Presentation time is organized by whether the last part (suffix) of Poster Session number is odd/even.

Odd number: 13:50-14:50 Even number: 14:50-15:50

Abstracts marked with * in the abstract number eligible for IUPAB2024 Student and Early Career Researcher Poster Award voting Ex) *25P-999

Protein: Structure

28P-001 Cryo-EM analysis of human GLUT9 Daiki Matsushita, Yongchan Lee, Yu Toyoda, Teppei Takada, Tomohiro Nishizawa Graduate School of Medical Life Science, Yokohama City University, Yokohama, Japan 28P-002 Molecular basis of substrate recognition in human y+LAT1-CD98hc complex Juntaro Nakahara, Yongchang Lee, Natsumi Yoshida, Pattama Wiriyasermkul, Ryo Ekimoto, Mitsunori Ikeguchi, Sushi Nagamori, Tomohiro Nishizawa Graduate School of Medical Life Science, Yokohama City University 28P-003 Particle formation for nanofiber elongation in Fibroin artificial

sequence Kento Yonezawa, Chan Kok Sim, Takehiro Sato, Haruya Kajimoto,

Kiichi Hayashi, Takuya Sawai, Yusuke Okamoto, Rakuri Aiba, Yuki Nakatani, Kenta Kimura, Yoichi Yamazaki, Sachiko Toma-Fukai, Yugo Hayashi, Hironari Kamikubo NAIST MS/NAIST CDG

28P-004 Cryo-EM analysis of mouse b0,+AT-rBAT complex Aoi Maeda, Yongchan Lee, Pattama Wiriyasermkul, Sushi Nagamori, Tomohiro Nishizawa Graduate School of Medical Life Science, Yokohama City University

28P-005	Cryo-EM Structure of MexB-MexY Chimera Protein MexBYB Multidrug Efflux Pump Jiye Wang, Kenta Tsutsumi, Ryosuke Nakashima, Kunihiko Nishino, Eiki Yamashita, Atsushi Nakagawa Institute for Protein Research, Osaka University, Osaka, Japan
28P-006	Investigation of the effect of ATP/ADP for formation of 2-Cys peroxiredoxin (Prx2) high molecular weight complex <u>Trang Ngoc Tran</u> , Ryusei Yamada, Hiroki Konno Graduate School of Frontier Science Initiative, Kanazawa University, Kanazawa, Japan
28P-007	Comprehensive analysis of different fold proteins with similar interfaces <u>Takumi Sekine</u> , Kazuo Fujiwara, Masamichi Ikeguchi Department of Biosciences, Soka University, Hochioji, Japan
28P-008	Solution structure of clock protein complex KaiA-KaiC Ken Morishima, Masahiro Shimizu, Ritsuki Sakamoto, Yasuhiro Yunoki, Rintarao Inoue, Masaaki Sugiyama Institute for Integrated Radiation and Nuclear Science, Kyoto University
28P-010	The brain metabolites, betaine and dimethyl glycine disrupt acetylcholinesterase activity and enhance the inhibitory effect of Donepezil, Rivastigmine, and Galantamine Laishram Rajendrakumar Singh, Kritika Kumari Dr. B. R. Ambedkar Center for Biomedical Research, University of Delhi, Delhi-110007, India
28P-011	Towards serial femtosecond crystallography of metalloproteins with sub-ångström details <u>Faisal Koua</u> , Tiankun Zhou, Jay-How Yang, Marcin Sikorski, Jayanath Koliyadu, Mohammed Vakili, Johan Bielecki, Richard Bean, Tokushi Sato, Adrian Mancuso European XFEL, Schenefeld, Schleswig-Holstein, Germany
28P-012	Structural basis of main proteases of HCoV-229E bound to inhibitor PF-07304814 and PF-07321332 Qisheng Wang Shanghai Advanced Research Institute, Chinese Academy of Sciences, Shanghai,

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Protein: Structure & Function

28P-014 Evidence for an alternative YidC-assisted insertion mode - Exploring a putative parallel YidC dimer.

Denis Knyazev, Lukas Winter, Andreas Vogt, Sandra Posch, Yavuz Öztürk, Christine Siligan, <u>Nikolaus Goessweiner-Mohr</u>, Nora Hagleitner-Ertugrul, Hans-Georg Koch, Peter Pohl Institute of Biophysics, Johannes Kepler University Linz, Gruberstrasse 40, A-4020 Linz, Austria

28P-015 Modeling of Photoswitchable Ligands Linked to Physiology Wieslaw A Nowak Department of Biophysics. Institute of Physics. N. Copernicus University in Torr

Department of Biophysics, Institute of Physics, N. Copernicus University in Torun, Poland

28P-016 Identifying functional hotspot residues for activation in M2 muscarinic receptor

Yuya Sugiura, Tatsuya Ikuta, Yuji Sumii, Hirokazu Tsujimoto, Ryoji Suno, Putri Nur Arina Binti MOHD ARIFF, So Iwata, Norio Shibata, Asuka Inoue, Takuya Kobayashi, Hideki Kandori, <u>Kota Katayama</u> Department of Life Science and Applied Chemistry, Nagoya Institute of Technology/ OptoBioTechnology Research Center, Nagoya Institute of Technology

28P-017 Aqp5 mutations in patients suffering from palmoplantar keratoderma (Bothnian type)

<u>Christine Siligan</u>, Helena Zich, Nikolaus Gössweiner-Mohr, Anna Stoib, Peter Pohl Institute of Biophysics, Johannes Kepler University, Linz, Austria

28P-018 pH-Induced Conformational Dynamics and Oligomeric Assembly of Peroxiredoxin 6: Insights into Catalytic Mechanisms

<u>Hamidur Rahaman</u>, Shahnaj Sharifun, Kakchingtabam Pushpa Department of Biotechnology, Manipur University, Indo Myanmar Road, Canchipur, Imphal, India-795003

28P-019 Exploring Covalent Bond Electron Densities in the Active Site of the EcoRV-DNA Complex through QM/MM Metadynamics

<u>Hiroki Sato</u>, Itaru Onishi, Mika Mitsumatsu, Ryotarou Matsuda, Masayuki Irisa Comp. Sci. and Sys. Eng., Kyushu Inst. Tech., Japan

28P-020 Fantastic Enzymes and where to find them Ehmke Pohl, Katy Cornish, Stefanie Freitag-Pohl, Arnthór Aevarsson Department of Chemistry, Durham University, Durham, DH1 3LE, UK 28P-021 Theoretical study on allosteric control mechanism of a luminescent reaction of bioluminescent protein Aequorin Tomohiro Ando, Toshiya Funahashi, Toru Nakatsu, Shigehiko Hayashi Grad. Sch. of Sci. Kvoto Univ. 28P-022 Correlating single molecule studies with Cryo-EM structures to understand the inner workings of ATP synthase. Meghna Sobti, Hiroshi Ueno, Simon Brown, Hiroyuki Noji, Alastair Stewart Molecular, Structural and Computational Biology Division, The Victor Chang Cardiac Research Institute, Darlinghurst, Australia/St Vincent's Clinical School, Faculty of Medicine, UNSW Sydney, Kensington, Australia. 28P-023 Molecular basis for heat-hypersensitive mutants of ryanodine receptor type 1 Liu Chujie, Yamazawa Toshiko, Oyama Kotaro, Murayama Takashi, Kobayashl Takuya, Harada Yoshie, Suzuki Madoka Institute for Protein Research, Osaka University 28P-024 Proton-coupled electron transfer dynamics and ligand binding in the mycobacterial respiratory supercomplex III2IV2 Ana Patricia Gamiz-Hernandez, Daniel Riepl, Terezia Kovalova, Sylwia M. Król, Dan Sjöstrand, Martin Högborn, Peter Brzezinski, Ville R. I. Kaila Department of Biochemistry and Biophysics, The Arrhenius Laboratories for Natural Sciences, Stockholm University, SE-106 91, Stockholm, Sweden. 28P-025 High-speed AFM observation of collagen degradation process by Grimontia hollisae collagenase Hayato Yamashita, Keisuke Tanaka, Yuko Ushiki-Kaku, Akihiro Tsuji, Shunii Hattori, Masavuki Abe Graduate School of Engineering Science, Osaka University 28P-026 Structural basis for the pH-dependent functional regulation of cytochrome b6f complex from Chlamydomonas reinhardtii Akihiro Kawamoto, Hatsuki Tanabe, Shin-Ichiro Ozawa, Hideaki Tanaka, Yuichiro Takahashi, Genji Kurisu Institute for Protein Research, Osaka University, Japan

28P-027 Structural insights into the elongation complex of RNA polymerase II paused at the +1 nucleosome entrance Masahiro Naganuma, Tomoya Kujirai, Haruhiko Ehara, Tamami Uejima, Tomoko Ito, Mie Goto, Mari Aoki, Masami Henmi, Sayako Miyamoto-Kohno, Mikako Shirouzu, Hitoshi Kurumizaka, Shun-ichi Sekine RIKEN Center for Biosystems Dynamics Research, Yokohama, Japan 28P-028 Positive allosteric modulation of cytochrome c oxidase activity Yuya Nishida, Takumi Tateno, Takemasa Nagao, Yasunori Shintani National Cerebral and Cardiovascular Center, Osaka, Japan 28P-029 The structure and function of the ghrelin receptor coding for drug actions Yuki Shiimura, Dohyun Im, Ryosuke Tany, Hidetsugu Asada, Ryoji Kise, Hideko Wakasugi-Masuho, Kazuma Matsui, Jun-ichi Kishikawa, Takayuki Kato, Masayasu Kojima, So Iwata, Ikuo Masuho Division of Molecular Genetics, Institute of Life Science, Kurume University/ Department of Cell Biology, Graduate School of Medicine, Kyoto University 28P-030 Cracking the code: A computational expedition into neurodegenerative polypeptides and innovative therapies Ioana Mariuca Ilie, Simone Ruggeri University of Amsterdam, The Netherlands 28P-031 Structural and Functional Elucidations of Druggable Viral Macrodomains Chun-Hua Hsu National Taiwan University, Taipei, Taiwan 28P-032 Synthesis of versatile neuromodulatory molecules by a gut microbial glutamate decarboxylase Pavani Dadi, Clint Pauling, Abhishek Shrivastava, Dhara D. Shah School of Mathematical and Natural Sciences, Arizona State University, Glendale, AZ, USA./Biodesign Center for Fundamental and Applied Microbiomics, Arizona State University, Tempe, AZ, USA. 28P-033 An integrated approach using seguential and structural features for precise prediction of protein-protein binding affinity Zhongliang Guo, Osamu Muto, Rui Yamaguchi Aichi Cancer Center Research Institute

