CBI2026

Best Poster Award

P09-06

Shinya Kawano

Gifu pharmaceutical university

Boltz-2-guided discovery of potent, selective AKR1B10 inhibitors

Like! Poster Award

P09-02

Rento Yabuki

Department of AI Systems Medicine, M&D Data Science Center, Institute of Integrated Research, Institute of Science Tokyo

Quantum-Chemically Supervised Pre-training for Cyclic Peptide–Protein Affinity Prediction

Excellent Poster Awards

P01-13

Seita Kawakami

Kagoshima University

Benchmark Evaluation of a First-Principles-Based Protein-Protein Docking Method for Epitope Prediction in Diverse VHH–Antigen Complexes

P02-02

Toshiaki Watanabe

DAIICHI SANKYO CO., LTD.

AI Agents for Formalizing Medicinal Chemistry Knowledge into Machine-Readable Chemical Transformation Rules

P02-18

Natsumi Hayakawa

Tohoku University

Preliminary Analysis for Developing a Transformer-Based Foundation Model of Disease Trajectories in a Japanese Population

P03-03

Taketo Tsuga

Institution for Protein Research The University of Osaka

Deciphering Tyrosine Kinase Substrate Specificity through Structure Prediction and Inverse Folding

P04-11

Jaeseok Kwon

kyushu institute of technology

Development of a multi-label toxicity prediction model for the safety evaluation of peptide drugs

P05-16

Yuki Kobayashi

Kyoto University

Development of a Time-Series Analysis Method for Single-Cell RNA Sequencing Data Constrained by Biological Models Using Four-Dimensional Variational Data Assimilation

P06-23

Yusuke Tateishi
Graduate School of Science and Technology, Kumamoto University

From Where to What: Chemistry-Informed Atom Attribution Framework for Physicochemical Interpretation of Molecular Activity Predictions

P07-07

Makoto Koide
Department of Biomedical Data Intelligence, Graduate School of Medicine, Kyoto University

Development and Evaluation of an AI Agent for Personalized Meal Planning Using Structured Recipe Data

P08-02

Mikiya Tsukuda
Mie university

Multi-Stimuli-Responsive DNA Hydrogels Integrated with Modular DNA Origami Nanoactuators

P09-04

Kosuke Takeuchi
DAIICHI SANKYO CO., LTD.

Accelerating Hit Expansion: From R-Group Enumeration to Automated Synthesis

P09-19

Tatsuya Yoshizawa
Yokohama city university

Can General-Purpose LLMs Edit Molecules Like Medicinal Chemists?