

28th(WED) Hall

9:00	Registration
9:50	Opening Ceremony
10:00	Session Shiroh Futaki(Institute for Chemical Research, Kyoto University) Intracellular delivery using arginine-rich peptides
11:00	Session Kazunori Kataoka(The University of Tokyo) Smartpolymeric micelles as nanocarriers for gene and drug delivery
12:00	Luncheon seminar INFOCOM CORPORATION
13:30	Oral Presentation Differentiation of Organ Availabilities Contributing to Low Oral Bioavailability by Sequential and Simultaneous Analyses: Intestinal Conjugative Metabolism Impacts on Oral Bioavailability Takashi MIZUMA
13:50	Oral Presentation Evidence for Functional Variation of the Death Receptor 3 (<i>DR3</i>) Gene That Inhibits Apoptosis and Predisposes to Progressive Joint Destruction in Rheumatoid Arthritis Shunichi SHIOZAWA
14:10	Oral Presentation Crystal structure analysis of a hydrolase form <i>Plasmodium falciparum</i> . Nobutada TANAKA
14:30	Coffee break
14:40	Session Kimito Funatsu(The University of Tokyo) Development of Total System for Molecular Design
15:30	Coffee break
15:50	Session Fumiyoshi Yamashita(Kyoto University) Current Status and Perspectives of <i>In Silico</i> Pharmacokinetics
16:50	Coffee break
17:00	Poster Session & Mixer 1(at HOUOU_NO_MA)

29th(thu) Hall

9:00	Registration
10:00	Session Junichi Higo (Tokyo University of Pharmacy and Life Science) Free-energy landscape of peptides in explicit water obtained from molecular dynamics simulation
11:00	IBM Session Blue Gene: protein folding and drug discovery Jed W. Pitera(IBM Research)
12:00	Luncheon seminar Sumisho Electronics Co.,Ltd
13:30	Session A.L.Hopkins(Pfizer Global Research & Development) The Druggable Genome: Mapping Chemical-Biological Space for Drug Discovery
14:30	Coffee break
14:40	Oral Presentation <i>Ki</i> Bank (A Database for Structure-Based Drug Design) and Its Application Shinji AMARI
15:00	Oral Presentation In Silico Screening by Pharmacophore Model Hidenori SESHIMO
15:20	Oral Presentation General Model for Estimation of the Inhibition of Protein Kinases Using Monte Carlo Simulations Yukio TOMINAGA
15:40	Coffee break
15:50	Session Julian Simon (Fred Hutchinson Cancer Research Center) Budding Yeast <i>Saccharomyces cerevisiae</i> : an Organism for Anticancer Drug Discovery
16:50	Coffee break
17:00	Poster Session & Mixer 2(at HOUOU_NO_MA)

30th(Fri) Hall

9:30	Open Registration
10:00	Session Makoto Taiji(Genomic Sciences Center, RIKEN) Petaflops special-purpose computer for molecular dynamics simulations and its application to protein science
11:00	Session Hideaki NATSUGARI(The University of Tokyo) Drug discovery through integration of synthetic organic chemistry and computational chemistry: how can I make better use of computational chemistry?
12:00	Lunch
13:10	Ceremony for poster awards
13:30	Oral Presentation On Kinase Interactome Networking Interacted with Cellular Nuclear Receptors Jian-Qin LIU
13:50	Oral Presentation The New Clustering Technique in Gene Discovery Data Analysis Ryo MUKAE
14:10	Oral Presentation Construction of the Web-accessible comprehensive database for drug discovery and development, “TP-Search” Kazuya MAEDA
14:30	Coffee break
14:40	Oral Presentation <i>In silico</i> proteome-wide affinity fingerprinting of known drugs Kazuo SATO
15:00	Oral Presentation Genome-wide homology modeling of protein-protein complex structures: alignment based prediction of protein-protein interaction Mayuko TAKEDA-SHITAKA
15:30	Closing ceremony