28P-034 Structure of the human 80S ribosome at 1.9 Å resolution – the molecular role of chemical modifications and ions in RNA

<u>Charles Barchet</u>, Samuel Holvec, Antony Lechner, Léo Fréchin, Nimali De Silva, Isabelle Hazemann, Philippe Wolff, Ottilie von Loeffelholz, Bruno Klaholz

Centre for Integrative Biology (CBI), Department of Integrated Structural Biology, IGBMC (Institute of Genetics and of Molecular and Cellular Biology), 1 rue Laurent Fries, Illkirch, France/Centre National de la Recherche Scientifique (CNRS) UMR 7104, Illkirch, France/Institut National de la Santé et de la Recherche Médicale (Inserm) U964, Illkirch, France/Université de Strasbourg, Strasbourg, France

Protein: Physical property

28P-035	Amyloid formation of the β2-microglobulin variants, D76N and V27M:Diverse diseases via a common assembly mechanismMasatomo So, Roberto Martinez, Nicolas Guthertz, Martin Wilkinson,Sheena RadfordUniversity of Leeds, Leeds, UK/Kyoto University, Kyoto, Japan
28P-036	Negative Charge Increment during Evolution of Ferritin Takumi Kuwata, Yusuke Murakami, Kazuo Fujiwara, <u>Masamichi Ikeguchi</u> Department of Biosciences, Soka University
28P-037	Prediction of detailed structures over the entire free energy landscape of protein folding using extended statistical mechanical models and restrained simulations Koji Ooka, Munehito Arai College of Arts and Sciences, The University of Tokyo, Tokyo, Japan.
28P-038	Structural studies of protein condensates prepared by ultracentrifugation/air-drying Ryuga Someya, Akira Nomoto, Suguru Nishinami, Hiroka Sugai, Kentaro Shiraki Institute of Pure and Applied Sciences, University of Tsukuba, 1-1-1 Tennodai, Tsukuba, Ibaraki 305- 8573, Japan

Protein: Function

28P-039	SsUrel is a pH-gated urea channel from Streptococcus salivarius
	Anna Stoib, Xenia Fischer, Sandra Posch, Felix Wolkenstein, Sahar Shojaei,
	Christine Siligan, Nikolaus Goessweiner-Mohr, Andreas Horner
	Institute of Biophysics, Johannes Kepler University Linz, Gruberstr. 40, 4020 Linz,
	Austria

28P-040 Effect of the inorganic phosphate on the iron oxidation/mineralization activity of Escherichia coli non-heme ferritin A Takumi Kuwata, Kazuo Fujiwara, Masamichi Ikeguchi

Dept. of Biosci. Grad. Sch. of Sci. and Eng. Soka Univ., Tokyo, Japan,

28P-041 Structural Changes of Poly(ethylene terephthalate) undergoing **Enzymatic Degradation** Daisuke Tadokoro, Tomoya Imai

Reserch Institute of Sustainable Humanosphere, Kyoto University

28P-042 Mode of action of virulence factors of intracellular pathogens studied with time-resolved and high-resolution atomic force microscopy Christian Nehls, Thomas Gutsmann Research Center Borstel, Division of Biophysics, Borstel, Germany, /Centre for Structural Systems Biology, Hamburg, Germany

Protein: Measurement & Analysis

28P-043	BioSAXS for solution protein structure analysis at SPring-8 Satoshi Nagao, Hiroyasu Masunaga, Nobutaka Shimizu, Masaki Yamamoto, Hiroshi Sekiguchi JASRI/SPring-8
28P-044	Biomolecular Interactions with the NanoTemper Dianthus Stefanie Freitag-Pohl, Dorata Gasparikova, Kate V. Sowerby, Abbey M. Butler, Ehmke Pohl Department of Chemistry, Durham University, Durham DH1 3LE, UK
28P-045	Easy and fast LLPS size estimation using microplate reader <u>Enomoto Mayu</u> , Suai Anzawa, Tadashi Kodama, Kyoko Furuita, Wataru Togawa, Ryoga Kobayashi, Naotaka Sekiyama, Yohei Miyanoiri, Toshimichi Fujiwara, Hidehito Toshio, Chojiro Kojima Yokohama National University

- 28P-046 Spatiotemporal and global profiling of DNA-protein interactions and substrates of lysine-modifying enzymes in living cells <u>Minjia Tan</u>, Hao Hu, Wei Hu, An-Di Guo, Linhui Zhai, Xiao-Hua Chen State Key Laboratory of Drug Research, Shanghai Institute of Materia Medica, Chinese Academy of Sciences, Shanghai, China
- 28P-047 EPR Spectroscopy Combined with Rapid Freeze-Quenching Reveals Relationship Between Temperature Dependence of Active Site Rearrangement and of Activity in Inorganic Pyrophosphatase <u>Masaki Horitani</u>, Hiroshi Sugimoto, Yuri Kasu Saga University

Protein: Design & Engineering

28P-048	DNA replication triggered by small-molecule for in vitro auto-selection
	of enzymes
	Thibault Philippe Laurent Di Meo, Yannick Rondelez, Hiroyuki Noji
	Noii Laboratory. Department of Applied Chemistry. Graduate School of Engineering.
	The University of Tokyo
28P-049	Generation of antibodies to an extracellular region of the transporters Glut1/Glut4 by immunization with a designed antigen
	<u>Makoto Nakakido,</u> Taichi Sumikawa, Ryo Matsunaga, Daisuke Kuroda,
	Satoru Nagatoishi, Kouhei Tsumoto
	The University of Tokyo
28P-050	Heterocomponent protein tube formation via "Nature Inspired Protein Assembly Design (NIPAD)"
	Masahiro Noji, Yukihiko Sugita, Makito Miyazaki, Yuta Suzuki
	Research Fellow of Japan Society for the Promotion of Science, Japan/Graduate School of Human and Environmental Studies, Kyoto University, Kyoto, Japan
28P-051	Understanding the tussle between aggregation-prone proteins and chaperons; toward the development of an enzyme immobilization
	platform
	Nilanjana Bose
	INDIAN INSTITUTE OF TECHNOLOGY, DELHI

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28P-052 De Novo Design of P-loop Harboring Protein

Takahiro Kosugi, Nobuyasu Koga

Institute for Molecular Science (IMS), National Institutes of National Sciences (NINS)/ Exploratory Research Center on Life and Living Systems (ExCELLS), National Institutes of National Sciences (NINS)/Molecular Science Program, SOKENDAI (The Graduate University for Advanced Studies)/PRESTO, Japan Science and Technology Agency

28P-053 De novo design of helical peptide binders targeting the KIX domain of CBP

Shunji Suetaka, Munehito Arai Department of Life Sciences, Graduate School of Arts and Sciences, The University of Tokyo, Tokyo, Japan

28P-054 Red fluorescent proteins engineered from green fluorescent proteins <u>Hiromi Imamura</u>, Shiho Otsubo, Mizuho Nishida, Norihiro Takekawa, Katsumi Imada Kyoto University

28P-055 A "Protein Structure Transformer" for integrative structural biology and molecular design

Lucien Fabrice Krapp, Fernando Meireles, Luciano Abriata, Matteo Dal Peraro Laboratory for Biomolecular Modeling, EPFL

28P-056 Genetically encoded fluorescent biosensors for cellular metabolism <u>Yusuke Nasu</u>, Yuki Kamijo, Robert Campbell Department of Chemistry, School of Science, The University of Tokyo/PRESTO, Japan Science and Technology Agency

28P-057 Role of "relaxed" peptide bond in protein structure and function Kaori Chiba, Masaru Hoshino, Hiromu Ohshima, Manami Suwa National Institute of Technology. Ibaraki college

Protein: Intrinsic disorder

28P-058 Deciphering the Role of GM1 Ganglioside-Bound Aβ Species in Alzheimer's Disease: Insights from Monoclonal Antibody 4396C and Advanced Biophysical Techniques

<u>Maho Yagi-Utsumi</u>, Satoru G. Itoh, Yui Kanaoka, Shogo Miyajima, Katsuhiko Yanagisawa, Katsuyuki Nishimura, Hisashi Okumura, Takayuki Uchihashi, Koichi Kato

Graduate School of Pharmaceutical Sciences, Nagoya City University/Exploratory Research Center on Life and Living Systems (ExCELLS), National Institutes of Natural Sciences

