Poster Presenter List (Names are in alphabetical order. Presentation order is not yet determined.)

- Benedicta Obenewaa Dankyi, IgG and IgM Responses to PfEMP1 Domains
 Associated with Semi-immunity to Clinical Malaria in Burkinabe Children
 Under Five
- Chalermpon Kumpitak, Age-Associated Gene Expression Profiling in Malaria Vector Anopheles Mosquitoes
- Chayanut Suansomjit, Malaria Resurgence in Thailand I: Cross-Sectional Surveys
- Chukwuma Stephen EZENWANNE, Plasmodium knowlesi Rosette Related Protein 1 is associated with the formation of rosettes
- Faridah Ousseini, A novel molecular assay for the identification of six Plasmodium species including Plasmodium ovalecurtisi and Plasmodium ovalewallikeri
- Hajime Takahashi, Construction of eukaryotic artificial cell-based sensors for simultaneous detection of multiple analytes at ambient temperature
- Hamada Mana, Isolation and characterization of rabbit monoclonal antibodies suitable for proximity-dependent protein biotinylation analysis
- Haruna Arai, Elucidation of the molecular mechanism of OPSIN3 in metastatic prostate cancer
- Hikari Ozawa, Enzymatic Lipidation Enhances the Antimicrobial Activity of Peptides
- Hinako Fujisaki, Analysis of proximity cetuximab-induced EGFR interactome using AirID
- Hiroshi Sakai, Systematic Identification of Pax1 Transcriptional Coactivators
- Huai Chuang, Inhibition of SICA-HUVEC Surface Expression in Plasmodium knowlesi by Brefeldin A and Torin 2
- Junpei Fukumoto, Ultrastructural expansion microscopy enable for detailed protein visualization in human malaria parasites
- Kana Fujihara, Development of Anti-CD5 VNAR Antibody for Targeted Nucleic Acid Delivery in T Cells
- Kanit Phumchuea, Malaria Resurgence in Northwestern Thailand II: Passive Case Detection
- Keisuke Konishi, Development of a Novel CRBN Modulator for Targeted Protein Degradation

- Kongpope Chareonsuk, Mitochondrial DNA Copy Number in Plasmodium falciparum is Stage-Dependent
- Maho Tanimoto, Development of VNAR Antibody-based Technology for Degradation of Target Membrane Proteins
- Mari Saida, Mitochondrial transfer from skeletal muscle to macrophages via extracellular vesicles
- Masahito Asada, Babesia bovis spherical body protein 3 is a crucial protein for ridge formation on infected red blood cells
- Masaki Takeguchi, An attempt to identify non-TbHpHbR proteins that bind to anti-TbHpHbR polyclonal antibodies in Trypanosoma brucei
- Masashi Takeda, Elucidating the function of androgen receptor in the heart.
- Masaya Nagano, Cryo-EM analysis of Axin2 homo-oligomer involved in the termination of the Wnt/β-catenin pathway
- Mayumi Tachibana, The ookinete crystalloid protein PH3 is involved in the crystalloid ultrastructure formation
- Miako Sakaguchi, Binding analysis of SICA-HUVEC domains in Plasmodium knowlesi by Cell ELISA
- Mie Kurata, Pathology-Based Optimization of Immunohistochemistry for Difficult-to-Detect Tumor Antigens: Addressing Unmet Needs in Precision Oncology
- Nattawan Rachaphaew, Factors predicting parasitemia and gametocytemia among patients with Plasmodium vivax infection in Thailand
- Nichakan Inthitanon, Malaria care-seeking behaviors and infection prevalence among short-term Myanmar migrants in Thailand
- Noritaka Saeki, ER α signaling differentially modulates inflammatory and proliferative responses in lining and sub-lining synovial fibroblasts in an inflammatory arthritis
- Palakorn Chintanawiwat, Genetic Signatures of Plasmodium vivax
 Circumsporozoite Surface Protein During Malaria Resurgence in Thailand
- Pattamaporn Petchvijit, Molecular Identification and Species Composition of Anopheles Larvae in Malaria-Endemic Region of Thailand
- Pinyapat Kongngen, A Human Hepatocyte Platform for advancing Antimalarial Drug and Vaccine development Against Pre-erythrocytic stage of Plasmodium vivax
- Piyarat Sripoorote, Sociodemographic and behavioral determinants of Plasmodium vivax-specific antibody responses among short-term Myanmar migrants in Thailand

- Qinaer Madingbieke, Toward the establishment of Toxoplasma mutator
- Rojrung Rattanaporn, Detection of Plasmodium lactate dehydrogenase in a bead-based multiplex assay
- Riko Uehara, Cell-Free Multistep Gene Regulatory Cascades Using Eukaryotic ON-Riboswitches Responsive to in Situ Expressed Protein Ligands
- Satoru Otowa, Functional analysis of slow-twitch fibers in Sox6-deficient skeletal muscle
- Shun Shiraishi, Identification and functional analysis of host proteins that interact with EV-A71 protease and polymerase
- Srisuda Keayarsa, D-glucose and fetal bovine serum supplementation support the erythrocytic stage development of Plasmodium knowlesi: ex vivo
- Thananya Jinato, Harnessing human tonsil organoids to evaluate vaccine driven immune activation
- Thant Zin Tun, Selective phospholipid binding by PH1 and its role in AMA1 translocation during Plasmodium yoelii erythrocyte invasion
- Tuyet-Kha Nguyen, Development and characterization of human monoclonal antibodies against Plasmodium vivax blood-stage antigens
- Wakaba Yagi, Regulation and physiology of yeast vacuolar amino acid transporter Avt3
- Xueling Kuang, The effect of MEK1 inhibitors on proliferation of the malaria parasite
- Yoshihiko Kuchitsu, The contribution of human STING major variants into the inflammatory responses in model cells for diseases caused by STING constitutive activation
- Yoshiya Murata, The first step towards establishing prenyltransferase assay system in Escherichia coli
- Yoshika Tanoue, Physiology of vacuolar accumulation of basic amino acids in budding yeast Saccharomyces cerevisiae
- Yusuke Kojima, Exploring Novel Regulatory Mechanisms of Lipoxygenases via Protein-Protein Interactions
- Yuta Yanagihra, Identification of novel Dnmt1 interactors

If you would prefer not to have your full name and title displayed, please contact us (pim2025@pim-sympo.jp). We are happy to abbreviate your information upon request.