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**

*27P-001	Structural Basis for the Functional Diversity in Mechanosensitive Channel OSCAs <u>Kio Horinaka</u> , Tatsuya Hagino, Tsukasa Kusakizako, Osamu Nureki Department of Biological Sciences, Graduate School of Science, The University of Tokyo, Tokyo, Japan
*27P-002	Development of an Oxidative Folding Promoter by Controlling Protein Recognition Properties Koki Suzuki, Ryoya Nojiri, Tomohide Saio, Takahiro Muraoka School of Engineering, Tokyo University of Agriculture and Technology
*27P-003	How well do Alphafold2 structures perform in Molecular Docking? Ben Hanks, John Douglas Tanner, Ben Corry Australian National UNiversity
*27P-004	<b>Cryo-EM Structure Analysis of hOCT2, Organic Cation Transporter 2</b> <u>Haruna Inuzuka</u> , Yongchan Lee, Tomohiro Nishizawa Yokahama City University
*27P-005	Magnetic field effects on structure of iron sulfur protein studied by EPR and SAXS Shogo Soga, Ryoma Kobayashi, Hirokazu Masai, Shinji Kohara, Kiminori Maeda, Mitsuhiro Hirai, Hiroki Nagashima, Shigeki Arai Graduate School of Science and Engineering, Saitama University, Saitama, Japan

*27P-006	Analysis of the aggregation characteristics of tau droplets under oxidizing and reducing conditions Yuki Michiue, Ayumi Masui, Keisuke Yuzu, Yumiko Ohhashi, Keiichi Yamaguchi, Yasushi Kawata, Eri Chatani Grad. Sch. Sci,. Kobe Univ
*27P-007	Determination of the hemocyanin structure from Concholepas concholepas using an X ray crystallography and Cryo EM combined approach Sebastian Manuel Muñoz, Michelle Salazar, Gabriel Vallejos, Augusto Manubens, Mathias Ellena, José Edwin Quesñay, Andre Ambrosio, Maria Inés Becker, Victor Castro-Fernandez, Victoria Guixé Laboratorio de Bioquímica y Biología Molecular, Facultad de Ciencias, Universidad de Chile. Santiago, Chile.
*27 <b>P-008</b>	Designing Self-assembling Protein Nanoparticle using computational method <u>JinWoong Song</u> , SeaHae Choi, Junsu Ko, Won-Kyu Lee, Juyong Lee College of Pharmacy, Seoul National University, Seoul, Republic of Korea
*27P-009	Structure-based discovery of dual pathway inhibitors for SARS-CoV-2 entry Haofeng Wang ShanghaiTech University
27 <b>P-010</b>	Efficient design of allosteric activators for Rsp5 E3 ligase using machine-learning tool ProteinMPNN Wei-Lin Lu Institute of Biological Chemistry, Academia Sinica
27P-011	<b>Structural analysis of dissimilatory sulfate reductase</b> Rio Hamada, Koji Nishikawa, <u>Hideaki Ogata</u> University of Hyogo

## 27P-013 Approach to in situ structural analysis using JEOL's Cryo-FIB-SEM and CRYO ARM

Tomoko Miyata, Miki Kinoshita, Fumiaki Makino, Yoshie Kushima, Reiko Yamauchi, Keiichi Namba Graduate School of Frontier Biosciences, Osaka University/JEOL YOKOGUSHI Research Alliance Laboratories, Osaka University

27P-014 Cryo-EM structure of full-length cargo receptor ERGIC-53 in complex with MCFD2

<u>Satoshi Watanabe</u>, Yoshiaki Kise, Kento Yonezawa, Mariko Inoue, Nobutaka Shimizu, Osamu Nureki, Kenji Inaba Tohoku University

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#### **Protein: Structure & Function**

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277-013	<u>using HS-AFM and computational simulation</u> <u>Yamamoto Sohma</u> , Kazusa Takeda, Holger Flechsig, Hiroki Konno College of Science and Engineering, School of Biological Science and Technology,
*27P-016	Kanazawa University, Kakuma-machi, Kanazawa 920-1192, Japan Structural Basis of How MGME1 Processes DNA 5' Ends to Maintain Mitochondrial Genome Integrity
	<u>Eric Yin-Chen Mao</u> , Chyuan-Chuan Wu Department of Chemistry, College of Science, National Cheng Kung University, Tainan, Taiwan
*27P-017	Unveiling dynamics of Adenosine A2a receptor coupled to G proteins Sari Hagimoto, Duy Tran, Akio Kitao Tokyo Institute of Technology
*27P-018	Structural basis for recruitment of peptidoglycan endopeptidase MepS by lipoprotein Nlpl Shen Wang Institute of Biochemistry and Molecular Biology, College of Medicine, National Taiwan

University, Taipei, Taiwan.

#### \*27P-019 Structural and functional analysis of PPL, a lectin from the poisonous mushroom Pleurocybella porrigen Daisuke Adachi Graduate School of Medical Life Science, Yokohama City University, 1-7-29 Suehiro, Tsurumi-ku Yokohama, 230-0045, Kanagawa, Japan \*27P-020 ERK1 is a noble topological factor to relax DNA supercoiling Sangmin Ju, Jaehyeon Jeong, Soo Jin Lee, Sanzhar Tarassov, Jeong Ho Jang, Heeyoun Bunch 1School of Applied Biosciences, College of Agriculture & Life Sciences, Kyungpook National University, Daegu, Republic of Korea \*27P-021 Reconstruction and Analysis of the Ancestral ATPase Aya Suzuki, Ryutaro Furukawa, Meghna Sobti, Hiroshi Ueno, Alastair G. Stewart, Satoshi Akanuma, Hiroyuki Noji Applied Chemistry, Graduate School of Engineering, The University of Tokyo, Tokyo, Japan \*27P-022 In situ structural analysis of Salmonella T3SS within the SCV Taiga Horii, Hiroko Takazaki, Yukihisa Hayashida, Yusuke V. Morimoto, Takayuki Kato Grad. Sch. Frontier Biosci., Osaka Univ., Japan/IPR, Osaka Univ., Japan \*27P-023 Structural basis of bifunctionality of mimosine synthase in plants Sayaka Tsuji, Shigeki Ogai, Masakazu Fukuta, Hirosuke Oku, Hiroshi Sugimoto, Masaki Horitani The United Graduate School of Agricultural Sciences, Kagoshima University \*27P-024 Revealing KcsA dynamics by single-particle analysis and molecular dynamics Kotaku Yano, Hiroko Takazaki, Takuo Yasunaga Graduate School of Computer Science and Systems Engineering, Kyushu Institute of Technology, Fukuoka, Japan \*27P-025 Molecular Mechanisms of Diverse Chemokine Recognition and **Downstream Signaling Selectivity of Chemokine Receptors** Fumiya K. Sano, Shirsha Saha, Sharma Saloni, Ramanuj Banerjee, Yoshiaki Kise, Wataru Shihoya, Osamu Nureki, Arun Shukla Grad. Sch. of Sci., The Univ. of Tokyo

*27 <b>P-027</b>	Deciphering Substrate Selectivity in SWEET Transporters: A Molecular Dynamics Perspective Aditi Laddha, Ramasubbu Sankararamakrishnan Department of Biological Sciences and Bioengineering, Indian Institute of Technology Kanpur, Kanpur, Uttar Pradesh, India
*27 <b>P-028</b>	Elucidation of Characteristic Cold-Adaptation Mechanism of Pyruvate Kinase from Psychrophilic Bacteria by X-ray Crystallography <u>Hansani Ekanayake</u> , Hiroshi Sugimoto, Masaki Horitani The United Graduate School of Agricultural Sciences, Kagoshima University, Japan
27P-029	Different Dimerization Behavior of Fluorescent Proteins, eGFP and eYFP <u>Yuna Kinoshita</u> , Haruko Hosoi Toho University
27P-030	Role of actin-binding loops in determining myosin velocity <u>Hideki Furusawa</u> , Takeshi Haraguchi, Kohji Ito Department of Biology, Graduate School of Science, Chiba University, Chiba 263- 8522, Japan
27P-031	Investigated the Amino Acid Region That Enables the fastest Movement in the Fastest Myosin Runa Komoto, Suzune Kato, Kohei Yosimura, Takeshi Haraguchi, Kohji Ito Department of Biology, Graduate School of Science, Chiba University, Chiba 263- 8522, Japan
27P-032	Real-Time, Site-Specific Observation of Chaperone-Mediated Protein Folding using Noncanonical Amino Acid Labeling <u>Munehiro Kumashiro</u> , Adarshi Welegedara, Haocheng Qianzhu, Elwy Abdelkader, Thomas Huber, Gottfried Otting, Tomohide Saio Institute of Advanced Medical Sciences, Tokushima University, Tokushima, Japan
27P-033	Search for specific regions of myosin responsible for moving actin through chiral curves <u>Yoshiki Takayama</u> , Kohei Yoshimura, Taisei Nagai, Takuma Imi, Takeshi Haraguchi, Kohji Ito Department of Biology, Graduate School of Science, Chiba University, Chiba 263- 8522, Japan

#### 27P-034 Reaction Pathways in DNA Hydrolysis of EcoRV Calculated by QM/ MM Metadynamics Itaru Onishi, Mika Mitsumatsu, Hiroki Sato, Ryoutarou Matsuda, Norio Yoshida, Fumio Hirata, <u>Masayuki Irisa</u> Comp. Sci. and Sys. Eng., Kyushu Inst. Tech., Japan 27P-035 Elucidating the Mechanism Underlying Atypical UBA7-UBE2L6 Disulfide Complex Formation Pei-Tzu Chen, Kuen-Phon Wu Institute of Biological Chemistry, Academia Sinica, Taipei, Taiwan 27P-036 Molecular mechanisms for smooth rotation of the flagellar rod within the LP ring

<u>Akio Kitao</u>, Tomoko Yamaguchi, Fumiaki Makino, Tomoko Miyata, Tohru Minamino, Takayuki Kato, Keiichi Namba School of Life Science and Technology, Tokyo Institute of Technology

#### **Protein: Physical property**

*27P-037	Human antimicrobial peptide LL-37 possesses unique multimerization
	properties compared to its orthologs in mouse and rat
	Mitsuki Shibagaki, Jeremia Chrisnanto, Dessalegn Tefera, Kotaro Tsukioka,
	Waka Ueda, Kohei Kano, Hao Gu, Fumi Hirai, Yasuhiro Kumaki,
	Hiroyuki Kumeta, Tomoyasu Aizawa
	Graduate School of Life Science, Hokkaido University, Sapporo, Hokkaido, Japan
1270 020	A Nenetach methodology of Liquid liquid phase concreted draplet