Membrane proteins

28P-060	Strategies for Cancer Therapy by Regulating Intracellular Dynamics of Antibody Drugs
	<u>Kazuya Kabayama,</u> Yoshiyuki Manabe, Atsushi Toyoshima,
	Kazuko Kaneda, Tadashi Watabe, Koichi Fukase
	Institute for Radiation Sciences, Osaka University/Department of Chemistry, Graduate School of Science, Osaka University/FRC, Graduate School of Science, Osaka University
28P-061	Differential molecular responses of PIEZO1 to membrane tension and ligand binding observed with diffracted X-ray tracking
	Mei Ishii, Kayoko Kawaguchi, Mayui Sugiura, Hiroshi Sekiguchi,
	Tatsuya Arai, Kazuhiro Mio, Yuji Sasaki
	The Graduate School of Frontier Sciences, The Univ. of Tokyo
28P-062	Thermodynamic Analysis of pH-Dependent Substrate Binding in the Multidrug Transporter, EmrE
	Kazumi Shimono, Keisuke Matsuda, Shoko Suzuki, Shuichi Miyamoto, Seiji Miyauchi
	Sojo University/Toho University
28P-063	INTRAMOLECULAR DOMAIN DYNAMICS OF LIGHT-HARVESTING
	PROTEIN LH1-RC OBSERVED BY THE DIFFRACTED X-RAY
	TRACKING METHOD
	Tatsunari Ohkubo, Tatsuya Arai, Hiroshi Sekiguchi, Kazuhiro Mio,
	Yuji C. Sasaki
	Grad. Sch. Med Life Sci, Yokohama City University/OPERANDO-OIL, AIST

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28P-064 NMR and modelling study of interaction of spider Cys-knot toxins with membrane and cationic ion-channels of P-loop superfamily Zakhar Shenkarev, Pavel Mironov, Eugene Kovalenko, Dmitrii Kulbatskii, Alexander Paramonov, Mikhail Shulepko, Maxim Zaigraev, Ekaterina Lyukmanova Department of Biology, MSU-BIT Shenzhen University, Shenzhen, China/Shemyakin-Ovchinnikov Institute of Bioorganic Chemistry, Moscow, Russia

28P-066 Diffusion and Oligomerization of GPCRs in Live Cells – The Impact of Ligands and Membrane Disruptors Claudiu Gradinaru University of Toronto, Canada

- 28P-067 Usability of the novel detergent NDT-C11 in cryoEM Christoph Gerle, Chimari Jiko, Jiannan Li, Chai Gopalasingam, Hideki Shigematsu, Pilseok Chae Riken SPring-8 Center, Sayo, Japan
- 28P-068 The functional role of the pleckstrin homology domains of dynamins in evolution and disease Isabel Pérez-Jover, Javier Espadas, Irune Ornos, Julene Ormaetxea Guisasola, Isaac Santos-Pérez, Vadim Frolov, <u>Anna Shnyrova</u> Biophysics Institute (CSIC, UPV/EHU), Leioa, Spain/Department of Biochemistry and Molecular Blology, University of the Basque Country, Leioa, Spain

DNA & DNA binding proteins

- 28P-069 The conformational analysis of DNA and nucleosome with doxorubicin analyzed by molecular dynamics simulation
 <u>Hisashi Ishida</u>, Hidetoshi Kono
 Institute for Quantum Life Science, National Institutes for Quantum Science and Technology

 28P-070 DNA-binding and -unwinding Dynamics of the nonhexameric
- 28P-070 DNA-binding and -unwinding Dynamics of the nonhexameric Escherichia coli UvrD helicase lacking C-terminal amino acids Hiroaki Yokota The Graduate School for the Creation of New Photonics Industries

28P-071 Role of Long-range Interactions in Protein-DNA Recognition

Anastasia A. Anashkina

Laboratory of DNA-protein interactions, Engelhardt Institute of Molecular Biology, Russian Academy of Sciences, 119991 Vavilov str, 32, Moscow, Russia/Department of Information and Internet Technologies, Institute of Biodesign and Modeling of Complex Systems, Sechenov University, 119048, Trubetskava Ulitsa, 8 b.2, Moscow, Russia

28P-072 In-cell NMR analysis on base-pair opening dynamics and interactions with ligands of nucleic acids in living human cells

Yudai Yamaoki, Takashi Nagata, Tomoki Sakamoto, Omar Eladl, Keiko Kondo, Masato Katahira Institute of Advanced Energy, Kyoto University/Graduate School of Energy Science, Kyoto University

RNA & RNA binding proteins

28P-073 Theoretical study on an enzymatic reaction of the hammerhead ribozyme

Ayaka Matsuyama, Masahiko Taguchi, Shigehiko Hayashi Kyoto University, Kyoto, Japan

DNA/RNA nanotechnology

29D-074

28P-074	RNA droplets perform 'AND' logic operation upon an input of targeted microRNAs
	Shin-ichiro M. Nomura. Yoshihiro Shimizu. Hirohide Saito.
	Masahiro Takinoue
	Tokyo Institute of Technology
28P-075	Use of aptamers to control nucleic acid phase separation <u>Samuel Hauf</u> , Yohei Yokobayashi Okinawa Institute of Science and Technology
28P-077	Experimental investigation of a modified Whiplash PCR driven by successive primer extension for massively parallel Implementation of DNA-based state machines Ken Komiya, Koji Sakamoto Japan Agency for Marine-Earth Science and Technology (JAMSTEC)

28P-078 Construction of giant unilamellar vesicle type molecular robots that uses cargo/component-holding DNA hydrogel Shoji Iwabuchi, Ryuji Kawano Tokyo University of Agriculture and Technology 28P-079 Dimeric DNA origami nanocapsules for controllable cargo accessibility

Yusuke Sakai, Joanna Markiewicz, Martyna Adamiak, Dmitry Ghilarov, Piotr Stepien, Jonathan G Heddle Malopolska Centre of Biotehcnology, Jagiellonian University, Poland

Chromatin & Chromosomes

28P-080 Brownian dynamics with exact solutions of diffusion in 3D for chromatin dynamics Yukitaka Ishimoto Grad. Sch. of Sci. Eng., Saga University

Water & Hydration & Electrolyte

28P-081	Nonthermal excitation effects of sub-terahertz radiation on transcription by RNA polymerase
	Masahiko Imashimizu AIST
28P-082	Role of hydration water on the stability of proteins

18P-082 Role of hydration water on the stability of proteins Mafumi Hishida Tokyo University of Science

 28P-083
 Liquid water structure by means of molecular dynamic simulation and machine-learning

 Taku Mizukami, Nguyen Viet Cuong, Dam Hieu Chi

 School of Materials Science, Japan Advanced Institute of Science and Technology

Morphogenesis and Development

28P-084 Epithelial Morphogenesis Analysis Using Texture Tensor Toshinori Namba, Kaoru Sugimura, Shuji Ishihara

Universal Biology Institute, The University of Tokyo, Tokyo, Japan/Graduate School of Arts and Sciences, The University of Tokyo, Tokyo, Japan

Molecular motor

28P-085	KIF6 is essential for male fertility through the ATP production pathway within sperm flagella <u>Tsukasa Makino</u> , Chizuru Ito, Takeshi Masuda, Kazuho Ikeda, Daisuke Takao, Yasushi Okada, Kiyotaka Toshimori, Masahide Kikkawa University of Tokyo
28P-086	Exploring Efficient Control of F1-ATPase
	Takahide Mishima, Yohei Nakayama, David A. Sivak, Shoichi Toyabe Department of Applied Physics, Graduate School of Engineering, Tohoku University
28P-087	Product inhibition slow down the moving velocity of processive chitinase and sliding-intermediate state blocks re-binding of product Yoshiko Tanaka, Takayuki Uchihashi, <u>Akihiko Nakamura</u> Shizuoka University/Institute for Molecular Science
28P-088	Decoding Volvox Swimming: Out-of-Phase Metachronal Waves Drive Oscillating Speed and Rotational Steering <u>Katsuya Shimabukuro</u> , Natsume Takeda, Tatsuya Suehiro, Naoki Uemura National Institute of Technology, Ube College
28P-090	The Force-Generating State of Myosin Detected by Quasielastic Neutron Scattering Satoru Fujiwara, Shinsaku Maruta, Yasunobu Sugimoto, Kai Nishikubo, Taiki Tominaga, Akio Inoue, Hidetaka Furuya, Katsuzo Wakabayashi, Toshiaki Arata National Institutes for Quantum Science and Technology
28P-091	Rotation-dependent inhibition and activation mechanism of ATPase inhibitory factor 1 for mitochondrial ATP synthase from atomistic simulation Ryohei Kobayashi, Kei-ichi Okazaki Research Center for Computational Science, Institute for Molecular Science, Aichi, Japan

28P-092 CryoEM structure of dimeric F1-like ATPase in Mycoplasma mobile suggests a rotary catalytic mechanism for the gliding motility <u>Takuma Toyonaga</u>, Takayuki Kato, Akihiro Kawamoto, Tomoko Miyata, Keisuke Kawakami, Junso Fujita, Tasuku Hamaguchi, Keiichi Namba, Makoto Miyata Grad. Sch. Sci., Osaka Metro. Univ., Osaka, Japan/OCARINA, Osaka Metro. Univ., Osaka, Japan/IMRAM, Tohoku Univ., Miyagi, Japan/AIMcS, Tohoku Univ., Miyagi, Japan

28P-093 Processive movement of myosin II HMM oligomers along actin filaments

<u>Taro QP Uyeda</u>, Hideya Hayashi

Department of Physics and Applied Physics, Graduate School of Advanced Science and Engineering, Waseda University

