihashi	Mammalian Neural Circuits Lab., NIG
1P-684	Elevated alcohol consumption and tolerance due to nucleocytoplasmic transport dysfunction	Yoshiatsu Aomine	Lab. for Advanced Brain Functions, IPR, Osaka Univ.
1P-703	Anti-Aβ antibody resolves the post-stroke neural reparative function impaired by Aβ pathology	Kento Otani	Department of Neuroinflammation and Repair, Medical Research Institute, Tokyo Medical and Dental University
1P-729	Transcriptional-metabolic reprogramming controls inflammatory memory in macrophages	Seonghyeon Hong	Kumamoto Univ.
1P-755	Cellular Landscape of Cytokine Storm in Severe Dengue	Mohamad Alkadi	Osaka University
1P-813	RNA binding protein ZCCHC24 maintains cancer stem like status in triple negative breast cancer	Yutaro Uchida	Institute of Science Tokyo, Department of Systems Biomedicine
1P-851	SKNY (SpatialKNifeY): Construction of a microenvironment analysis algorithm based on cancer cell clusters and its application to single cell spatial omics data of breast cancer	Shunsuke A Sakai	Division of Translational Informatics, Exploratory Oncology Research and Clinical Trial Center, National Cancer Center