\*27P-038 A Nanotech methodology of Liquid-liquid phase separated droplet regulation with Butterfly-shaped Gold Nanomaterials <u>Tomohiro Nobeyama</u>, Koji Takata, Megumi Mori, Yoichi Yamada, Tatsuya Murakami, Kentaro Shiraki Faculty of Pure and Applied Sciences, University of Tsukuba, Tsukuba, Japan

#### \*27P-039 Nonionic Amino acid Interactions Evaluated Through Solubility Akira Nomoto, Shunsuke Tomita, Kentaro Shiraki Institute of Pure and Applied Sciences, University of Tsukuba/Health and Medical Research Institute, National Institute of Advanced Industrial Science and Technology

#### \*27P-040 Fly-Casting-Like Capture and Translocation of KIF1A by C-Terminal Tail of Tubulin Koki Adachi, Mitsunori Takano Dept. Pure & Appl. Phys., Grad. Scl. Adv. Sci. & Eng., Waseda Univ., Tokyo, Japan

\*27P-041 αB-crystallin prevents aging of α-synuclein droplets Kenji Fujitsuka, Keisuke Yuzu, Yuki Michiue, John A. Carver, Eri Chatani Graduate School of Science, Kobe University, Kobe, Japan

#### 27P-042 Differences in microstructural changes during tensile deformation between hair shapes <u>Hironori Kimura</u>, Kota Yamamoto, Kazuyuki Suzuta Milbon Co., Ltd

#### **Protein: Function**

*27P-043	Development of Cell-free Screening Method for Terminal deoxynucleotidyl transferase for Enzymatic DNA synthesis <u>Takashi Ohmizu</u> , Hiroshi Ueno, Hiroyuki Noji University of Tokyo
*27P-044	Analysis of the physiological significance of dual-localization of Hfd1 in yeast <u>Yuta Konishi</u> , Haruka Sakaue, Hironori Takeda, Toshiya Endo Kyoto Sangyo Univ., Division of Life Science
*27P-045	Identification of multiple responsible genes for abnormal cold acclimation of C. elegans lectin mutants <u>Moe Tezuka</u> , Misaki Okahata, Akane Ohta, Atsushi Kuhara Faculty of Science and Engineering Konan University & Institute for Integrative Neurobiology, Kobe, Japan
27P-046	Reconstitution of ER glutathione transport system <u>Ryuta Sakamoto</u> , Chika Tsutsumi, Ryosuke Tahara, Kazuhiro Nagata, Ryo Ushioda Laboratory of Molecular and Cellular Biology, Faculty of Life Sciences, Kyoto Sangyo University
27P-047	Investigating the catalytic mechanism of Sars-CoV-2 MPro Stephan Kleine-Doepke, Pedram Mehrabi, Caitlin Hatton Universität Hamburg, Germany

#### Protein: Measurement & Analysis

*27P-048	Supramolecular chirality in DFNKF amyloid fibrils derived from human calcitonin by VCD
	<u>Shinryu Isa,</u> Toki Fujino, Raja Prema, Daisuke Sato, Akira Naito, Hisako Sato, Izuru Kawamura
	Yokohama National University, Yokohama, Japan
*27P-049	Measurement of structural flexibility of enzymes using spin labeling- ESR
	<u>Akane Yato,</u> Rio Asaka, Keiichi Watanabe, Masaki Horitani The United Graduate School of Agricultural Sciences, Kagoshima University
27P-050	Muon in Structural Biology: Visualization of proton and electron transfer by the elementary particle "Muon"
	<u>Tamiko Kiyotani</u> , Ichiro Tanaka, Masatoshi Hiraishi, Nobuo Niimura Showa Pharmaceutical University
27P-051	Optimization of Cryo 3D-CLEM for in situ Structural Analysis <u>Hiroko Takazaki</u> , Misaki Arie, Taiga Horii, Takayuki Kato Institute for Protein Research, Osaka University, Osaka, Japan.
27P-052	Investigating CRMP2 isoforms multimerization dynamics by High- Speed AFM
	<u>Djamel Eddine Chafai</u> , Saho Kitagawa WPI Nano Life Science Institute (WPI-NanoLSI), Kanazawa University, Kakuma-machi, Kanazawa 920-1192, Japan
27P-053	Designing an alternative protocol to detect the antigen-antibody reaction using EPR and aggregated AuNPs as paramagnetic probes
	Luis Celedón Ornelas, Alma Nelly Díaz Herreros,
	José Silvestre Figueroa Mendoza, Marco Alonso Arellano Alcántara,
	Belén Chávez Ramírez, Stephany Natasha Arellano Ahumada,
	Daniel Ramírez Rosales
	Instituto Politécnico Nacional

#### Protein: Design & Engineering

*27P-054	De novo design of a protein containing one left-handed βαβ-motif. <u>Naoki Tomita</u> , Hiroto Murata, Hiroki Onoda, Leonard Chavas, George Chikenji Dept. of Appl. Phys., Grad. Sch. of Eng., Nagoya Univ., Aichi, Japan
*27P-055	Machine-learning-assisted multiple maturation of antibody fragment: simultaneous improvement of target-binding, bacterial expression, and thermal stability <u>Tomoyuki Ito</u> , Sakiya Kawada, Hikaru Nakazawa, Akikazu Murakami, Mitsuo Umetsu Grad. Sch. Eng., Tohoku Univ., Sendai, Japan
*27P-056	Construction of heptameric de novo peptide nanopore by chimera proteinization Ayaka Nakada, Kota Natio, Rina Ogawa, Misa Yamaji, Yoshikazu Tanaka, Ryuji Kawano Department of Biotechnology and Life Science, Tokyo University of Agriculture and Technology, Tokyo, Japan
*27P-057	Library design aiming for the development of covalent binding antibody mimetics <u>Yuki Tokunaga</u> , Ryo Matsunaga, Kohei Tsumoto School of Engineering, The University of Tokyo, Japan
*27P-058	Miniaturized cyclic peptides derived from CDR-H3 of antibodies exhibit binding activities to SARS-CoV-2 RBD Yoshiki Yasuda, Satoru Nagatoishi, Ryo Matsunaga, Daisuke Kuroda, Kouhei Tsumoto Department of Chemistry and Biotechnology, school of Engineering, The University of Tokyo, Tokyo, Japan
*27P-059	<b>De novo nanobody binder design by generative AI models</b> <u>Hakyung Lee</u> , Juyong Lee Department of Molecular Medicine and Biopharmaceutical Sciences, Seoul National University
27P-061	Towards further enhancement of the activity of the minimal luciferase picALuc Tadaomi Furuta, Yuki Ohmuro-Matsuyama School of Life Science and Technology, Tokyo Institute of Technology

June 27 [Thu]

#### 27P-062 Structural analysis unveils the enhanced stability of Al-designed ubiquitin-fold proteins Kuen-Phon Wu, Wei-Lin Lu, Wei-Jen Chuang Institute of Biological Chemistry, Academia Sinica, Taipei, Taiwan

#### Protein: Intrinsic disorder

*27P-063	Comprehensive Analysis of Intrinsically Disordered Proteins in the Marsupial
	<u>Shiho Aoki</u> , Wataru Onodera, Toru Asahi
	Waseda University, Dept. of Advanced Sci. and Eng., Tokyo, Japan
*27P-064	Elucidating fusion dynamics of FUS protein droplets using
	fluorescence microscopy and optical tweezers
	<u>Syamil Muharror Ahsanul Husna,</u> Atsumi Hando, Saori Kanbayashi,
	Satoshi Takahashi, Kiyoto Kamagata
	Department of Chemistry, Graduate School of Science, Tohoku University/Institute of Multidisciplinary Research for Advanced Materials, Tohoku University
*27P-065	Balancing stability, dynamics and kinetics in phase separation of
	intrinsically disordered proteins
	<u>Guoqing Zhang</u> , Xiakun Chu
	The Hong Kong University of Science and Technology (Guangzhou)
27P-066	Characterization, regulation, and design of protein droplets
	<u>Kiyoto Kamagata</u> , Ryo Kusano, Atsumi Hando, Nanako Iwaki,
	Maulana Ariefai, Keisuke Ikeda, Tomoshi Kameda
	Institute of Multidisciplinary Research for Advanced Materials, Tohoku University,
	Sendai, Japan

#### Heme proteins

#### \*27P-067 Two distinct conformations in apo forms of bacterial heme ABC transporter <u>Machika Kataoka</u>, Ayaho Abe, Chai Gopalasingam, Christoph Gerle, Hideki Shigematsu, Masaki Yamamoto, Hiroshi Sugimoto

Graduate School of Science, University of Hyogo, Japan./RIKEN SPring-8 Center, Hyogo, Japan.

#### 27P-068 Dramatic Effects of Chemical Modifications on the Function of a Classical Allosteric Protein by Pin-Point Changes in Hydrophobicity Antonio Tsuneshige Department of Frontier Bioscience, and Research Center for Micro-Nano Technology, Hosei University, and Tokyo, Japan

#### Membrane proteins

*27P-069	Vibrational spectroscopic study of chemical interaction between <i>K</i> -opioid receptor (KOR) and ligands having morphinan structure <u>Ryo Nishikawa</u> , Kota Katayama, Seiya lwata, Ryoji Suno, Chiyo Suno, Takuya Kobayashi, Hideki Kandori Graduate School of Engineering. Nagoya Institute of Technology, Aichi, Japan
*27P-070	Role of ANT1 in proton transport: New insights into the mechanism of fatty acid anion sliding at the protein-lipid interface Sanja Vojvodić, Juergen Kreiter, Mario Vazdar, Elena E. Pohl Physiology and Biophysics, University of Veterinary Medicine, Vienna, Austria
*27P-071	Cryo-EM Structural Analysis of Enterococcus hirae V-ATPase with Improved Resolution Yuan-E Lee, Raymond Burton-Smith, Akihiro Otomo, Takeshi Murata, Ryota lino, Kazuyoshi Murata ExCELLS/NIPS, Okazaki, Japan
*27P-072	Solid-state NMR analysis of wild-type and mutant Schizorhodopsin proteins <u>Akito Kitaguchi</u> , Seiya Tajima, Toshio Nagashima, Toshio Yamazaki, Hideki Kandori, Keiichi Inoue, Izuru Kawamura Yokohama National University, Japan
*27P-073	Zn2+ Ion Transportation Mechanisms of TRPC6 Channels: All-Atom Molecular Dynamics Simulation Sirin Sittiwanichai, Kowit Hengphasatporn, Yasuteru Shigeta Center for Computational Sciences, University of Tsukuba, 1-1-1 Tennodai, Tsukuba,