Single Molecule Biophysics

28P-094	CHEMICAL FRICTION ALONG THE MINOR GROOVE OF DNA FACILITATES ENZYMATIC TRANSLOCATION OF λ EXONUCLEASE VIA ELECTROSTATIC RATCHET Gwangrog Lee, Jungmin Yoo, HyeokJin Cho, Jejoong Yoo
28P-095	Single-molecule level tracking of the CCT/TRiC chaperonine mediated functional cycle
	<u>Kazutaka Araki</u> , Takahiro Watanabe-Nakayama, Daisuke Sasaki, Yuji Sasaki, Kazuhiro Mio OPERANDO-OIL, AIST, Chiba, Japan
28P-097	Multidisciplinary Platforms to Study Biological Questions Hsiufang Fan Institute of Biomedical Science, National Sun Yat-sen University, Taiwan
28P-098	deepFLUOR: Deep Learning Classification of Single-Molecule Signals Jinseob Lee, Byungju Kim, Yeongkyoung Park, Yoonki Kim, Jongbong Lee School of Interdisciplinary Bioscience and Bioengineering, POSTECH, Republic of Korea

Cell biology: Adhesion

28P-099	Stress propagation in a living cell
	Ayama Tokuyasu
	Grad. Sch. Nanobioscience, Yokohama city university, Kanagawa, Japan

28P-101 S100A11 promotes focal adhesion disassembly via myosin II-driven contractility and Piezo1-mediated Ca2+ entry Tareg Omer Mohammed, You-Rong Lin, Kai Weissenbruch, Kien Xuan Ngo, Yanjun Zhang, Noriyuki Kodera, Martin Bastmeyer, Yusuke Miyanari, Azuma Taoka, <u>Clemens M. Franz</u> WPI Nano Life Science Institute, Kanazawa University, Kanazawa, Japan

Cell biology: Motility

28P-102	Wave dynamics and collective behavior of swimming flagellar apparatus isolated from the green algae C. reinhardtii <u>Azam Gholami</u> , Sai Venkata Ramana, Albert Bae NYU Abu Dhabi, UAE
28P-103	Symbiotic bacteria break through narrow passage by flagellar wrapping Aoba Yoshioka, Tetsuo Kan, Kazutaka Takeshita, Hirofumi Wada, Yoshitomo Kikuchi, <u>Daisuke Nakane</u> Department of Engineering Science, The University of Electro-Communications
28P-104	The effect of TGF-β-induced EMT on the establishment of epithelial collective migration <u>Tomoaki Nagai</u> , Hirokazu Kaji, Michiru Nishita Department of Biochemistry, Fukushima Medical University School of Medicine
28P-106	Characteristics and mechanics of the crawling of the tested amoeba Arcella sp. Genta Matsumoto, Katsuhiko Sato, Toshiyuki Nakagaki, <u>Yukinori Nishigami</u> Graduate School of Life Science, Hokkaido University, Sapporo, Japan/Research Institute for Electronic Science, Hokkaido University, Sapporo, Japan

28P-107	Shape coupled bifurcation of an amoeba cell brings ballistic movement in amoeboid migration Hiroyuki Ebata, Yukinori Nishigami, Hisanori Fujiwara, Satoru Kidoaki, <u>Masatoshi Ichikawa</u> Department of Physics, Kyoto University, Japan
28P-108	Ciliary waveform conversion is induced by the shape change of doublet microtubule accompanied by the modification of outer-arm dynein motor activity <u>Toshiki Yagi</u> , Ai Sumiyoshi, Shogo Sawada Prefectural University of Hiroshima
28P-109	Viewing the swimming motion of a unicellular organism in extreme environmental conditions. Masayoshi Nishiyama Kindai University
28P-110	The Calcium Sensitive Helical Arrangement of Axonemal Components in Chlamydomonas Flagella Hitoshi Sakakibara, Kenta Ishibashi, Hiroyuki Iwamoto, Hiroaki Kojima, Kazuhiro Oiwa Bio-ICT Lab. NICT, Hyogo, Japan
28P-111	Utilizing Wavelet Analysis Features for the Simplified Prediction of Enhanced Cellular Stress Fluctuations on the Matrix with Stiffness Heterogeneity Satoru Kidoaki IMCE, Kyushu University
28P-112	Gliding direction of Mycoplasma mobile correlates with the curved configuration of its cell shape <u>Kana Suzuki</u> , Daisuke Nakane, Takayuki Nishizaka Gakushuin University
28P-113	Functional exploration of Candidatus Izimaplasma MreB using the minimal synthetic bacterium JCVI-syn3B. Ryu Takaishi, Mone Mimura, Hana Kiyama, Makoto Miyata Osaka Metropolitan University

Cell biology: Cytoskeleton & Membrane skeleton

- 28P-114Preparation of Dictyostelium discoideum NAA80 knockout strain
Tomoko Tsuji, Akira Nagasaki, Taro Uyeda
Department of Physics, Faculty of Advanced Science and Engineering, Waseda
University, Tokyo, Japan
- 28P-115
 Visualization of intracellular structure of D. discoideum during unicellular and multicellular phases

 Yuki Gomibuchi, Yukihisa Hayashida, Yusuke V. Morimoto, Takuo Yasunaga

Grad. Sch Comp. Sci and Sys. Eng., Kyushu Inst. Tech., Fukuoka, Japan

- 28P-116 Direct measurement of the physical properties of ER Mao Ikeda Yokohama City University
- 28P-117 Visualization of GTP hydrolysis in microtubules <u>Tomohiro Shima</u>, Hanjin Liu Graduate School of Science, The University of Tokyo
- 28P-118 Void space around microtubules <u>Shinji Kamimura</u>, Tomohiro Shima, Yasushi Okada, Hiroyuki Iwamoto Department of Biological Sciences, Faculty of Science and Engineering, Chuo University, Tokyo, Japan

Cell biology: Signal transduction & Cell membrane

- 28P-120 Application of Single-Molecule Tracking to Drug Discovery <u>Daisuke Watanabe</u>, Michio Hiroshima, Masahiro Ueda Laboratory of Single Molecule Biology, Graduate School of Frontier Biosciences, Osaka University; Suita, Osaka, Japan/Laboratory for Cell Signaling Dynamics, Center for Biosystems Dynamics Research, RIKEN; Suita, Osaka, Japan
- 28P-121 Investigation of cellular localization of opioid receptors: A combined biochemical assay and microscopy study Ming Chi Chen, Guan Yu Zhuo, Tzu Yu Lin, Shih Ting Lin, Daniel Tzu Li Chen, Cynthia Wei Sheng Lee Institute of Translational Medicine and New Drug Development, China Medical University, Taichung 40402, Taiwan

28P-122 Quantification of repellent response of single E. coli cell through the change in polar localization of adaptation enzyme CheB and flagellar motor rotation Hajime Fukuoka, Yumiko Uchida, Yong-Suk Che, Akihiko Ishijima Grad. Sch. Frontier Biosci., Osaka Univ. 28P-123 Lamellipodia-like membrane protrusions maintain the integrity of epithelial cell-cell junctions Yosuke Senju

Research Institute for Interdisciplinary Science (RIIS), Okayama University

28P-124 Unilateral-bidirectional regulation of electrical synapse formation in C. elegans

Zan Wu, Lin Pang, <u>Mei Ding</u> Institute of Genetics and Developmental Biology, Chinese Academy of Sciences

Biological & Artificial membrane: Structure & Property

- 28P-125 Curcumin Exerts the Membrane Raft Modulating Activity via Phase Separation and Induces CD44 Shedding in Tumor Cells <u>Toshiyuki Murai</u>, Yoshikazu Masaki, Kazuma Yasuhara Osaka University
- 28P-126 The effect of different lateral packing stress in acyl chains on KcsA orientation and structure in lipid bilayer Eri Saki H. Hayakawa, Misuzu Ueki, Elmukhtar Alhatmi, Shigetoshi Oiki, Fuyuki Tokumasu, Masayuki Iwamoto, Drake C. Mitchell Div. of Medical Zoology, Dept. of Infection and Immunity, Jichi Medical Univ.
- **28P-127** Probing the supramolecular aggregation state of bacterial endotoxin to reveal the basis of biological recognition and endotoxin masking in drug formulations

Andra B Schromm, Nicolas Gisch, Wilmar Correa, Walter Richter, Guillermo Martinez-de-Tejada, Klaus Brandenburg, Friedrich von Wintzingerode Division of Immunobiophysics, Research Center Borstel, Leibniz Lung Center, Borstel, Germany

Biological & Artificial membrane: Dynamics

28P-128 Non-equilibrium patterns in phase-separated lipid membranes under shear flow Tsutomu Hamada, Shino Mizuno, Hiroyuki Kitahata

Japan Advanced Institute of Science and Technology

Biological & Artificial membrane: Excitation & Channels

- 28P-129 Towards elucidating the tension effects on water flux across lipid bilayers and aquaporins: An attempt using water-in-oil microdroplets Misuzu Ueki, Takahisa Maki, <u>Masayuki Iwamoto</u> Dep. Mol. Neurosci., Facul. Med. Sci., Univ. Fukui
- 28P-130 Light-evoked channel activity using photolipids Rohit Yadav, Juergen Pfeffermann, Nikolaus Goessweiner-Mohr, <u>Peter Pohl</u> Johannes Kepler University, Linz, Austria

