

17 March (9:00-12:00) Oral Presentation Program

Chair: Naoya Yamaguchi (Facul. Biotech., Fukui Prefectural Univ.)		Chair: Daisuke Tsugama (Grad. Sch. Agric. Life Sci., Univ. Tokyo)		Chair: Hideki Takanashi (Grad. Sch. Agric. and Life Sci., Univ. Tokyo)		Chair: Kazumi Furukawa (National Inst. of Tech., Numazu College)		Chair: Yoshiyuki Yamagata (Fac. Agr., Kyushu Univ.)		Chair: Takaki Yamauchi (Biosci. Biotech. Cent., Nagoya Univ.)	
9:00	101 ☆Kuroda, R., Y. Kitomi, K. Sugimoto, Y. Uga (Inst. Crop. Sci., NARO)	201 ☆Chigira, K.1, C. Nomura2, A. Abe2, T. Ookawa1 (1.Graduate School of Agriculture, Tokyo University of Agriculture and Technology, 2.Iwate Biotechnology Research Center)	301 ○Kobayashi, A.1, H. Asai2, K. Maruyama2, F. Nakaoka1, S. Watanabe1, G. Chaya1 (1.Fukui Agri.Exp. Stn., 2.JIRCAS)	401 ○Takase, Y., T. Kuboyama (Col. Agr., Ibaraki U.)	501 ○Yoshimichi, F.1, A. Tomita2, A. Hairmansari3, N. Nafisah4, R. Hermanasari4, H. Saito1, T. Kazama5, K. Toriyama6, T. Sato6,7 (1.JIRCAS, 2.Faculty of Agriculture, Okayama Univ., 3.Research Center for Food Crops, National Research and Innovation Agency (BRIN), 4.Indonesian Center for Rice Research, 5.Graduate School of Agriculture, Kyushu Univ., 6.Graduate School of Agricultural Science, Tohoku Univ., 7.RIKEN Nishina Center)	601 ☆Lulie Desalegne, M., N. Munguambe Emilia, S. Zheng, D. Fujita (Grad. Sch. Agr., Saga Univ.)					9:00
	The extreme deep-rooted allele of qSOR1 in rice contributes to deep-rooting in dicotyledonous plants	Investigation of genotyping method for QTL analysis of rice using iSeq 100 sequencer	Characteristics of Lao rice genetic resources cultivated in Fukui.	QTL analysis of the stigma exertion rate in the F2 population between 'Koshihikari' and Oryza rufipogon W1944	Effects of QTLs for soil-surface rooting in rice (Oryza sativa L.) with respect to salinity and iron-toxicity stresses in paddy fields	Genetic analysis of tiller number on local rice varieties					
9:15	102 ☆Ogawa, G.1, G. Chaya1,2, S. Segami3, Y. Iwasaki1, K. Miura1 (1.Dep. Biosci., Fukui Pref. Univ., 2.Fukui Agri. Exp. Stn., 3.Res. Inst. Env. Agr. Fish., Osaka Pref.)	202 ○Komatsu, A.1, M. Ohtake1, M. Nagata1, M. Kondo2 (1.Institute of Agrobiological Sciences, NARO, 2.Graduate School of Bioagricultural Sciences, Nagoya University)	302 ☆Lim, S.1, A. Onoda1, C. Orn1,2, H. Iwamoto1, R. Ishikawa1, H. Saito3, Y. Sato4, T. Ishii1 (1.Grad. Sch. Agr. Sci., Kobe Univ., 2.Cambodian Agr. Res. Dev. Inst., 3.Trop. Agr. Res. Front, JIRCAS, 4.Natl. Inst. Genet.)	402 ☆Ono, S.1, E. Toda2, T. Tezuka3, A. Agata3,4, A. Kinoshita1, Y. Sato3, T. Okamoto1 (1.Dept. Biol., Tokyo Met. Univ., 2.Dept. Biol., Univ. Tokyo, 3.Dept. Gen. Evol., NIG, 4.Bioagric. Sci., Nagoya Univ.)	502 ☆Sato, T., T. Yamada (Grad. Sch. Agr., Tokyo U. Agr. Tech.)	602 ☆Ninomiya, T.1, H. Nguyen1,2, S. ZHENG1, D. Fujita1 (1.Grad. Sch. Agr., Saga Univ., 2.Grad. Sch. Agr. Sci., Kagoshima Univ.)					9:15
	Genetic analysis of short grain rice mutants, srs2 and srs6.	Field trial test of Gn1a-knockout rice line with enlarged sink capacity by genome-editing	Variations in grain traits among local rice varieties collected more than half-century ago in Indo-China countries	Production of hybrids and amphidiploids between cultivated rice and closely and distantly related wild Oryza to cultivated species by in vitro fertilization	Evaluation of protein aggregation in high-temperature treated seedlings of rice core collection cultivars and its relationship with physiological disorder	Analysis of qVBN11 for vascular bundle number at panicle neck in rice					
9:30	103 ○Tsugane, K.1, M. Maekawa2 (1.IBBP Center, Nat. Inst. Bas. Bio., 2.Inst. Plant Sci. Res., Okayama University)	203 ☆Iba, M.1, A. Henry3, Y. Kitomi2, M. Quintana3, M. Natividad3, A. Centeno3, K. Irie1, Y. Uga2 (1.Grad. Sch. Tokyo Univ. Agri., 2.NICS, 3.IRRI)	303 ☆Nohara, T.1, S. Yoshida2, K. Tanaka3, O. Aung4, H. Kikuno5, K. Irie5, K. Wakui2 (1.Agr. Science, Grad Agri, Tokyo Univ of Agri, 2.Dept of Bio Res Dev, Grad Agri, Tokyo Univ of Agri, 3.NODAI Genome Res Ctr, Tokyo Univ of Agri, 4.Dept of Agri Res, Min of Agri, Livest and Irrig, Myanmar, 5.Fac of Intl Agri and Food Stud, Tokyo Univ of Agri)	403 Maryenti, T.1, S. Koshimizu2, T. Ishii3, K. Yano4, ○T. Okamoto1 (1.Dept. Biol. Sci., Tokyo Met. Univ., 2.Dept. Informatics, NIG, 3.Tottori Univ., ALRC, 4.Dept. Biol. Sci., Meiji Univ.)	503 Koyama, T.1, T. Watanabe2, J. Murakami3, A. Nagano4,5, ○T. Yoshikawa2 (1.Fac. Agri., Kyoto Univ., 2.Grad. Sch. Agri., Kyoto Univ., 3.Fac. Agri., Kibi Int. Univ., 4.Fac. Agr., Ryukoku Univ., 5.IAB, Keio Univ.)	603 ☆YA, M.1, S. Zheng1, T. ANAI1,2, D. Fujita1 (1.Grad. Sch. Agr., Saga Univ., 2.Grad. Sch. Agr., Kyushu Univ.)					9:30
	Analysis of the dominant large-grained rice mutant Lgg	Effect of pyramiding root QTLs on root morphological traits in hydroponic and root distribution in field condition	A Report of B chromosome in B. juncea populations collected in southern Shan State in Myanmar.	Genome composition of wheat-rice hybrid (Oryzawheat) produced by in vitro fertilization system	Changes in blast disease response along with vegetative phase transition in rice	Development and characterization of early heading mutant lines of glutinous rice variety 'Hiyokumochi'.					9:45