Ibaraki, 305-8577, Japan

## \*27P-074 Structural Basis for Signaling and Drug-Induced Activation of the Trk Receptors

<u>Erik Kot</u>, Sergey Goncharuk, Ekaterina Vasilieva, Alexandra Shabalkina, María Franco, Ekaterina Lyukmanova, Alexander Arseniev, Andrea Benito-Martínez, Mario Costa, Antonino Cattaneo, Marcal Vilar, Konstantin Mineev Shenzhen MSU-BIT University, Shenzhen, China

#### 27P-075 New Lipid-Bilayer Nanodiscs for Membrane-Protein Biophysics <u>Sandro Keller</u>, David Glueck, Lena Bauernhofer, Loretta Eggenreich, Carolyn Vargas Biophysics, Institute of Molecular Biosciences (IMB), University of Graz, Austria

27P-076 Coupling of ATP reactions with allocrite transport in heme ABC transporter; BhuUV-T, revealed by time-resolved spectroscopy. <u>Tetsunari Kimura</u>, Ayaka Naka, Akiho Hara, Yasuhiro Kobori, Yoshitsugu Shiro, Hiroshi Sugimoto Kobe University, Graduate School of Science, Department of Chemistry/Kobe

University, Molecular Photoscience Research Center

#### 27P-077 Understanding the Structure and Receptor Selectivity of Histamine H4 Receptor

Dohyun Im, Jun-ichi Kishikawa, Yuki Shiimura, Yukihiko Sugita, Takeshi Noda, Takayuki Kato, Hidetsugu Asada, So Iwata Department of Cell Biology, Graduate School of Medicine, Kyoto University, Kyoto, Japan

#### 27P-078 Generation of human TMEM16F-specific affibodies

Eunyoung Kim, Jinho Bang, Ji Hye Sung, Jonghwan Lee, Sunghyun Kim, Byoung-Cheol Lee

Korea Brain Research Institute, Neurovascular Unit Research Group, Daegu 41068, Korea

#### **DNA & DNA binding proteins**

*27P-079	High-speed AFM analysis of effects of drugs on the dynamic DNA morphologies interacting with MDP1, dormancy induction protein of
	Mycobacterium tuberculosis
	Kaho Nakamoto, Rei Moriya, Kenichi Umeda, Akihito Nishiyama,
	Sohkichi Matsumoto, Noriyuki Kodera
	Grad. Sch. Math. & Phys., Kanazawa Univ.

- \*27P-080 Single-molecule imaging of MDP1, dormancy induction protein of Mycobacterium tuberculosis, with high-speed AFM Yuna Goto, Kaho Nakamoto, Kenichi Umeda, Akihito Nishiyama, Sohkichi Matsumoto, Noriyuki Kodera Grad. Sch. Math. & Phys., Kanazawa Univ.
- \*27P-081 Template-free oligonucleotide synthesis by Terminal Deoxynucleotidyl Transferase in a microreactor array Yusuke Miyata, Hiroshi Ueno, Hiroyuki Noji Department of Applied Chemistry. School of Engineering. The University of Tokyo
- 27P-082 Phosphorylation and histone peptides reduce main- but not sidechain dynamics of N-terminal intrinsically disordered region of HP1 during phase separation, as studied by conventional and TOAC spin labels

Isao Suetake, Toshiki Takei, Tomoaki Sugishiata, Shun Ito, Kazunobu Sato, Yuichi Mishima, Kohei Muraoka, Toru Kawakami, Yoh Matsuki, Toshimichi Fujiwara, Takeji Takui, Makoto Miyata, Hironobu Hojo, <u>Toshiaki Arata</u> Inst. Protein Res., Osaka Univ., Japan/Grad. Sch. Sci., Osaka Metropolitan Univ.,

**RNA & RNA binding proteins** 

Japan

27P-083 Short repeat RNA suppresses aggregation of ALS-causative protein TDP-43 and its 25 kDa carboxy-terminal fragment <u>Ai Fujimoto</u>, Akira Kitamura Graduate school of Life Science, Hokkaido University, Hokkaido, Japan

#### **DNA/RNA** nanotechnology

*27P-084	How to engineer a fast-moving DNA-nanoparticle motor with long run length and high unidirectionality? <u>Takanori Harashima</u> , Akihiro Otomo, Ryota lino Institute for Molecular Science, National Institutes of Natural Sciences/Graduate Institute for Advanced Studies, SOKENDAI
*27P-085	Timing-controlled dynamics of DNA droplet-based artificial cell Tomoya Maruyama, Masahiro Takinoue Department of Life Science and Technology, Tokyo Institute of Technology, Japan
*27P-086	Regulation of molecular distribution in lipid vesicles based on artificial DNA cortex <u>Takuro Yoshinaga</u> , Koki Shibata, Yusuke Sato Department of Intelligent Systems Engineering, Kyushu Institute of Technology, Japan
*27P-087	Mechanical properties of artificial cells with DNA cytoskeleton Kazutoshi Masuda, Miho Yanagisawa Graduate School of Arts & Science, The University of Tokyo, Tokyo, Japan
*27P-088	Lipid nanoparticle fusion with a phospholipid membrane Jan Šimek, Nestor Mora, Radek Šachl J. Heyrovsky Institute of Physical Chemistry
*27 <b>P-</b> 089	Spontaneous film-like DNA structure formation at the oil-air interface Daichi Tominaga, Shogo Hamada, Yusuke Sato Department of Intelligent Systems Engineering, Kyushu Institute of Technology, Japan

#### **Nucleic acid: Others**

- \*27P-090 The effect of Temperature and Pressure on the structural transition from the quadruplex to random coil of VEGF Hiroto Yamasaki, Toshiki Nakao, Minoru Kato Ritsumeikan University
- 27P-091 Modelling complex and large RNA structures to advance RNA biology and therapeutics <u>Naoto Hori</u>, James A. Robins, Huong T. Vu School of Pharmacy, University of Nottingham

#### **Chromatin & Chromosomes**

*27 <b>P-09</b> 2	DNA Unwinding analysis of N-terminal tailless nucleosomes using
	nanopore measurements

Satoshi Ogihara, Hikaru Nozawa, Takumi Oishi, Munetaka Akatsu, Hitoshi Kurumizaka, Sotaro Uemura Department of Biological Sciences, Graduate School of Science, The University of Tokyo

#### 27P-093 Effect of RNA expression on chromatin phase separation : Molecular Dynamics simulation

Shaya Shiraishi, Yuuki Norizoe, Takuya Saito, Takahiro Sakaue Department of Physical Sciences, Aoyama Gakuin University

#### Electronic

\*27P-095 Electrochemical activity of catalytic amyloids: self-assembly of (XH)4 peptides and hemin on graphite electrodes <u>Marie Sugiyama</u>, Luo Wei, Ayhan Yurtsever, Takeshi Fukuma, Yuhei Hayamizu Tokyo Institute of Technology, Tokyo, Japan

#### Water & Hydration & Electrolyte

- \*27P-096 Prediction of hydration structures over membrane proteins using deep learning in combination with the emprical hydration distribution Kochi Sato, Mao Oide, Masayoshi Nakasako Department of Physics, Keio University, Kanagawa, Japan/SPring-8 Center, RIKEN, Hyogo, Japan/SPRING, Japan Science and Technology Agency, Tokyo, Japan
- 27P-097 Hydration and Fluctuation Dynamics of a Membrane Transport Protein-Glucose Complex

Tatsuki Kawauchi, Tomohiko Hayashi, Mitsunori Ikeguchi Graduate School of Science and Technology, Niigata University

#### Molecular genetics & Gene expression

27P-098 Nine-banded armadillo transcriptome and chromatin accessibility at single-cell reveal persistent identity signatures in concordance with cell population variations Risa Karakida Kawaguchi, Sara Ballouz, Maria T Pena, Leon French, Frank M. Knight, Linda B. Adams, Jesse Gillis Center for iPS Cell Research and Application, Kyoto University/Cold Spring Harbor Laboratory

#### **Morphogenesis and Development**

\*27P-099 Deep learning approach to investigate tissue hydraulics during ovarian follicle development.

Jake Turley, Kim Whye Leong, Chii Jou Chan Mechanobiology Institute, National University of Singapore

#### Muscle

27P-100 Morphological discrimination of isolated sarcoplasmic reticulum vesicles in different Ca2+ concentrations using deep learning <u>Katsuya Saito</u>, Kenji Etchuya, Jun Nakamura, Chikara Sato, Makiko Suwa Biological Science Course, Graduate School of Science and Engineering, Aoyama Gakuin University, Kanagawa, Japan

#### Molecular motor

\*27P-101 Microscopic Choreography: Unraveling Molecular Properties of Cytoplasmic Dynein Shaping Collective Motion of Microtubules in vitro

#### Yosuke Harada, Kazuhiro Oiwa

Department of Life Science, Graduate School of Science, University of Hyogo, Hyogo, Japan/Advanced ICT Research Institute, National Institute of Information and Communications Technology, Hyogo, Japan

#### \*27P-102 Biochemical Characterization of C. elegans Kinesin Bmk-1

<u>Toru Kurosaka</u>, Shunsuke Kumagai, Fofou Yonta Tostani, Shinsaku Maruta Department of Biosciences, Graduate School of Science and Engineering Soka University, Hachioji, Tokyo JAPAN

*27P-103	Rejuvenating actin filaments: Direct observation of nucleotide exchange in actin filaments enhanced by myosin II Kenta Toshino, Taro QP Uyeda Dept. Pure & Appl. Physics, Grad. Sch. Adv. Sci. & Eng., Waseda Univ.
*27P-104	Molecular dynamics observation of rotational motion in the stator unit of the flagellar motor <u>Takumi Matsumoto</u> , Yukinari Kamiyama, Mitsunori Takano Dept. of Pure & Appl. Phys., Grad. Sch. of Adv. Sci. and Eng., Waseda Univ., Tokyo, Japan
*27 <b>P</b> -105	<b>Negative differential resistance of bio-molecular motor F1-ATPase</b> <u>Haruto Kimura</u> , Shoichi Toyabe, Yohei Nakayama Department of Applied Physics, Graduate School of Engineering, Tohoku University
27P-106	V-ATPase rotation probed by Janus nanoparticle Akihiro Otomo, Jared Wiemann, Swagata Bhattacharyya, Yan Yu, <u>Ryota lino</u> Institute for Molecular Science, National Institutes of Natural Sciences, Okazaki, Japan/Graduate Institute for Advanced Studies, SOKENDAI, Hayama, Japan
27P-107	Detecting conformations of F1-ATPase to elucidate the rotation mechanism Kenta Suga, Fumika Ogura, Hiroki Yamashita, Hiroki Kaizu, Ayari Tagawa, Mitsuhiro Sugawa, Nobukiyo Tanaka, Tomoko Masaike Department of Applied Biological Science, Faculty of Science and Technology, Tokyo University of Science, Japan
27P-108	Cooperation among c-subunits of FoF1-ATP synthase in rotation- coupled proton translocation by hetero-mutated c-ring <u>Noriyo Mitome</u> , Shintaroh Kubo, Sumie Ohta, Hikaru Takashima, Yuto Shigefuji, Toru Niina, Shoji Takada Faculty of Education, Tokoha University
27P-109	DNA Hybridization kinetics in Active Matter self-assembly Mst Rubaya Rahsid, Yamashina Takefumi, Kawamata Ibuki, Marie Tani, Masatoshi Ichikawa, Akira Kakugo Department of Physics, Kyoto University, Japan