Membraneless Organella, autophage, Liquid-liquid phase separation

28P-131	The role of promyelocytic leukemia protein (PML) in the regulation of calcium homeostasis in HeLa cells Alexander Fonin, Rinat Sharipov Laboratory of Protein Structural Dynamics, Stability and Folding, Institute of Cytology, St. Petersburg, 194064, Russia
28P-132	Effect of F-actin, myosin and its fragments on the morphology and stability of PEG/DEX droplets <u>Tatsuyuki Waizumi</u> , Hiroki Sakuta, Mahito Kikumoto, Masahito Hayashi, Kanta Tsumoto, Kingo Takiguchi, Kenichi Yoshikawa Grad. Sch. Sci., Nagoya Univ.
28P-133	Translation-coupled genomic RNA replication in fibril-stabilized all-aqueous droplet colonies Ryo Mizuuchi, Hidekazu Sono, Keiji Murayama, Norikazu Ichihashi Department of Electrical Engineering and Bioscience, Faculty of Science and Engineering, Waseda University, Tokyo, Japan/JST, FOREST, Saitama, Japan

28P-134 PRESSURE AND TEMPERATURE EFFECTS ON FUS LIQUID DROPLET OF AMYOTROPHIC LATERAL SCLEROSIS PATHOLOGICAL VARIANT, R495X

Yutaro Shiramasa, Ryu Yamamoto, Fuka Sasaki, Soichiro Kitazawa, Tomoshi Kameda, <u>Ryo Kitahara</u> Graduate School of Pharmacy, Ritsumeikan University/College of Pharmaceutical Sciences, Ritsumeikan University

28P-136 Micropolarity governs the structural organization of biomolecular condensates

Songtao Ye, Chia-Heng Hsiung, Andrew Latham, Bin Zhang, <u>Xin Zhang</u> Department of Chemistry, Westlake University, 600 Dunyu Road, Hangzhou 310030, Zhejiang Province, China

Chemoreception

28P-137 Structural and biochemical analyses of SatA, a periplasmic binding protein involved in chemotactic response to serine with MIp3 in Vibrio cholerae. <u>Miyuki Aoyama</u>, Norihiro Takekawa, So-ichiro Nishiyama, Hirotaka Tajima, Ikuro Kawagishi, Katsumi Imada

Dept. of Macromol. Sci., Grad. Sch. of Sci., Osaka Univ.

Neuroscience & Sensory systems

 28P-138 Post-synaptic Effects of CPTX on Excitatory Synapses
 Boxiao Zhao, Akito Hattori, Shigeo Sakuragi, Hiroko Bannai, Michisuke Yuzaki
 School of Advanced Science and Engineering, Waseda University, Tokyo, Japan

 28P-139 Regulation of intracellular tau dynamics using optogenetic tools Shigeo Sakuragi, Akito Hattori, Boxiao Zhao, Yoshihiro Sakata, Gen Matsumoto, Akihiko Takashima, Yoshiyuki Soeda, Hideaki Yoshimura, Hiroko Bannai

Neuronal circuit & Information processing

- 28P-141 Fast Intrinsic Optical Signal (FIOS) measurements of brain slices: no-stain, label-free and non-invasive fast optical signal measurements Yoko Tominaga, Maki Koike-Tani, Tomomi Tani, Takashi Tominaga Inst. of Neurosci., Tokushima Bunri Univ., Sanuki, Japan
- 28P-142 Visualizing demyelination effects on interhemispheric communication with voltage-sensitive dye imaging in cuprizone-induced multiple sclerosis model mice

<u>Kyoka Tsukuda</u>, Michiko Miwa, Makiko Taketoshi, Yoko Tominaga, Kentaro Nakashima, Takashi Tominaga Grad. Sch. of Pharm. Sci., Tokushima Bunri Univ., Japan/Inst. of Neurosci., Tokushima Bunri Univ., Japan

Behavior

28P-143 The Implications of microRNA, CaMK2A, and MeCP2 Signaling on Adolescent Cognitive Ability <u>Ting-Kuang Yeh</u>, Li-Ching Lee, Pei-Jung Lin, Chun-Yen Chang National Taiwan Normal University

Photobiology: Vision & Photoreception

28P-144	The importance of water in membrane receptor function
	Anthony Watts
	Biochemistry Department, University of Oxford, Oxford, OX1 3QU UK

28P-145Time-resolved detections of substrate release and uptake reactions of
the light-driven chloride pump halorhodopsin
Chihaya Hamada, Keisuke Murabe, Takashi Tsukamoto, Takashi Kikukawa

Grad. Sch. Life Sci., Hokkaido Univ., Sapporo, Japan/Fac. Adv. Life Sci., Hokkaido Univ., Sapporo, Japan

28P-146 Molecular characterization of opsins from a nematode Keiichi Kojima, Yuki Tanioka, Keita Sato, Yosuke Nishimura, Susumu Yoshizawa, Hideyo Ohuchi, Takahiro Yamashita, Yuki Sudo Fac. Med. Dent. Pharm. Sci. Okayama Univ., Okayama, Japan

28P-147	Study on the Mechanisms of High Fluorescence of Archaearhodopsin-3 (AR3) Mutants Masae Konno, Krystyna Herasymenko, Stefan Haacke, Keiichi Inoue The Institute for Solid State Physics, University of Tokyo, Japan
28P-148	Analysis of the mechanism of photoreceptor RcPYP complex formation Yoichi Yamazaki, Yuna Kawabuchi, Kento Yonezawa, Sachiko Toma-Fukai, Hironari Kamikubo Division of Materials Science, Nara Institute of Science and Technology
28P-149	Driving force of proton pump rhodopsins revealed by electrophysiological study Satoshi Tsunoda, Akari Okuyama, Shoko Hososhima, Hideki Kandori Graduate School of Engineering, Nagoya Institute of Technology, Nagoya, Japan/ OptoBio Technology Research Center, Nagoya Institute of Technology, Nagoya, Japan
28P-150	Solid-state NMR characterization of histidine residues in Themoplasmatales archaeon heliorhodopsin Sari Kumagai, Toshio Nagashima, Toshio Yamazaki, Kota Katayama, Hideki Kandori, <u>Izuru Kawamura</u> Yokohama National University
28P-151	Characterization of the magnetic and geometrical structure of radical pairs in Serum Albumin by electron spin resonance <u>Hiroki Nagashima</u> , Masaki Kashiwazaki, Shuhei Arai, Kiminori Maeda Graduate School of Science and Engineering, Saitama University/Institute for Quantum Science and Technology, National institutes for Quantum Science and Technology
28P-152	Novel green/red light-sensing mechanism in the phytochrome- superfamily protein Takayuki Nagae, Yuya Fujita, Tatsuya Tsuchida, Takanari Kamo, Ryoka Seto, Masako Hamada, Hiroshi Aoyama, Ayana Sato-Tomita, Tomotsumi Fujisawa, Toshihiko Eki, Yohei Miyanoiri, Yutaka Ito, Takahiro Soeta, Yutaka Ukaji, Masashi Unno, Masaki Mishima, <u>Yuu Hirose</u>

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Poster Sessions

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Photobiology: Photosynthesis

28P-153 Molecular Docking Simulations at Quinone Binding Site in Photosynthetic Reaction Centers

<u>Ayumu Takagi</u>, Shigeru Itoh, Akihiro Kimura, Hirotaka Kitoh Grad. Sch. Sci. Eng., Kindai Univ.

28P-154 Fluorescence spectra of a photosynthetic carbonyl carotenoid, siphonaxanthin: Dual fluorescence observed only at ambient temperature in polar solvents

Kazuhiro Yoshida, Soichiro Seki, Yumiko Yamano, Tetsuichi Wazawa, Takeharu Nagai, <u>Ritsuko Fujii</u> Grad. Sch. Sci., Osaka Metropolitan Univ., Osaka, Japan/Grad. Sch. Sci., Osaka City Univ., Osaka, Japan/ReCAP, Osaka Metropolitan Univ., Osaka, Japan

28P-155 Cryo-EM structure of marine green algal LHCII utilizing blue-green light

Soichiro Seki, Tetsuko Nakaniwa, Pablo Castro-Hartmann, Kasim Sader, Akihiro Kawamoto, Hideaki Tanaka Tanaka, Qian Pu, Genji Kurisu, Ritsuko Fujii

Division of Molecular Materials Science, Graduate School of Science, Osaka City University, 3-3-138, Sugimoto, Sumiyoshi-ku, Osaka 558–8585, Japan

28P-156 Biohydrogen production from whiskey waste liquid by two-stage fermentation

<u>Masahiro Hibino</u>, Kousei Miyamoto Div. Sust. Enviro. Eng., Muroran Inst. Tech.

28P-157 Energy gradient of the β82 chromophores established by the linker proteins in Synechocystis PCC 6803 Phycobilisome Rod Hiroto Kikuchi

Dept. of Phys., Nippon Medical School

28P-158 Estimation of Local Antenna Sizes of Photosystem I in Chlamydomonas Cells

Xianjun Zhang, Yuki Fujita, Rin Taniguchi, Shen Ye, <u>Yutaka Shibata</u> Department of Chemistry, Graduate School of Science, Tohoku University

Photobiology: Optogenetics & Optical control

- 28P-159 The effect of blue light on the proliferation of E. coli cells <u>Nagomi Matsumoto</u>, Osamu Hisatomi Graduate School of Science, Osaka University, Toyonaka, Osaka, Japan
- 28P-160 Photocontrol of small GTPase Ras fused with a photoresponsive protein Nobuyuki Nishibe, Ziyun Zhang, Kazunori Kondoh, Shinsaku Maruta Department of Biosciences, Graduate School of Science and Engineering, Soka University Hachioji, Tokyo Japan
- 28P-161 Electrophysiological characterization of light-activated protontransporting heliorhodopsins

Shoko Hososhima, Satoshi Tsunoda, Hideki Kandori Department of Life Science and Applied Chemistry, Nagoya Institute of Technology, Showa-ku, Nagoya 466-8555, Japan/OptoBioTechnology Research Center, Nagoya Institute of Technology, Showa-ku, Nagoya 466-8555, Japan

28P-162 Reconstitution of a light-activatable transcription factor, Photozipper, with extrinsic chromophores

Osamu Hisatomi, Nagomi Matsumoto Graduate School of Science, Osaka University, Toyonaka, Osaka, Japan

Radiobiology & Active oxygen

- 28P-163 YAP/Aurora A-mediated ciliogenesis regulates ionizing radiationinduced senescence via Hedgehog pathway in tumor cells Jinpeng He, Wei Ma, Li Wei, Tongshan Zhang, Junrui Hua, Jufang Wang Institute of Modern Physics, Chinese Academy of Sciences/University of Chinese Academy of Sciences
- 28P-164 Cadmium tolerance, accumulation and translocation in sweet sorghum irradiated by carbon beam Xicun Dong Institute of Modern Physics, Chinese Academy of Sciences

Origin of life & Evolution

28P-165	Analysis of evolutionary constraints using bacterial experimental evolution Astushi Shibai, Sumpei Sato, <u>Chikara Furusawa</u> Center for Biosystems Dynamics Research, RIKEN/Universal Biology Institute, The University of Tokyo
28P-166	Mega-phylogenetic evolution of complex adaptive traits in thousands of bacterial species <u>Takao K Suzuki</u> , Wataru Iwasaki Graduate School of Frontier Sciences, the University of Tokyo
28P-167	Quantum evolution form electronic state of macro-biomolecules Masanori Yamanaka Nihon University
28P-168	The Relation Between Biology and Physics: Origins of Life Research and its Philosophical Implications Julieta Macome History and Philosophy of Science Department, Cambridge University

Synthetic biology & Artificial cells

28P-169	Spatially separated transcription and translation in the artificial cell with the artificial organelle
	<u>Kanji Tomohara,</u> Yoshihiro Minagawa, Hiroyuki Noji
	Graduate School of Engineering, The University of Tokyo, Tokyo, Japan
28P-170	Synthetic minimal cells with various vesicle-polymer compositions: Toward implementing evolution
	Department of Physics, Graduate School of Science, Tohoku University, Japan.
28P-171	Molecular tools aiming at arbitrary manipulation of micro-structures in living cells Hideki Nakamura
	Hakubi Center for Advanced Research, Kyoto University, Kyoto, Japan,/School of Engineering, Kyoto University, Kyoto, Japan/JST PRESTO, Supra-Assembly of Biomolecule, Tokyo, Japan

28P-172 Phase separation-induced actin bundle elongates filopodia-like tube on giant liposome from inside Masahito Hayashi, Tomoyuki Kaneko LaRC, Dept. Frontier Biosci. Hosei Univ., Tokyo, Japan/LaRC, FB, Grad. Sch. Sci. & Eng., Hosei Univ Tokyo, Japan 28P-173 Pattern diversity emerges from a simple gene network Xiongfei Fu, Jingwen Zhu, Pan Chu Key Laboratory for Quantitative Synthetic Biology, Shenzhen Institute of Synthetic

Key Laboratory for Quantitative Synthetic Biology, Shenzhen Institute of Synthetic Biology, Shenzhen Institutes of Advanced Technology, Chinese Academy of Sciences, Shenzhen 518055, China/University of Chinese Academy of Sciences, Beijing 100049, China

Genome biology

28P-174 Insights into chromatin organization obtained by a rapid classification of A/B compartments from Hi-C data <u>Takashi Sumikama</u>, Hisashi Miura, Ichiro Hiratani, Takeshi Fukuma PERSTO, JST/Kanazawa University

Computational biology: Bioinformatics

28P-175	Computational Evaluation of the Human Health Effects of the Main Compounds Found in Artemisia dracunculus
	<u>Andrijana Pujicic,</u> Adriana Isvoran
	Department of Biology-Chemistry, West University of Timisoara, 16 Pestalozzi, 300115 Timisoara
28P-178	Development of Prediction Methods for Class A GPCR and G-protein Coupling Selectivity Using Deep Learning
	<u>Kento Fujishima</u> , Kenji Etchuya, Ikuo Masuho, Makiko Suwa
	Biol. Sci., Grad. Sci. Eng., Aoyama Gakuin Univ., Kanagawa, Japan.
28P-179	Consortium of "Consistent substitutions" on Influenza A(H1N1) viral proteome and its possible consequences on human host-viral interactions: A study using Multiple Sequence Alignments, text mining, and Molecular Dynamics Simulations
	<u>Debashree Bandyopadhyay,</u> Syeda Lubna
	Birla Institute of Technology and Science, Pilani, Hyderabad Campus

28P-180 Analysis of Partial Structural Similarity of ribonuclease and chymotrypsin based on their amino acid sequences <u>Takeshi Kikuchi</u>, Ahasanul Kabir, Takuya Takahashi Ritsumeikan University

Computational biology: Molecular simulation

28P-181	Gas-phase Structural Analysis of Biomolecules using Coarse-grained Molecular Dynamics Simulation Kazumi Saikusa, Satoko Akashi, <u>Sotaro Fuchigami</u> School of Pharmaceutical Sciences, University of Shizuoka
28P-183	2D-replica exchange simulation of membrane permeation process of cyclic hexapeptides <u>Tsutomu Yamane</u> , Masateru Ota, Mitsunori Ikeguchi Riken Center for Computational Science
28P-184	Classification of conformational dynamics of high mannose- type oligosaccharides by molecular simulation and data clustering <u>Yue Zhang</u> , Takumi Yamaguchi School of Materials Science, Japan Advanced Institute of Science and Technology
28P-185	The role of water and cholesterols in APP cleavage by gamma- secretase <u>Jinyoung Byun</u> , Juyong Lee College of Pharmacy, Seoul National University, Seoul, Republic of Korea
28P-186	Decoupling processes of the Adenosine A2A receptor from G-proteins through the lens of dPaCS-MD simulations Duy Phuoc Tran, Louis-Philippe Picard, Alexander Orazietti, Sari Hagimoto, Adnan Sljoka, R. Scott Prosser, Akio Kitao School of Life Science and Technology, Tokyo Institute of Technology, Tokyo 152- 8550, Japan
28P-187	PROTAC-mediated ternary complex structure distribution profiles using enhanced sampling methods Genki Kudo, Takumi Hirao, Takatsugu Hirokawa, Ryunosuke Yoshino Physics Department, Graduate School of Pure and Applied Sciences, University of Tsukuba

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28P-188 3D Structure Prediction of the Odorant-bound Olfactory Receptor Takumi Hirao, Yusuke Ihara, Chiori Ijichi, Genki Kudo, Ryunosuke Yoshino, Takatsugu Hirokawa Doctoral Program in Medical Sciences, Graduate School of Comprehensive Human Sciences, University of Tsukuba 28P-189 Quantitative Evaluation of Protein-Compound Substructure Interaction with Inverse Mixed-Solvent Molecular Dynamics Simulation Keisuke Yanagisawa, Ryunosuke Yoshino, Genki Kudo, Takatsugu Hirokawa Department of Computer Science, School of Computing, Tokyo Institute of Technology/Middle Molecule IT-based Drug Discovery Laboratory (MIDL), Tokyo Institute of Technology

28P-190 Binding Free Energy Shifts of Protein Complexes due to Amino Acid Mutations

Kazutomo Kawaguchi, Hidemi Nagao Institute of Science and Engineering, Kanazawa University

28P-191 Computational estimation of the free energy change of peptide- bond rotation induced by reduction of "plant-type" ferredoxin

Tomoki Nakayoshi, Yusuke Ohnishi, Hideaki Tanaka, Genji Kurisu, Yu Takano Graduata School of Information Sciences, Hirochima City, University/Faculty

Graduate School of Information Sciences, Hiroshima City University/Faculty of Pharmacy, Meijo University

28P-192 Dependence of the abnormal open states patterns in the ATXN2 gen on the number of CAG repeats

<u>Stepan Dzhimak</u>, Mikhail Drobotenko, Oksana Lyasota, Jose Luis Hernandez-Caceres, Yuriy Nechipurenko, Alexandr Svidlov, Anna Dorohova

Laboratory of Problems of Stable Isotope Spreading in Living Systems, Southern Scientific Center of the Russian Academy of Sciences, Rostov-on-Don, Russia/ Department of Radiophysics and Nanothechnology, Kuban State University, Krasnodar, Russia,

28P-193 GPU-accelerated coarse-grained MD simulator and its application to postsynaptic density

<u>Yutaka Murata</u>, Shoji Takada Dept. Biophysics, Div. Biology, Grad. Sch. of Science, Kyoto University

28P-195	Vibrational Dynamics of Water Molecules in FUS Protein Condensates: Molecular Interpretation <u>Yotaro Takeda</u> , Tatsuya Ishiyama, Eiji Yamamoto Department of System Design Engineering, Keio University, Japan
28P-196	Collagen-collagen interactions: Triple helix to helix-helix to fibrils. <u>George Anthony Pantelopulos</u> , Robert Best National Institutes of Health
28P-197	Analysis of Antigen-Antibody Interface Based on MD Simulations: Toward Antibody Design Takefumi Yamashita Hoshi University/The University of Tokyo
28P-198	Simulation-based prediction and elucidation of the pathogenic mechanism of deafness in GJB2-encoded Cx26 channel protein <u>Cheng-Yu Tsai</u> , Ying-Chang Lu, Yen-Hui Chan, Chuan-Jen Hsu, Pei-Lung Chen, Chen-Chi Wu, Lee-Wei Yang Graduate Institute of Medical Genomics and Proteomics, National Taiwan University College of Medicine, Taipei, Taiwan/Department of Otolaryngology, National Taiwan University Hospital, Taipei, Taiwan
28P-199	Applications of Tree-Search-MD to drug target proteins: conformational changes between inactive and active structures of a kinase and a ligand binding to a GPCR Yukina Nakai, <u>Toru Ekimoto</u> , Tsutomu Yamane, Masao Inoue, Naoki Ogawa, Sun-Yong Park, Kei Terayama, Mitsunori Ikeguchi Yokohama City University
28P-200	Theoretical Insights into Drug Resistance Mechanisms of HIV-1 Protease: Residue Interaction Network Analysis Keidai Yamase Chiba Institute of Technology
28P-201	Molecular dynamics simulations of lipid adsorption by PLA2 of snake venom. <u>Tatsuhiro Kawashima</u> , Ryuta Imayoshi, Kazutomo Kawaguchi, Hidemi Nagao Graduate School of Natural Science and Technology, Kanazawa University, Japan