#### **Single Molecule Biophysics**

#### \*27P-110 Microsecond single molecule dynamics measurement of SARS-CoV-2 Spike protein using Diffracted X-ray Tracking

<u>Daisuke Sasaki</u>, Tatsuya Arai, Hiroshi Sekiguchi, Kazuhiro Mio, Yuji Sasaki Graduate School of Frontier Sciences, The University of Tokyo

## \*27P-111 REGULATORY MECHANISMS OF KINESIN FUNCTION AT VARYING PH

Suvranta Tripathy, <u>Fawaz Baig</u>, Hassan Bazzi University of Michigan Dearborn

#### \*27P-112 Impact of mutations on cadherin 23 functions and leads to hearingloss disease

Gaurav Kumar Bhati, Surbhi Garg, Pritam Saha, Sabyasachi Rakshit Department of Chemical sciences, Indian Institute of Science Education and Research Mohali, India

## **\*27P-113** Direct observation of a single DNA molecule responding for the AC electric field and different physical environment.

Yunosuke Fuji, Shin Tkano, Takuma Yoshinaga, Yuuta Moriyama, Toshiyuki Mitsui Grad. Sch. of Sci. and Eng. Aoyama Gakuin Univ.

## \*27P-114 Photothermal assisted ultra-low concentration detection using nanopore sensing

<u>Hirohito Yamazaki</u>, Kota Kaito Top Runner Incubation Center for Academia-Industry Fusion, Nagaoka University of Technology, Nagaoka/Department of Mechanical Engineering, Nagaoka University of Technology

## 27P-115 Single-molecule analysis of the behavioral dynamics of EGFR cancer mutants with resistance to anticancer drugs

Michio Hiroshima, Masahiro Ueda Osaka University/RIKEN BDR

## 27P-116 Mechanistic insight into the mechanical unfolding of integral membrane proteins

Hao Yu Huazhong University of Science and Technology

#### **Cell biology: Adhesion**

27P-117 Study of adhesion factor in Acanthamoeba bunch formation caused by Hokutovirus infection Yuto Shimada, Masaharu Takemura, Kazuyoshi Murata

Exploratory Research Center on Life and Living Systems, National Institutes of Natural Sciences

#### **Cell biology: Motility**

#### \*27P-118 The mechanical properties of fibroblasts in co-culture system Arata Nagai, Kaito Kojima, Hiromu Kuwabara, Yuuta Moriyama, Toshiyuki Mitsui Grad. Sch. of Sci. and Eng. Aoyama Gakuin Univ.

**\*27P-119** Mechanism of bacterial actin driven motility reconstituted in a minimal synthetic bacterium

Hana Kiyama, Shigeyuki Kakizawa, Daichi Takahashi, Makoto Miyata Graduate School of Science, Osaka Metropolitan University, Osaka, Japan

\*27P-120 Light-Induced Control of Archaellum Rotation in Haloacterium salinarum

Ishii Kazuki, Ayaka Ihara, Daisuke Nakane, Takayuki Nishizaka Gakushuin University

\*27P-121 Haloplasma motility reconstituted in a minimal synthetic bacterium, JCVI-syn3B

Mone Mimura, Hana Kiyama, Shingo Kato, Yuya Sasajima, Atsuko Uenoyama, Shigeyuki Kakizawa, André Antunes, Tomoko Miyata, Fumiaki Makino, Keiichi Namba, Makoto Miyata Grad. Sch. Sci., Osaka Metropolitan Univ., Japan

\*27P-122 Gliding machinery of Mycoplasma mobile observed by electron cryotomography <u>Minoru Fukushima</u>, Tomoko Miyata, Takuma Toyonaga, Keiichi Namba,

Makoto Miyata Grad. Sch. Sci., Osaka Metropolitan Univ., Osaka, Japan

\*27P-123 Rapid response of bacterial motility with pressure change Seiichiro Kinoshita, Masayoshi Nishiyama Grad.Sch.Sci. and Eng., Kindai Univ.

*27P-124	Visualization and analysis of MreBs driving Spiroplasma motility in minimal synthetic bacterium Yoshiki Tanaka, Hana Kiyama, Takuma Toyonaga, Makoto Miyata Grad. Sch. Sci., Osaka Metro Univ.
*27P-125	In vitro analysis of the bacterial actin MreB molecule that gives swimming motility to the minimal synthetic bacterium JCVI-syn3B. Satoshi Kanamori, Daichi Takahashi, Yuhei Tahara, Hana Kiyama, Makoto Miyata Graduate School of Science, Osaka Metropolitan University
27P-126	<b>CryoEM structures of the growing end of the bacterial flagellar hook.</b> <u>Haruto Takeuchi</u> , Sae Hashimoto, Tomoko Miyata, Fumiaki Makino, Keiichi Namba, Norihiro Takekawa, Katsumi Imada Dept. of Macromol. Sci., Grad. Sch. of Sci., Osaka Univ.
27P-127	Activation of the PomA/B flagellar stator by a site-specific chemical modification in the plug segment Hiroaki Koiwa, Akihiro Otomo, Yuki Tajimi, Tatsuro Nishikino, Michio Homma, Takayuki Uchihashi, Ryota lino, <u>Seiji Kojima</u> Department of Biological Science, Graduate School of Science, Nagoya University
27P-128	Analysis of the Virio alginolyticus lateral flagellar motor genes, lafT and lafU <u>Kazuki Yokoyama</u> , Norihiro Takekawa, Seiji Kojima Department of biological science, Graduate school of Science, Nagoya University
27P-129	Structural change of ATPase ring complex of the flagellar type III export apparatus revealed by cryoEM analysis and high-speed AFM Norihiro Takekawa, Asako Usui, Yuki Tajimi, Miki Kinoshita, Tohru Minamino Takayuki Uchihashi, Katsumi Imada Dept. of Macromol. Sci., Grad. Sch. of Sci., Osaka Univ.

#### Cell biology: Cytoskeleton & Membrane skeleton

\*27P-130 Cryo-ET of vertebrate cilia revealed that Calaxin stabilizes the docking of outer arm dyneins onto ciliary doublet microtubule <u>Hiroshi Yamaguchi</u>, Motohiro Morikawa, Masahide Kikkawa Department of Cell Biology & Anatomy, Grad. Sch. Med., The University of Tokyo, Tokyo, Japan

*27P-131	Capping and severing mechanisms of Cytochalasin D to actin filament by TIRF observation <u>Takahiro Mitani</u> , Shuichi Takeda, Ikuko Fujiwara, Hajime Honda Dep. of Material Sci. and Bioeng., Nagaoka Univ. of Tech., Niigata, Japan.
*27P-132	Physically specific domain at the plasma membrane induced by transmembrane phospholipid movement during myoblast cytokinesis <u>Akira Murakami</u> , Kotaro Hirano, Junya Sano, Kohki Okabe, Yuji Hara School of Pharmaceutical Sciences, University of Shizuoka
*27P-133	Elucidating the Role of Spiroplasma fibril protein using synthetic bacterium, JCVI syn3 <u>Ali Ahsan</u> , Hana Kiyama, Makoto Miyata Osaka Metropolitan University, Graduate School of Science
27P-134	Domain characterization of Archaea gelsolin for inhibiting actin polymerization by TIRF and crystal structure observations <u>Horyo Mizuki</u> , Shuichi Takeda, Robert Robinson, Ikuko Fujiwara Materials Sciences and Bioengineering, Nagaoka University of Technology
27P-135	Thermodynamic Analysis of Cofilin–F-actin Interaction <u>Hideyuki Komatsu</u> , Nayu Itou, Sinobu Sato, Shigeori Takenaka Department of Bioscience and Bioinformatics, Kyushu Institute of Technology
27P-136	The phase separation of EB and TEN2 promotes inhibitory synapse formation Sotaro Ichinose, Hirohide Iwasaki Department of Anatomy, Gunma University Graduate School of Medicine, Gunma, Japan
Cell biology:	Signal transduction & Cell membrane
*27P-137	Aquaporin-3 and aquaporin-5 differentially modulate cell stiffness and

cell-cell adhesion and promote cell migration Catarina Pimpão, Filomena A. Carvalho, Inês V. da Silva, Nuno C. Santos, Graça Soveral

> Research Institute for Medicines (iMed.ULisboa), Faculty of Pharmacy, Universidade de Lisboa, 1649-003 Lisbon, Portugal/Department of Pharmaceutical Sciences and Medicines, Faculty of Pharmacy, Universidade de Lisboa, 1649-003 Lisbon, Portugal

Poster Sessions

June 27 [Thu]

- \*27P-138 Aquaporin-3 is involved in inflammasome activation contributing to the settings of inflammatory response in THP-1 cells Inês V. da Silva, Angela Casini, Pablo Pelegrin, Graça Soveral Research Institute for Medicines (iMed.ULisboa), Faculty of Pharmacy, Universidade de Lisboa, 1649-003 Lisboa, Portugal
- \*27P-139 Positive feedback regulation of excitable Ras by RasGEFX for spontaneous signal generation in cell migration Koji Iwamoto, Satomi Matsuoka, Masahiro Ueda Grad. Sch. Sci., Osaka Univ, Osaka, Japan
- 27P-140 Phosphatidylserine enhances membrane localization and lateral diffusion of active form of Ras for excitability Satomi Matsuoka, Da Young Shin, Michio Hiroshima, Hiroaki Takagi, Masahiro Ueda Graduate School of Frontier Biosciences, Osaka University/Graduate School of Science, Osaka University/Center for Biosystems Dynamics Research, RIKEN