28P-202 Analysis of Protein Simulations Using Relaxation Mode Analysis Avori Mitsutake Meiji University 28P-203 How the Membranes Fuse: From Spontaneous to Induced Hongxia Guo Institute of Chemistry, Chinese Academy of Sciences 28P-204 Impact of glycosylation on the structural features and hydration effects of glycoproteins Haeri Im, Song-Ho Chong, Isseki Yu, Yuji Sugita RIKEN Cluster for Pioneering Research, Wako, Japan 28P-205 Developing an IDP-Specific Force Field by Optimizing CMAP **Parameters** Haozhe Guo, Chen Song Peking-Tsinghua Center for Life Sciences, Academy for Advanced Interdisciplinary Studies, Peking University, Beijing, China/Center for Quantitative Biology, Academy for Advanced Interdisciplinary Studies, Peking University, Beijing, China 28P-206 Coarse-Grained Co-transcriptional Folding Simulation of RNA Switch Akito Taneda Hirosaki Universitv

Computational biology: Biological modeling and simulation

28P-207	Dynamic transitions in microtubules: role of flared ends and lattice repair in catastrophes and rescues
	Nikita Gudimchuk, <u>Veronika Aleksandrova</u> , Mikhail Anisimov
	Lomonosov Moscow State University, Moscow, Russia
28P-208	Analysis of fracture patterns in a vertex model including detachment
	of cells
	<u>Nozomi Fujita,</u> Yuichi Togashi
	Ritsumeikan University
28P-210	Learning force field parameters from ensemble-averaged data with a
	differentiable approach
	<u>Yohei Sako</u> , Yasuhiro Matsunaga
	Graduate School of Science and Engineering, Saitama University, Saitama, Japan.

28P-211	Controlled Drug Delivery from Polymeric Surfaces: Harnessing Sonochemical Methods for Fluorouracil Nanoparticle Synthesis Paulina Chytrosz-Wrobel, Monika Golda-Cepa, Piotr Kubisiak, <u>Waldemar Kulig</u> , Lukasz Cwiklik, Andrzej Kotarba Department of Physics, University of Helsinki, Helsinki, Finland
28P-212	Molecular modelling, homo-oligomerisation and membrane interactions of hepatitis E virus pORF1 replication polyprotein <u>Thibault Tubiana</u> , Sonia Fieulaine, Stéphane Bressanelli Université Paris-Saclay, CEA, CNRS, Institute for Integrative Biology of the Cell (I2BC), 91198, Gif-sur-Yvette, France
28P-213	Improving Structure-Based Virtual Screening using AlphaFold2 with Multi-State Modeling Woong-Hee Shin, Jinung Song, Junsu Ha, Juyong Lee, Junsu Ko Department of Biomedical Informatics, Korea University College of Medicine
28P-214	Flexible Fitting of Coarse-Grained Models to AFM Images of Intrinsically Disordered Proteins <u>Sakura Homma</u> , Yasuhiro Matsunaga Saitama University
28P-215	Structure formations induced by a non-reciprocal cell-cell interactions in a multicellular system. <u>Biplab Bhattacherjee</u> , Masayuki Hayakawa, Tatsuo Shibata Laboratory for Physical Biology, RIKEN Center for Biosystems Dynamics Research, Kobe, Japan.
28P-216	Universal existence of power-law correlations in homogeneous states of anisotropic active matter models <u>Hiroyoshi Nakano</u> , Kyosuke Adachi Institute for Solid State Physics, University of Tokyo
28P-217	Searching for the BET interactome through AI and Molecular dynamics simulations Alberto Perez University of Florida/Riken visiting JSPS fellow
28P-219	Membrane fusion as a pathway to fission Russell k w Spencer, Marcus Müller Georg-August Universitat Goettingen

Poster Sessions

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Computational biology: machine learning for molecules or cell systems

28P-220 Optimization of In Vitro Glycolytic Pathway Using Machine Learning Naosato Takagi, Daisuke Kiga Waseda University, Tokyo, Japan

28P-221 Development of the super-resolution Cryo-EM based on the Generative Adversarial Networks Xinyuan Li, Takayuki Kato Graduate School of Frontier Biosciences (FBS), Osaka University

28P-222 Deep Learning-Based Water Molecule Displacement Prediction Method for Improving the Accuracy of Drug Discovery Docking Software

> Yuki Ito, Masateru Ohta, Mitsunori Ikeguchi, Takashi Yoshidome Department of Applied Physics, Graduate School of Engineering, Tohoku University, Japan

28P-223 Construction of a physical reservoir computing device using active matter made from a swarm of biomolecular motors <u>Yiming Gong</u>, Gikyo Usuki, Arif Md. Rashedul Kabir, Kazuki Sada, Ibuki Kawamata, Nathanael Aubert-Kato, Masatoshi Ichikawa, Akira Kakugo Graduate School of Science, Kyoto University, Kyoto, Japan

Mathematical & Theoretical biology

28P-224 Spatial point processes with molecular density-dependent association-dissociation and cluster formation in signal transduction on the plasma membrane Hiroaki Takagi Nara Medical University, Nara, Japan

28P-226 Professor

Chanho Park, Junil Kim, <u>Julian Lee</u> Department of Bioinformatics and Life Science, Soongsil University

Data Sharing and Open Science

- 28P-227 Crafting an Individual-Centric Genomics Platform Senkei Umehara, Atsushi lida, Ken Yagi GENEX, Inc. (Tokyo, Japan)
- 28P-228 Serious accumulation of sequence errors in international public database searched by analyzing deposited plasmids in RIKEN-BRC Gene bank with high-throughput sequencing Yoshihiro Miwa, Tetsushi lida, Junko Kijima, Shingo Nozaki, Shotaro Kishikawa Gene-Eng-Div, BRC, RIKEN

Ecology & Environment

28P-229 Ionic-strength and pH dependent reactivities of ascorbic acid and cysteine toward ozone in microdroplets studied by aerosol optical tweezers Yuan-Pin Chang

Department of Chemistry, National Sun Yat-sen University, Kaohsiung, Taiwan.

Nonequilibrium state & Biological rhythm

28P-230	From cellular chirality to large-scale chirality: Emergence of chiral spiral in migrating cellular system
	Masayuki Hayakawa, Biplab Bhattacherjee, Lihao Guo,
	Hidekazu Kuwayama, Tatsuo Shibata
	Laboratory for Physical Biology, RIKEN Center for Biosystems Dynamics Research, Kobe, Japan
28P-231	Integrated Analysis of Circadian Clock in cyanobacteria
	<u>Masaaki Sugiyama,</u> Ken Morishima, Yasuhiro Yunoki, Rintaro Inoue
	Institute for Integrated Radiation and Nuclear Science, Kyoto University
28P-232	Computational Study of Peak Position in One Dimensional
	Mesoscopic Reaction Diffusion System
	<u>Ryuta Imayoshi</u> , Kazutomo Kawaguchi, Hidemi Nagao
	Graduate School of Natural Science and Technology, Kanazawa University

Measureme	nts
28P-234	Extracellular Potential Measurement of Cardiomyocytes in Hyperkalemic Conditions Kentaro Kito, Masahito Hayashi, <u>Tomoyuki Kaneko</u> LaRC, FB, Grad. Sch. Sci. & Eng., Hosei Univ., Tokyo, Japan
28P-235	Single Molecule Analysis of Perforin Dynamics Using Nanopore Measurements. <u>Sotaro Nakamura</u> , Kazuhiro Kobayashi, Ryo lizuka, Hideaki Kato, Sotaro Uemura The University of Tokyo
28P-236	IR super-resolution micro-spectroscopy of keratin proteins in human nails <u>Ayaka Nagaoka</u> , Hirona Takahashi, Tetsuya Ida, Makoto Sakai Okayama University of Science
28P-237	Measurement of photocatalytic hydrogen production in titanium/ manganese oxide film/hydrogenated amorphous silicon thin film stack using flavan molecules <u>Yutaka Tsujiuchi</u> , Kohei Saito, Kazunori Takada, Koyu Akiyama, Hiroshi Masumoto Akita UNIV/Tohoku UNIV
28P-238	Current control using external blue-green light in an amino acid- containing gel stacked device in contact with a hydrogenated amorphous silicon thin film