#### Biological & Artificial membrane: Structure & Property

- \*27P-141 Nanofluidic model cell membrane platform for molecular analysis of membrane-bound proteins Yu Yoshimura, Nanami Nagatsuka, Ryota Komatsu, Shin-ichi Yusa, Kenichi Morigaki Graduate School of Agricultural Science, Kobe Univ, Hyogo, Japan
   \*27P-142 Integrated model membrane arrays generated by self-spreading of lipid bilayers Masako Fujii, Kenichi Morigaki Grad. of Agri. Sci., Kobe Univ., Hyogo, Japan.
   \*27P-143 Field model for multistate lateral diffusion of various transmembrane
- \*27P-143 Field model for multistate lateral diffusion of various transmembrane proteins observed in living Dictyostelium cells Kazutoshi Takebayashi, Yoichiro Kamimura, Masahiro Ueda Center for Biosystems Dynamics Research (BDR), RIKEN
- \*27P-144 Membrane shapes, liquid-liquid interfaces, and elastocapillarity Lukas Hauer, Katharina Sporbeck, Amir H. Bahrami, Roland L. Knorr Humboldt-Universitaet zu Berlin, Berlin, Germany

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## **Poster Sessions**

27P-145	Study on the mechanism of double cooperative effect of antimicrobial peptide LL-37 with HNP1 by leakage assay
	<u>Yuta Sekiya,</u> Kaori Sugihara
	The University of Tokyo, Institute of Industrial Science
27P-146	Induction of Apoptosis by Ceramide Derivatives and Its Potential Mechanisms through Domain Formation
	<u>Nobuaki Matsumori,</u> Koya Tsujimura, Miho Yakabe, Hideaki Kano
	Kyushu University

27P-147 Structural effects of Cholesterol, Lanosterol, and Oxysterol on Model Biomembranes

Ayumi Okayama, Tatsuya Hoshino, Kohei Wada, <u>Takahashi Hiroshi</u> Biophysics Laboratory, Division of Pure and Applied Science, Gunma University, Maebashi, Japan

#### **Biological & Artificial membrane: Dynamics**

*27P-148	Generation of autonomous rotors
	Veerpal Kaur, Charu Taneja, Subha shree, Abhishek Chaudhuri,
	Sabyasachi Rakshit
	Veerpal Kaur, Department of Chemical Sciences, Indian Institute of Science Education and Research Mohali, Mohali, Punjab, India
*27P-149	Molecular dynamics investigation of the dynamical response of the interfacial waters near DPPC bilayer to Hyaluronic acid
	<u>Anirban Paul</u> , Jaydeb Chakrabarti
	S. N. Bose National Centre for Basic Sciences, Kolkata, India
*27P-150	Optical Trapping of Membrane Proteins on the Supported Lipid Bilayers
	<u>Yasushi Tanimoto</u> , Shunya Moriyama, Kyoko Masui, Chie Hosokawa Graduate School of Science, Osaka Metropolitan University

27P-151 Structurally Stable Phospholipid Membrane Tube Developed by Self-assembly of Peptide Receptors Noriyuki Uchida, Ryu Ishizaka, Anju Kawakita, Masaki Okumura, Takahiro Muraoka Tokyo University of Agriculture and Technology

#### **Biological & Artificial membrane: Transport & Signal transduction**

27P-153 Reconstituting G protein-coupled receptors into a supported lipid bilayer using meta-stable peptide nanodiscs <u>Fumio Hayashi</u>, Masato Koezuka, Kenich Morigaki Grad Sch Sci, Kobe University

#### Membraneless Organella, autophage, Liquid-liquid phase separation

- \*27P-155 TMAO and urea effects on liquid-liquid phase separation of fused in sarcoma Keiji Kitamura, Ayano Ohshima, Fuka Sasaki, Yutaro Shiramasa, Soichiro Kitazawa, Ryo Kitahara Graduate School of Pharmacy, Ritsumeikan University, Shiga, Japan
- \*27P-156 Coarse-Grained Molecular Dynamics Study of Coacervate Formation using Elastin-like Polypeptides with Varying Hydrophobicity <u>Haruto Takegahara</u>, Yasunori Okamoto, Kenichi Funamoto, Takuya Mabuchi Graduate School of Biomedical Engineering, Tohoku University
- \*27P-157 Raman spectroscopic study of liquid-liquid phase separation in Lysozyme/Ovalbumin mixture system Taiga Sano, Toshiki Nakao, Minoru Kato Ritsumeikan University
- 27P-158 Liquid-liquid phase separation of the P53 core domain Amanda Santos Palma, Carlos Henrique Inácio Ramos, Leandro Ramos Souza Barbosa University of São Paulo, São Paulo, Brazil
- 27P-159 Theoretical studies of protein accumulation during mitosis with Flory-Huggins free energy Yuuki Karube, Yuuki Norizoe, Takuya Saito, Takahiro Sakaue Department of Physical Sciences, Aoyama Gakuin University

#### 27P-160 Quantitative Analytical Method Based on Machine Learning by Classification of Condensate Forming Cells by Glycolytic Enzymes in Saccharomyces cerevisiae Natsuko Miura, Ryuta Saito, Yuki Yoshimura, Kohei Tanaka, Michihiko Kataoka Graduate School of Agriculture, Osaka Metropolitan University/Graduate School of Life and Environmental Sciences, Osaka Prefecture University

#### **Neuroscience & Sensory systems**

#### \*27P-161 Controlling Tau Aggregation Using Light-Induced Cellular Models of Tau Oligomers

Tomoya Uchida, Naoki Kato, Shigeo Sakuragi, Akito Hattori, Yoshiyuki Soeda, Hideaki Yoshimura, Akihiko Takashima, Hiroko Bannai Waseda University, School of Advanced Science and Engineering, Tokyo, Japan

#### 27P-162 Specification of Smallest Neural Cell Colony Size for Measurement of Firing or Burst Firing

<u>Takumi Yamaguchi</u>, Kentaro Kito, Masahito Hayashi, Tomoyuki Kaneko LaRC, Dept. Frontier Biosci., Hosei Univ., Tokyo, Japan

#### Neuronal circuit & Information processing

#### 27P-164 Classifying Dynamics of Ising Interaction Networks by Structure of Traffic Diagrams Yoshiaki Horiike, Shin Fujishiro, Masaki Sasai

Department of Applied Physics, Nagoya University, Nagoya, Japan/Department of Neuroscience, University of Copenhagen, Copenhagen, Denmark

#### Behavior

#### \*27P-165 Quantitative Description and Investigation into the Mechanism of Gravitactic Swimming Behavior in Coral Larvae Asuka Takeda-Sakazume, Junko Honio, Kanae Matsushima.

Sachia Sasano, Yuuko Wada, Minori Oshima, Shoji A. Baba, Kei Yura, Yoshihiro Mogami, Masayuki Hatta Graduate School of Humanities and Sciences, Ochanomizu University, Tokyo, Japan/

Graduate School of Humanities and Sciences, Ochanomizu University, Tokyo, Japan/ Faculty of Core Research Natural Sciences Division, Ochanomizu University, Tokyo, Japan

#### Photobiology: Vision & Photoreception

*27P-166	The structural dynamics study of green-cone pigment by using spectroscpies <u>Mizusa Kani</u> , Sayaka Ohashi, Takuma Sasaki, Hiroo Imai, Hideki Kandori, Kota Katayama Grad. Sch. Eng., Nagoya Inst. Tech., Aichi, Japan
*27P-167	FTIR study of mutants of primate red and green pigments Sayaka Ohashi, Hiroo Imai, Hideki Kandori, Kota Katayama Grad. Sch. Eng., Nagoya Inst. Tech., Aichi, Japan
*27P-168	Activation mechanism of light-sensitive Gs protein-coupled receptor, jellyfish rhodopsin Shino Inukai, Mitsumasa Koyanagi, Akihisa Terakita, Hideki Kandori, Kota Katayama Graduate School of Engineering, Nagoya Institute of Technology.
*27P-169	Spectroscopic analysis of the photoreaction of TAT rhodopsin in the presence of calcium ion <u>Teppei Sugimoto</u> , Kota Katayama, Hideki Kandori Graduate School of Engineering, Nagoya Institute of Technology, Japan,
27 <b>P-170</b>	Free energy profile analysis of natural anion channelrhodopsin GtACR1 in each state of the photocycle <u>Takafumi Shikakura</u> , Cheng Cheng, Shigehiko Hayashi Graduate School of Science, Kyoto University, Kyoto, Japan
27P-171	Production of a light-driven Cldependent Na+ pump: Implications for the binding and transport of distinctive ions <u>Manami Hashimoto</u> , Kano Suzuki, Marie Kurihara, Taiki Nakamura, Keiichi Kojima, Susumu Yoshizawa, Yasuhisa Mizutani, Takeshi Murata, Yuki Sudo Grad. Sch., Med. Dent, and Pharm. Sci., Okayama Univ., Okayama, Japan.
	Grad. Sch., Med. Dent, and Pharm. Sci., Okayama Univ., Okayama, Japan.

#### **Photobiology: Photosynthesis**

^Z/P-1/Z	Light factor-dependent Growth of Yellow Chlamydomonas
	Okviyoandra Akhyar, Soichiro Seki, Kazuhiro Yoshida, Chiyo Takagi,
	Yasuhiro Kamei, Ritsuko Fujii
	Research Center for Artificial Photosynthesis (ReCAP), Osaka Metropolitan University, Japan
*27P-173	Robustness of photosynthetic light-harvesting antenna chlorosome against structural heterogeneity
	<u>Shun Arai</u> , Tomomi Inagaki, Jiro Harada, Chihiro Azai, Toru Kondo Tokyo Institute of Technology
*27P-174	Energy Transfer Pathway in Chlorophyll-f Containing Photosystem I Revealed by Single-Molecule Spectroscopy
	<u>Rin Taniguchi</u> , Toshiyuki Shinoda, Tatsuya Tomo, Ye Shen, Yutaka Shibata Department of Chemistry, Tohoku University, Miyagi, Japan,
27P-175	Post-translational conversion of amino acids in the O2-evolving
27P-175	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from alighatic amino acide
27P-175	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids Hatsune Mizue, Takehiro Suzuki, Takumi Matsubara, Tomomi Kitaiima-Ibara.
27P-175	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids <u>Hatsune Mizue</u> , Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae,
27P-175	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids <u>Hatsune Mizue</u> , Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi
27P-175	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids <u>Hatsune Mizue</u> , Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi Department of Physics, Graduate School of Science, Nagoya University
27P-175 27P-176	<ul> <li>Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids</li> <li>Hatsune Mizue, Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi</li> <li>Department of Physics, Graduate School of Science, Nagoya University</li> <li>Modification of chlorophyll pigments in photosynthetic light-</li> </ul>
27P-175 27P-176	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids Hatsune Mizue, Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi Department of Physics, Graduate School of Science, Nagoya University Modification of chlorophyll pigments in photosynthetic light- harvesting proteins
27P-175 27P-176	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acidsHatsune Mizue, Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi Department of Physics, Graduate School of Science, Nagoya UniversityModification of chlorophyll pigments in photosynthetic light- harvesting proteins Yoshitaka Saga, Shota Kawato, Kohei Hamanishi, Moe Sumura
27P-175 27P-176	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids Hatsune Mizue, Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi Department of Physics, Graduate School of Science, Nagoya University Modification of chlorophyll pigments in photosynthetic light- harvesting proteins Yoshitaka Saga, Shota Kawato, Kohei Hamanishi, Moe Sumura Kindai University
27P-175 27P-176	Post-translational conversion of amino acids in the O2-evolving complex of photosystem II: Formation of carboxylate ligands from aliphatic amino acids Hatsune Mizue, Takehiro Suzuki, Takumi Matsubara, Tomomi Kitajima-Ihara, Minako Hirano, Yuichiro Shimada, Yuki Kato, Naoshi Dohmae, Takumi Noguchi Department of Physics, Graduate School of Science, Nagoya University Modification of chlorophyll pigments in photosynthetic light- harvesting proteins Yoshitaka Saga, Shota Kawato, Kohei Hamanishi, Moe Sumura Kindai University

\*27P-177 Effect of photoactivated adenylyl cyclase expression in Salmonella Keisuke Sakai, Yusuke V. Morimoto Graduate School of Computer Science and Systems Engineering, Kyushu Institute of Technology, Fukuoka, Japan

#### \*27P-178 Novel Optogenetic Strategy for Regulating Insulin Signaling in the Deep Tissues of Living Mice Qi Dong, Mizuki Endo, Takeaki Ozawa The University of Tokyo

- 27P-179Relationship between Responsiveness of Cardiomyocytes Stimulated<br/>by Laser Irradiation and Cell Population Status<br/>Takaaki Nishikawa, Furuie Yasumasa, Kentaro Kito, Masahito Hayasi,<br/>Tomoyuki Kaneko<br/>LaRC, Dept. Frontier Biosci., Hosei Univ., Tokyo, Japan
- 27P-180 The "fifth" color switch of microbial rhodopsin Rei Abe-Yoshizumi, Hideki Kandori Grad. Sch. of Eng., Nagoya Inst. of Tech.

#### Radiobiology & Active oxygen

27P-181 Cell-killing caused by direct and indirect actions of high-LET particles in Boron Neutron Capture Therapy (BNCT) Ryoichi Hirayama, Yu Sanada, Akiko Uzawa, Yoshitaka Matsumoto, Atsushi Ito, Shin-ichiro Masunaga, Hiroki Tanaka, Yoshinori Sakurai, Minoru Suzuki, Sumitaka Hasegawa Institute for Quantum Medical Science, National Institutes for Quantum Science and Technology

#### **Origin of life & Evolution**

*27P-182	Droplets in PEG / salt solution as primitive compartments at the origin
	of life
	Yota Tabata, Masahito Hayashi, Tomoyuki Kaneko

LaRC, FB, Grad. Sch. Sci. & Eng., Hosei Univ., Tokyo, Japan

and Technology, Tsukuba, Japan

**\*27P-183** Adaptive Laboratory Evolution of Minimal Genome Bacterium to Low Temperature

Masaki Mizutani, Minoru Moriyama, Ryuichi Koga, Takema Fukatsu, Shigeyuki Kakizawa Bioproduction Research Institute, National Institute of Advanced Industrial Science

\*27P-184 Stability, structure, and interactions of prebiotic fatty acid membranes <u>Taren Elizabeth Buddle Ginter</u>, Akiko Baba, Masayuki Imai, Maikel Rheinstädter, Kosuke Fujishima Earth-Life Science Institute, Tokyo Institute of Technology, Tokyo, Japan/School of Life Science and Technology, Tokyo Institute of Technology, Tokyo, Japan

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#### Synthetic biology & Artificial cells

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^272-185	Artificial Cell Compartment by Thermal Control <u>Mirai Sasaki</u> , Yoshihiro Minagawa, Hiroyuki Noji Department of Applied Chemistry, The University of Tokyo, Tokyo, Japan
*27P-186	Microtubule/kinesin complexes spontaneously emerge vortices in cell-sized droplet generated by water/water phase separation <u>Hiroki Sakuta</u> , Naoki Nakatani, Takayuki Torisawa, Yutaka Sumino, Kanta Tsumoto, Kazuhiro Oiwa, Kenichi Yoshikawa Universal Biology Institute, University of Tokyo/Graduate School of Arts and Sciences, University of Tokyo
*27P-187	Regulation of Stochastic Cell Re-differentiation Ratio of Genetic Toggle Switch with Minute Expression Balancing Control of Repressor Proteins Sota Okuda, Kohei Uetsuka, Masaki Takeda, Daisuke Kiga School of Electrical Engineering and Bioscience, Department of Advanced Science and Engineering, Waseda University, Tokyo, Japan
*27P-188	Zombie cells produced from the minimal synthetic bacterium JCVI- syn3B <u>Nanase Oda</u> , Hana Kiyama, Makoto Miyata Graduate School Science, Osaka Metropolitan University, Japan
*27P-189	Large coiled-coil protein of Mycoplasma pneumoniae induces morphological changes in a minimal synthetic bacterium by inhibiting septum formation <u>Muhammad Algiffari</u> , Hana Klyama, Daisuke Nakane, Tsuyoshi Kenri, Makoto Miyata Graduate School of Science, Osaka Metropolitan University, Osaka, Japan

#### 27P-190 Efficient Proliferation of Synthetic Minimal Cells with Low Energy Costs

Ken Takagi, Minoru Kurisu, Toshihiro Kawakatsu, Masayuki Imai Department of Physics, Tohoku University, Sendai, Japan

#### **Computational biology: Bioinformatics**

- \*27P-191 Elucidation of mechanistic details of copper chaperoning to Superoxide Dismutase (SOD) using a novel free-energy computation technique and cross-validated with Molecular Dynamics Simulations Sharayu Umakant Ghodeswar, Debashree Bandyopadhyay Birla Institute of Technology and Science, Pilani - Hyderabad Campus, Telangana, India
- 27P-192 Computational analysis of OPRD1-OPRM1 heterodimer ligands <u>Ryota Takishima</u>, Aoi Fukushima, Wataru Nemoto Grad. Sci. & Eng., Tokyo Denki Univ., Saitama, Japan

#### Computational biology: Molecular simulation

- \*27P-193 A Gradient-Based Approach for Optimizing Molecular Structures using Atomic Force Microscopy Images and Normal Mode Analysis Xuan Wu, Osamu Miyashita, Florence Tama Department of Physics, Nagoya University
- \*27P-194 PINning down the elevator-type mechanism of auxin transport Lorena Zuzic, Bjørn Panyella Pedersen, Birgit Schiøtt Department of Chemistry, Aarhus University, Aarhus, Denmark
- \*27P-195 The Effect of Tricaprylin Surface on The Lid Region Dynamics of Candida antarctica Lipase B <u>Tegar Nurwahyu Wijaya</u>, Akio Kitao School of Life Science and Technology, Tokyo Institute of Technology, Tokyo, Japan/ Department of Chemistry, Universitas Pertamina, Jakarta, Indonesia

*27P-196	Development of drug discovery platform technology based on a generalized-ensemble simulation method -Evaluation of SARS CoV-2 PLpro candidate inhibitors- Masashi Muramoto, Suzuka Saitou, Simon Hikiri, Junichi Higo, Takuya Takahashi Graduate School of Life Sciences, Ritsumeikan University, Kusatsu, Japan.
*27P-197	Unraveling the Catalytic Mechanism of EPS1 in Salicylic Acid Biosynthesis Using Computational Modeling <u>Tianjie Li</u> , Yi Wang The Chinese University of Hong Kong
*27P-198	Complementary Analysis between 4D Crystallography and Extensive MD Simulation Captures Transient IF1-Ribosome Dynamics in Translation Initiation Shun Yokoi, Ilkin Yapici, E. Han Dao, Ebru Destan, Esra Ayan, Alaleh Shafei, Fatma Betul Ertem, Cahine Kulakman, Merve Yilmaz, Bilge Tosun, Halilibrahim Ciftci, Abdullah Kepceoglu, Jerome Johnson, Omur Guven, Ali Ergul, Brandon Hayes, Yashas Rao, Christopher Kupitz, Frederic P. Poitevin, Mengling Liang, Mark S. Hunter, Pohl Milon, Raymond G. Sierra, Ayori Mitsutake, Soichi Wakatsuki, Hasan DeMirci Department of Physics, School of Science and Technology, Meiji University, Kanagawa Japan/Biological Sciences Division, SLAC National Accelerator Laboratory, CA, USA/
*27P-199	The Regulatory Role of p53 C-Terminal Domain Acetylation in Modulating the Dynamics of SIR2's NAD+ Binding Pocket Zhen Bai, Tatsuhiro Kimizono, Akio Kitao Tokyo Institute of Technology
*27P-200	An Open Source de novo Drug Design Workflow with Active Learning and Enamine REAL Ben Cree Newcastle University
*27P-201	Studying the role of protonation in the (de)activation mechanism of class A GPCRs João Vitorino, Carlos Barreto, Irina Moreira, Miguel Machuqueiro BiolSI: Biosystems and Integrative Sciences Institute, Faculdade de Ciências, Universidade de Lisboa, Portugal

*27P-202	Molecular Dynamics simulation of the complex of the multiple distinctive structurral regions in the WNV envelope and human monoclonal antibody. <u>Masahito Yoshikawa</u> , Hideyuki Masaki, Ryuuichi Kato, Tatsuhiko Ozawa, Naoyuki Miyashita Department of Biological System Engineering, Graduate School of Biology-Oriented Science and Technology, KINDAI University
*27P-203	Identifying and Characterizing Ligands for Mutant p53 as Potential Breast Cancer Therapy <u>Rifqa Fikriya Rahasri</u> , Kazutomo Kawaguchi, Hidemi Nagao Kanazawa University
*27P-204	Molecular Docking, Molecular Dynamics, And MM-PBSA Analysis of Quinolone Antibiotics Againts FmtA of Staphylococcus aureus <u>Citra Hasanah</u> , Hidemi Nagao, Kazutomo Kawaguchi Graduate School of Natural Science and Technology, Kanazawa University
*27P-205	Computational analysis on binding structure of limonin to a bitter taste receptor TAS2R38 <u>Masamune Kashihara</u> , Daiki Hayashi, Shigenori Tanaka, Yoshiko Aihara Graduate School of Agricultural Science, Kobe University, Kobe, Japan
27P-206	Molecular Dynamics Study of the Unfolding Processes of Proteins with Highly Similar Native Structure Souta Kadowaki, Takashi Yoshidome Department of Applied Physics, Graduate School of Engineering, Tohoku University, Japan
27 <b>P-20</b> 7	Dynamic structure analysis of superoxide dismutase 1 protein upon Cys111 oxidation using molecular dynamics simulation <u>Yuta Hori</u> , Ayaka Sato, Kowit Hengphasatporn, Yasuteru Shigeta Center for Computational Sciences, University of Tsukuba
27P-208	Investigation of the effect of the 2-OH group of in Arabidopsis thaliana ceramide on plant cell membranes using MD simulation <u>Tsujii Keigo</u> , Minoru Nagano, Simon Hikiri, Takuya Takahashi Graduate School of Life Sciences, Ritsumeikan University, Kusatsu, Japan