Kohei Saito, Yutaka Tsujiuchi, Hiroshi Masumoto

Akita UNIV

Density-dependent state transitions and periodic advective flow in an

active actomyosin system

Tomoka Kashiwabara, Yusuke T. Maeda Dept. of Phys. Kyushu Univ., Fukuoka, Japan

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Poster Sessions

Bioimaging	
28P-240	Coupling between vinculin and retrograde actin flow visualized by live-cell single-molecule imaging Ying Liu, Naoki Watanabe, Sawako Yamashiro Laboratory of Single-Molecule Cell Biology, Kyoto University Graduate School of Biostudies, Kyoto, Japan
28P-241	Data-Driven Approaches in Single-Molecule Trajectory Analysis of Protein Mobility in Live Cells Yuma Ito, Makio Tokunaga School of Life Science and Technology, Tokyo Institute of Technology
28P-242	Monitoring the biofilm development of Escherichia coli BL21 Alexander Karl Bullen, Tomohiro Shima University of Tokyo
28P-243	Label-free direct screening of "spectral biomarkers" of colorectal tumour-specific P. anaerobius via Raman mapping in combination with data mining <u>Pooja Manik Badgujar</u> , Yu-Chung Yu-Chung Lin, Zhe-Rui Zhe-Rui Lin, Kuan-Ting Wu, Chia-Liang Cheng Department of Physics, National Dong Hwa University, Hualien 97401, Taiwan
28P-244	Revisiting the 105 gap issue in cellular thermal biology by label-free mid-infrared photothermal microscopy <u>Keiichiro Toda</u> , Masaharu Takarada, Genki Ishigane, Hiroyuki Shimada, Venkata Ramaiah Badarla, Kohki Okabe, Takuro Ideguchi The Univ. of Tokyo (Science), Tokyo, Japan
28P-245	Visualization of exocytosis using video-rate bioluminescence imaging Satoru Yokawa, Shinji Fukuda, Takahiro Suzuki, Tadahide Furuno Department of Analytical Chemistry and Biophysics, School of Pharmacy, Aichi Gakuin University
28P-246	Topography considerations for high-speed atomic force microscopy based force mapping on bacteria <u>Christian Ganser</u> , Shigetaka Nishiguchi, Takayuki Uchihashi National Institutes of Natural Sciences, ExCELLS

Poster Sessions

June 28 [Fri]

28P-247	Optical Freshness Evaluation Method of Raw Fish Meat Yasuhiro Maeda, Go Shioi, Tomonobu Watanabe Laboratory for Comprehensive Bioimaging, RIKEN, BDR, Japan
28P-248	Nanopipette-based single-cell stimulation with non-thermal atmospheric pressure plasma <u>Han Gia Nguyen</u> , Linhao Sun, Shinya Kumagai, Shinji Watanabe Grad. Sch. Nano Life Sci., Kanazawa University, Japan
28P-249	The role of receptor oligomerization in signal transduction investigated through single-molecule analysis <u>Hideaki Yoshimura</u> , Takeaki Ozawa Department of Chemistry, School of Science, The University of Tokyo, Tokyo, Japan
28P-250	Real-Time Imaging of Granzyme Secretion During CTL Assault on Cancer Cells Zhuohao Yang, Yuto Kurisu, Koji Nagaoka, Kazuhiro Kakimi, Takashi Funatsu, Yoshitaka Shirasaki RCAST, Univ. Tokyo
28P-251	Advancing Severe Asthma Research through Live-Cell Imaging of Secretion Activity Yoshitaka Shirasaki, Mai Yamagishi, Kaede Miyata, Yumiko Tanaka, Hiroki Kabata, Misato Irie, Rie Baba, Takashi Kamatani, Kazuyo Moro, Koichi Fukunaga, Sotaro Uemura Research Center for Advanced Science and Technology, The University of Tokyo
28P-252	Imaging of mitochondrial ATP in mouse sperm before and after capacitation using AMPK activators and inhibitors <u>Takashi W Ijiri</u> , Yuika Asanuma, Masamichi Yamamoto Setsunan University
28P-253	Numerous-color simultaneous imaging with dozens of bioluminescence colors <u>Mitsuru Hattori</u> , Yuki Hiruta, Takeharu Nagai SANKEN, Osaka University, Japan

Bioengineering

28P-255	Spiral Formation of Microtubules Driven by Kinesin Motors
	<u>Douglas Ng'ang'a,</u> Takahiro Nitta
	Applied Physics Course, Faculty of Engineering, Gifu University

28P-256 Sensitive detection of Salmonella with CRISPR–Cas13a system <u>Svitlana Kovalchuk</u>, Yoshihiro Minagawa, Hiroyuki Noji The University of Tokyo, Tokyo, Japan/National University of Food Technologies, Kyiv, Ukraine

28P-257 Spectroscopic signature responsible for the life activity of regenerating worm A. viride studied using Raman spectroscopy and Two-Photon Fluorescence Lifetime Imaging

<u>Chia-Liang Cheng</u>, Pooja Badgujar, Pei-Yang Huang, Artashes Karmenyan, Viktor Nikolayev, Jiun-Hong Chen Department of Physics, National Dong Hwa University, Hualien 97401, Taiwan

28P-258 Anticancer peptides delivery systems effects on model and natural lipid membranes

Bogdan Zorila, Diana Lavinia Stan, Roberta (Stoica) Moisa, <u>Mihaela Bacalum</u> Department of Life and Environmental Physics, Horia Hulubei National Institute for Physics and Nuclear Engineering

28P-259 Spatiotemporal changes in single cell rheology of developing embryos unveiled by atomic force microscopy <u>Takahiro Kotani</u>, Yuki Miyata, Yosuke Tsuboyama, Yuki Fujii, Takaharu Okajima Graduate School of Information Science and Technology, Hokkaido University

Crystal growth & Crystallization technique

28P-260 Assembly of Cage-Shaped Protein Dps Using Functional Peptides <u>Mitsuhiro Okuda</u>, Gabriela Pretre Meiji University/CIC-nanoGUNE/Komie Corp.

Virus structure, function, SARS-CoV-2

28P-261 Rational in silico design and structure analysis of SARS-CoV-2 neutralizing antibody UT28K

<u>Shunsuke Kita</u>, Tatsuhiko Ozawa, Kouki Ikeda, Liuan Chen, Yuki Anraku, Hideo Fukuhara, Emiko Igarashi, Yumiko Saga, Noriko Inasaki, Jiei Sasaki, Yuhei Kirita, Takao Hashiguchi, Hideki Tani, Hiroyuki Kishi, Hideki Niimi, Katsumi Maenaka

Facul. Pharm. Sci., Hokkaido Univ., Japan

28P-262 Verification of the effect of ligand and receptor flexibility on inhibitory activity by MD simulation

<u>Suzuka Saito</u>, Masashi Muramoto, Simon Hikiri, Junichi Higo, Takuya Takahashi Graduate School of Life Sciences, Ritsumeikan University, Kusatsu, Japan.

28P-263 N-substituted anthranilic acid derivatives as PPI inhibitors between Syntenin-1 PDZ domain and SARS-CoV-2 Env protein

<u>Hidekazu Hiroaki</u>, Ryusei Hamajima, Youichi Suzuki, Eiji Morita, Hong Wu, Yoshihiko Fujioka, Takeshi Tenno Graduate School of Pharmaceutical Sciences, Nagoya University/Center for One

Medicine Innovative Translational Research, Tokai National Higher Education and Research System/BeCellBar LLC, Nagoya, Aichi, Japan

28P-264 Molecular mechanisms of SARS-CoV-2 resistance to nirmatrelvir and the countermeasures

Haitao Yang Shanghai Institute for Advanced Immunochemical Studies, ShanghaiTech University, Shanghai, China

Mechanosensing and Mechanobiology, Biological Temperature

28P-265 Analysis on the Role of the Periplasmic Loop of the Bacterial Mechanosensitive Channel MscL Yasuyuki Sawada, Takeshi Nomura, Masahiro Sokabe

Institute of Materials Innovation, Institutes of Innovation for Future Society, Nagoya University

28P-266	Activation of chloride ion channel CLIC1 by mechanical external force using AFM in breast cancer cell <u>Ayana Yamagishi</u> , Samrat Mukherjee, Chikashi Nakamura National Institute of Advanced Industrial Science and Technology (AIST)/Tokyo University of Agriculture and Technology
28P-267	Mechanical properties of nestin tail domain analyzed by tensile test using AFM Ayana Yamagishi, Rina Tokuoka, Daijiro Takeshita, Chiaki Yoshikawa, Tomohiko Yamazaki, Taro Uyeda, <u>Chikashi Nakamura</u> AIST/Tokyo University of Agriculture and Technology
28P-268	Elucidation of the mechanism of intracellular temperature variation by high-speed temperature mapping Masaharu Takarada, Takashi Funatsu, <u>Kohki Okabe</u> Graduate School of Pharmaceutical Sciences, The University of Tokyo
Biophysics of	of disease
28P-269	Changes in the properties of rbcs in the process of extracorporeal membrane oxygenation by scanning flow cytometry

Ekaterina Yastrebova, Valeri Maltsev, Gleb Moroz V.V. Voevodsky Institute of Chemical Kinetics and Combustion of the Siberian Branch of the RAS

28P-270 Morphology of Cancer Organoids Reproduced by 3D Phase-Field Model

<u>Kotaro Kawamura</u>, Toshikaze Chiba, Keita Yanagiya, Yutaka Oya, Toshihiro Kawakatsu, Tatsuaki Tsuruyama, Masayuki Imai Department of Physics, Tohoku University, Aoba, Sendai, Japan

Miscellaneous topics

28P-271 Investigation of boson peak like behaviors appeared in cysteine and related amino acids <u>Hirofumi Nema</u>, Yasuhiro Fujii, Akitoshi Koreeda Ritsumeikan University

28P-272 Interactions of Model Antimicrobial Peptides with Lipid Membranes Normand Voyer, Pierre-Alexandre Paquet-Côté, François Otis, Jochen Bürck, Patrick Lagüe, Anne S. Ulrich 1Département de chimie and PROTEO, Université Laval, Québec, Canada