27P-209	Dynamin-1 membrane tubule constriction mechanism revealed by coarse-grained MD simulations MD. Iqbal Iqbal Mahmood, Shintaroh Kubo, Kei-ichi Okazaki Research Center for Computational Science, Institute for Molecular Science, National Institutes of Natural Sciences, Okazaki, 444-8585, Japan.
27P-210	Heterogeneous organization in phase-separated transcription factors: Residue-revel molecular simulations <u>Azuki Mizutani</u> , Cheng Tan, Yuji Sugita, Shoji Takada Grad. Sch. Of Science, Kyoto Univ., Kyoto, Japan
27 <b>P</b> -211	Molecular dynamics simulation of the PWW domain of LEDGF protein and histone tail H3K36 <u>Hinako Suzuki</u> , Itoh Satoru, Hisashi Okumura Shinshu University/Institute for Molecular Science
27 <b>P-</b> 212	<b>MD-based in silico screening using supercomputer Fugaku</b> <u>Tomoya Nabetani</u> , Toru Ekimoto, Tsutomu Yamane, Mitsunori Ikeguchi Graduate School of Medical Life Science, Yokohama City University
27P-213	Effects of sodium ions on conformational changes of the adenosine A2A receptor by molecular simulations <u>Akihiro Arisawa</u> , Ayori Mitsutake Meiji University,School of Science and Technology
27P-214	Optimal transport maps for targeted free energy estimation <u>Tsuyoshi Kawai</u> , Yasuhiro Matsunaga Graduate School of Science and Engineering, Saitama University, Saitama, Japan
27P-215	Surface oleophilicity induced by UV-hydroxylation of titanium Gehoon Chung, Wonjoon Moon, Byeong-Min Lee, Shin Hye Chung Seoul National University School of Dentistry/Seoul National University Dental Research Institute

## **\*27P-216** Looking for non-opioid analgesics using stochastic titration CpHMD with AMBER14SB

<u>João G. N. Sequeira</u>, Adrian E. Roitberg, Miguel Machuqueiro BiolSI: Biosystems and Integrative Sciences Institute, 1749-016 Lisboa, Portugal

June 27 [Thu]

#### Prediction of cross-fitness for adaptive evolution to different environmental conditions: Consequence of phenotypic dimensional reduction. Takuya Sato, Chikara Furusawa, Kunihiko Kaneko BDR, RIKEN \*27P-218 Theoretical model of cell shape control by cytoskeleton Vivek Semwal, Biplab Bhattacherjee, Michiko Takeda, Yu-Chiun Wang, Tatsuo Shibata Laboratory for Physical Biology, RIKEN Center for Biosystems Dynamics Research, Kobe, Japan \*27P-219 Universally conserved Mg-pinch motif in NTP processing enzymes Balint Dudas, Denes Berta, Edina Rosta Department of Physics and Astronomy, University College London (UCL), United Kingdom

\*27P-217

\*27P-220 A binding site for phosphoinositide modulation of voltage gated sodium channels described by multiscale simulations Yiechang Lin, Elaine Tao, James Champion, Ben Corry Australian National University

#### 27P-221 Development of an Efficient Estimation Method for Maximum Tolerated Dose by Reinforcement Learning

Ryosuke Takami, Koji Tabata, Yayoi Wada, Masahiro Sonoshita, Tamiki Komatsuzaki

Research Institute for Electronic Science, Hokkaido University, Sapporo, Japan/ Institute for Chemical Reaction Design and Discovery (WPI-ICReDD), Hokkaido University, Sapporo, Japan/The Institute of Scientific and Industrial Research, Osaka University /Institute for Open and Transdisciplinary Research Initiatives, Osaka University

\*27P-222 Study of the Allosteric Mechanism of Human Mitochondrial Phenylalanyl-tRNA Synthetase by Transfer Entropy via an Improved Gaussian Network Model and Co-evolution Analyses

#### Zhongjie Han, Chunhua Li

Center for Quantitative Biology, Academy for Advanced Interdisciplinary Studies, Peking University, Beijing, China/Peking-Tsinghua Center for Life Sciences, Academy for Advanced Interdisciplinary Studies, Peking University, Beijing, China/Faculty of Environmental and Life Sciences, Beijing University of Technology, Beijing, China

Singapore

# 27P-223 Nucleosome-resolution modeling and simulation of singlegene level chromatin organization mechanisms <u>Gu Chenyang</u>, Shoji Takada, Giovanni Brandani Grad. Sch. Sci., Kyoto university, Kyoto, Japan 27P-224 Allosteric drugs: new principles and design approaches Wei-Ven Tee, Igor N Berezovsky Bioinformatics Institute (BII), Agency for Science, Technology and Research (A\*STAR), 30 Biopolis Street, #07-01, Matrix, Singapore 138671/Department of Biological Sciences (DBS), National University of Singapore (NUS), 8 Medical Drive, 117579,

#### 27P-225 Study of Liquid–liquid Phase Separation of Tau fragment K18 via Coarse-grained Simulation

#### Zhuqing Zhang, Qinglin Yan College of Life Sciences, University of Chinese Academy of Sciences, Beijing 100049, China

#### 27P-226 Design principles of microtubule-associated proteins: exploring the role of lever arms and linker regions under directional loads Ilya B. Kovalenko, Vladimir A. Fedorov, Ekaterina G. Kholina, Philipp S. Orekhov, Egor M. Pozdnyakov, Fazoil I. Ataullakhanov, <u>Nikita Gudimchuk</u> Lomonosov Moscow State University, Moscow, Russia/Center for Theoretical

Lomonosov Moscow State University, Moscow, Russia/Center for Theoretical Problems of Physico-Chemical Pharmacology, Russian Academy of Sciences, Moscow, Russia

#### Computational biology: machine learning for molecules or cell systems

#### \*27P-227 A Machine Learning Approach to Classify Force Curves of Nuclear Elasticity Measurements.

MD Fahim Newaz Division of Nano Life Science, Kanazawa University, Kanazawa 920-1192, Japan

#### \*27P-228 Development of an Efficient Estimation Method for Maximum Tolerated Dose by Reinforcement Learning Ryosuke Takami, Koji Tabata, Yayoi Wada, Masahiro Sonoshita, Tamiki Komatsuzaki

Research Institute for Electronic Science, Hokkaido University, Sapporo, Japan

\*27P-229 RVINN: Inference of gene regulation dynamics in the mRNA life cycle using Physics-Informed Neural Networks Osamu Muto, Zhongliang Guo, Rui Yamaguchi Nagoya University/Aichi Cancer Center Research Institute

#### Mathematical & Theoretical biology

- \*27P-230 Active thermodynamic force drives mitochondrial equidistant distribution in axons <u>Masashi K. Kajita</u>, Yoshiyuki Konishi, Tetsuhiro Hatakeyama Department of Applied Chemistry and Biotechnology, Faculty of Engineering, University of Fukui, Fukui, Japan
- \*27P-231 Global propagation of single-gene deletion effects through stoichiometry conservation relations <u>Genta Chiba</u>, Ken-ichiro Kamei F., Arisa Oda, Kunihiro Ohta, Yuichi Wakamoto Grad. Sch. Arts and Sci. Univ. Tokyo, Tokyo, Japan
- \*27P-232 Pattern propagation driven by surface curvature <u>Ryosuke Nishide</u>, Shuji Ishihara The University of Tokyo
- 27P-233 Entangled gene regulatory networks with cooperative expression endow responses to unforeseen environmental changes Masayo Inoue Graduate School of Engineering, Kyushu Institute of Technology

#### Nonequilibrium state & Biological rhythm

- \*27P-234 Exploring dense active dynamics in suspension of ciliate Tetrahymena based on all cell tracking Kohei Okuyama, Masatoshi Ichikawa Department of Physics, Kyoto University
- 27P-235 Emergence of spontaneous oscillations in a liquid film of bacterial swimmers Lei-Han Tang

Hong Kong Baptist University, Hong Kong, China

## **Poster Sessions**

Measurements		
*27P-236	Single EVs detection and analysis using a glass nanopore Izumi Shibayama, Kohei Hayashi, Ryuji Kawano Department of Biotechnology and Life Science, Tokyo University of Agriculture and Technology	
*27P-237	Ultra-low-noise and wide-bandwidth current detection for enhanced scanning ion conductance imaging rate in scanning ion conductance microscopy Shoma Kamei, Shinji Watanabe Division of Nano Life Science, Graduate School of Frontier Science Initiative, Kanazawa University	
27P-238	Aptamer-based Al-driven nanopore measurement for the simultaneous detection of biomarkers for the cancer diagnosis <u>Ryo Akita</u> , Lysenko Artem, Shunsuke Ono, Hikaru Nozawa, Tatsuhiko Tsunoda, Sotaro Uemura Graduate School of Science, The University of Tokyo	
27P-239	Analysis of the conformational dynamics of oligosaccharides using ion mobility spectrometry <u>Hao Feng</u> , Takumi Yamaguchi School of Materials Science, Japan Advanced Institute of Science and Technology	
Bioimaging		
*27P-240	A bright and highly-response Ca2+ biosensor based on mScarlet: Progress toward fluorescence lifetime imaging Shosei Imai, Ryan Fink, Takuya Terai, Olivia A. Masseck, Robert E. Campbell Department of Chemistry, Graduate School of Science, The University of Tokyo, Tokyo, Japan.	
*27P-241	Hydroxyquinoline-derived Multifunctional Small Molecule Turn-On Fluorescent Probe as a Theranostic Agent for Alzheimer's Disease Priyam Ghosh, Parameswar Iyer Department of Chemistry, Indian Institute of Technology Guwahati, Guwahati, Assam 781039, India	

Poster Sessions

June 27 [Thu]

## \*27P-242 Precision-enhanced 1,000-fold faster 3D quantum thermometry in vivo

<u>Yurina Nakane</u>, Haruka Maeoka, Ryuki Imamura, Ryuji Igarashi, Shin Usuki, Takuma Sugi

Program of Biomedical Science, Graduate School of Integrated Sciences for Life, Hiroshima University, Japan

## \*27P-243 Elucidation of neuronal differentiation mechanisms by thermal signaling through control of intracellular local temperature Shunsuke Chuma, Kohki Okabe, Yoshie Harada

Department of Biological Sciences, Graduate School of Science, Osaka University, Osaka, Japan, /Institute for Protein Research, Osaka University, Osaka, Japan

#### \*27P-244 Imaging and modeling of glycolytic oscillations

<u>Saaya Hario</u>, Shosei Imai, Yudai Iyoda, Hikaru Sugimoto, Takuya Terai, Shinya Kuroda, Robert E. Campbell Department of Chemistry, Graduate School of Science, The University of Tokyo, Tokyo, Japan

## \*27P-245 Development of selective plane activation structured illumination microscopy

<u>Kenta Temma</u>, Ryosuke Oketani, Toshiki Kubo, Kazuki Bando, Shunsuke Maeda, Kazunori Sugiura, Tomoki Matsuda, Rainer Heintzmann, Tatsuya Kaminishi, Koki Fukuda, Maho Hamasaki, Takeharu Nagai, Katsumasa Fujita

Department of Applied Physics, Osaka University/AIST Advanced Photo-BIO OIL, AIST-Osaka University/Institute for Open and Transdisciplinary Research Initiatives, Osaka University

## \*27P-246 High-speed, high-resolution computational phase microscopy visualizing organelles

<u>Yugo Inutsuka</u>, Yasushi Okada The University of Tokyo/RIKEN

## 27P-247 Extraction of dependent spatial or spectral features from different disease states in Raman images

<u>Ryoya Kondo</u>, Yuta Mizuno, Jean-Emmanuel Clement, Kentaro Mochizuki, Katsumasa Fujita, Yoshinori Harada Grad. Sch. Chem. Sci. Eng., Hokkaido Univ.

*27P-248	Quantification of Spatial and Spectral Information Dependent on Measurement Methods and Disease States in Raman Images Ryoya Kondo, Yuta Mizuno, Jean-Emmanuel Clement, Kentaro Mochizuki, katsumasa Fujita, Yoshinori Harada, Tamiki Komatsuzaki Grad. Sch. Chem. Sci. Eng., Hokkaido Univ.
27P-249	Simultaneous measurement of average size and number of biomolecular condensates using spatial image correlation spectroscopy (SICS) Yuta Hamada, Akita Kitamura Graduate School of Life Science, Hokkaido University, Sapporo, Japan
27P-250	Imaging of Biomolecules by Constant Thermal Fluctuation Mode Atomic Force Microscopy Daisuke Yamamoto Faculty of Science, Fukuoka University
27P-251	Development of small nanodiamonds that can be observed by optically detected magnetic resonance inside cells. <u>Hirotaka Okita</u> , Shingo Sotoma, Yuki S Kato, Yukiho Shimazaki, Hiroshi Abe, Seiichi Saiki, Madoka Suzuki, Yoshie Harada Institute for Protein Research Osaka University
27P-252	Volumetric imaging of micrometer-scale cellular dynamics in centimeter-scale multicellular systems <u>Taro Ichimura</u> , Taishi Kakizuka, Keiko Itano, Kaoru Seiriki, Hitoshi Hashimoto, Yuki Sato, Hiroya Itoga, Shuichi Onami, Takeharu Nagai Institute for Open and Transdisciplinary Research Initiatives, Osaka University, Japan
27P-253	Multi-color fluorescence lifetime biosensors for quantifying Ca2+, ATP, and GTP/GDP ratio in live cells Cong Quang Vu WPI-NanoLSI, Kanazawa University

#### Bioengineering

## \*27P-254 Cell-free synthesis of hydrophobic peptides that form nanopores in bilayer lipid membranes

<u>Shoko Fujita</u>, Izuru Kawamura, Ryuji Kawano Department of Biotechnology and Life Science, Tokyo University of Agriculture and Technology, Tokyo, Japan

#### 27P-255 Morphological Difference in Hydrogel Induced Cancer Stem Cell in Synovial Sarcoma Model Cells

Zannatul Ferdous, Jean-Emmanuel Clément, Jian Ping Gong, Shinya Tanaka, Masumi Tsuda, Tamiki Komatsuzaki Institute for Chemical Reaction, Design and Discovery (WPI-ICReDD), Hokkaido University

#### \*27P-256 Novel approach for anticancer peptides carried by nanoparticles Roberta Moisa Horia Hulubei National Institute for Physics and Nuclear Engineering

#### \*27P-257 Stereo 3D reconstruction of a dragonfly flapping motion and its quantification using fine grid spotlight Natsuki Yamamoto

Akita Prefectural University

#### 27P-258 Microscopic toxicity assay of human organoids in microfluidic devices advanced by quantum beam technologies

Kotaro Oyama, Tomoko G Oyama, Hiroki Hamaguchi, Yusuke Kimura, Atsushi Kimura, Kimio Yoshimura, Masaaki Omichi, Yuuji Ueki, Akihiro Hiroki, Hiroyuki Hoshina, Yasuhiro Oshima, Michiyo Suzuki, Shinichiro Mori, Noriaki Seko, Noriko Ishioka, Mitsumasa Taguchi Takasaki Institute for Advanced Quantum Science, National Institutes for Quantum Science and Technology, Gunma, Japan

#### Crystal growth & Crystallization technique

## \*27P-259 High-Speed AFM investigation of structured fats' crystallization dynamics

<u>Anis Chikhoune</u>, Jan Kyselka, Djamel Eddine Chafai Équipe PVNTA, Laboratoire ALIMENTS, École Supérieure des Sciences de l'Aliment et des Industries Agroalimentaires (ESSAIA), Avenue Ahmed Hamidouche Route de Beaulieu, El Harrach 16200, Alger, Algeria

27P-260 Advancing Structural Biology: Innovations and Applications of In Vivo Macromolecular Crystallography at Nagoya University Etsuko Tokunaga, Swagatha Ghosh, Hiroki Onoda, Yasufumi Umena, Leonard M.G. Chavas NUSR, Nagoya Univ., Aichi, Japan/Dept. of Appl. Phys., Grad. Sch. of Eng., Nagoya Univ., Aichi, Japan

#### Virus structure, function, SARS-CoV-2

- \*27P-261 Cryo-EM structure of infectious and non-infectious Human Astrovirus and insights into its maturation process Kentaro Hiraka, Raymond Burton-Smith, Chihong Song, Kana Miyamoto, Kei Haga, Reiko Todaka, Kazuhiko Katayama, Kazuyoshi Murata National Institute for Physiological Sciences, National Institutes of Natural Sciences/ Exploratory Research Center on Life and Living Systems (ExCELLS), National Institutes of Natural Sciences
- \*27P-262 Structural basis for antiviral activity of a nucleoside analogue targeting dengue virus RNA-dependent RNA polymerase Shiori Ito, Shunsuke Kita, Kentaro Uemura, Yuki Iwama, Takashi Tadokoro, Hirofumi Sawa, Akihiko Sato, Akira Matsuda, Katsumi Maenaka Facul. Pharm. Sci., Hokkaido Univ., Japan
- 27P-263 DEVELOPING BROAD SPECTRUM ANTIVIRALS: PEPTIDE-PORPHYRIN CONJUGATES ACTION, FROM MOLECULAR SCALE TO IN VIVO

Miguel A. R. B. Castanho IMM, Instituto de Medicina Molecular, Lisbon, Portugal

27P-264 Conformational dynamics of SARS-CoV-2 spike protein investigated by single molecule fluorescence spectroscopy Yuji Itoh, Taisei Mori, Tateki Suzuki, Takao Hashiguchi, Satoshi Takahashi

IMRAM, Tohoku Univ., Miyagi, Japan/Grad. Sch. Life Sci., Tohoku Univ., Miyagi, Japan

Poster Sessions

#### Mechanosensing and Mechanobiology, Biological Temperature

## \*27P-265 Investigation of the mechanism of neurite outgrowth using nuclear heating

Yukiho Shimazaki, Shunsuke Chuma, Kohki Okabe, Yoshie Harada Department of Biological Sciences, Graduate School of Science, Osaka University, Osaka, Japan/Institute for Protein Research, Osaka University, Osaka, Japan

- \*27P-266 Small-molecule FLIM sensors for visualization of temperature in calcium cycling of sarcoplasmic reticulum <u>Takeru Yamazaki</u>, Kayoko Nomura, Toshiko Yamazawa, Satoshi Arai WPI Nano Life Science Institute (WPI-NanoLSI), Kanazawa Univ., Ishikawa, Japan
- 27P-267 Differential roles of a periplasmic tension sensor and a cytoplasmic one in the channel opening of MscL <u>Takeshi Nomura</u>, Yasuyuki Sawada, Masahiro Sokabe School of Human Science and Environment, University of Hyogo, Hyogo, Japan

#### **Biophysics of disease**

#### \*27P-268 Tau Inclusions in Soma Induce Neuronal Death in Human iPSCderived Neurons

Naoki Kato, Sumihiro Maeda, Hideyuki Okano, Hiroko Bannai School of Advanced Science and Engineering, Waseda University

\*27P-269 Reversible tangle formation of Alzheimer's disease-fold Tau filaments by conformational changes of the fuzzy coat region Shingo Tamai, Takashi Nomura, Ryohei Kojima, John Burke, Atsushi Yamagata, Mikako Shirouzu, Takeshi Fukuma, Motomasa Tanaka Lab. for Protein Conformation Diseases, RIKEN CBS/Biomedical Sciences & Engineering Track, Tokyo Medical and Dental University

# **27P-270** Lead and Mercury poisoning promote cardiac dysfunction in isolated hearts affecting cardiac ion channels and intracellular calcium homeostasis.

<u>Gonzalo R. Ferreira</u>, Romina Cardozo, Axel Santander, Luisina Chavarria, Santiago Sastre, Milagros Benitez, Nicolas Mujica, Lucia Dominguez, Garth Lamb Nicolson

Laboratory of Ion channels, Biological Membranes and Cell Signaling. Department of Biophysics. Facultad de Medicina. Montevideo. Universidad de la Republica. Uruguay.

#### **Miscellaneous topics**

- \*27P-271 Novel antibacterial agents to treat Multidrug resistant bacteria causing wound infections in diabetic patients <u>Mithali Raj Marla</u>, Shailaja Raj Marla, Maria shajan Kamineni Institute of Medical Sciences
- 27P-272 An Interactive 3-D Graph Tool to Visualize Electromagnetic Waves on Web Browsers for Physics Education Satoshi Yamaguchi, Masayuki Irisa Comp. Sci. and Sys. Eng., Kyushu Inst. Tech., Japan