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

*26P-001	Unravelling Protein Complexity with 3Di-based Structural Entropy Zecheng Zhang, Qian-Yuan Tang Hong Kong Bapist University
*26P-002	Frustration-Fluctuation Correspondence in Enzymes Yuxiang Zheng, Qian-Yuan Tang Hong Kong Baptist University
*26P-003	Cryo-EM Structure of the hERG Channel Complexed with a K+ Channel Blocker Yasuomi Miyashita, Toshio Moriya, Masato Kawasaki, Satoshi Ogasawara, Naruhiko Adachi, Satoshi Yasuda, Tetsuichiro Saito, Toshiya Senda, Takeshi Murata Developmental Biology, Graduate School of Medicine, Chiba University/Membrane Protein Research Center, Graduate School of Science, Chiba University
*26P-004	Crystal Structure of Pectocin M1 from Pectobacterium carotovorum: Unveiling Diverse Conformations and Binding Interactions during the Initial Step of Pectocin M Uptake through the Ferredoxin Uptake System Nawee Jantarit, Hideaki Tanaka, Genji Kurisu Protein Crystallography Laboratory, Institute for Protein Research, Osaka University, Suita, Osaka Japan/Department of Macromolecular Sciences, Graduate School of Science, Osaka University, Toyonaka, Osaka, Japan

*26P-005	Symmetry Matched Protein – Macrocycle Assembly
	<u>Colin Wren</u> , Ronan J. Flood, Niamh M. Mockler, Martin Savko, Qiang Shi, Peter B. Crowley
	School of Biological and Chemical Sciences, University of Galway, H91 TK33, Galway, Ireland
*26P-006	Structure of Francisella tularensis subsp. novicida Cas9 in the catalytically poised state
	<u>Shinsuke Higashiyama</u> , Ryoya Nakagawa, Hisato Hirano, Osamu Nureki Department of Biological Sciences, Graduate School of Science, University of Tokyo, Tokyo, Japan
*26P-007	Cryo-EM structure of a photosystem I supercomplex from oleaginous green alga Coccomyxa subellipsoidea
	<u>Pi-Cheng Tsai</u> , Fusamichi Akita, Jian-Ren Shen Research Institute for Interdisciplinary Science, Okayama University, Okayama, Japan
*26P-008	Structural analysis of the photosystem I-light harvesting I supercomplex from a cryptophyte alga Rhodomonas sp. Wenyue Zhang Graduate school of natural science and technology & Research Institute for Interdisciplinary Science, Okayama University, Okayama, Japan.
*26P-009	Structural analysis of brain-associated proteins in complex with novel PET radiotracers Kaede Goto, Junta Tomono, Shozo Furumoto, Nobuyuki Okamura, Ryuichi Harada, Takeshi Yokoyama, Yoshikazu Tanaka Grad. Sch. Life Sci., Tohoku Univ., Miyagi, Japan.
26P-010	Cryo-EM structure of the zinc-activated channel (ZAC) in the Cys-loop receptor superfamily Fei Jin, Yi-Yu Lin, Ru-Chun Wang, Yimeng Zhao, Cheng Shen, Danqi Sheng, Muneyoshi Ichikawa, Ye Yu, Jin Wang, <u>Motoyuki Hattori</u> Fudan University, Shanghai, China
26P-011	The conformation and its thermal stability of antibiotic peptide alamethicin in alcohol solution studied by NMR Yoshinori Miura Center for Advanced Instrumental Analysis, Kyushu University

26P-012 A novel blue-carotenoprotein from sponge <u>Momose Kuroda</u>, Yui Fujita, Momoko Ishida, Satoko Matsunaga, Mitsuru Jimbo, Takeshi Yokoyama, Yoshikazu Tanaka, Ryuichi Sakai NIT, Hakodate College

26P-013 Structure-activity Relationship of a novel enzyme derived from marine Streptomyces

<u>Takumi Oshiro</u>, Shuta Uehara, Yoshikazu Tanaka, Takuya Ito, Yoshio Kodera, Takashi Matsui Grad. Sch. Sci., Kitasato Univ.

Protein: Structure & Function

*26P-015	Proton conduction mechanism in FO rotary motor studied by quantum molecular dynamics simulation
	Yukinari Kamiyama, Dan Parkin, Junichi Ono, Yoshifumi Nishimura,
	Hiromi Nakai, Mitsunori Takano
	Grad. Scl. Adv. Sci. & Eng., Waseda Univ., Tokyo, Japan
*26P-016	Time-resolved crystallography for the study of a B12-dependent photoreceptor using X-ray free-electron lasers
	Ronald Rios Santacruz, Giorgio Schiro, David Leys, Nigel Scrutton,
	Tosha Takehiko, Kensuke Tono, Derren Heyes, Harshwardhan Poddar, Martin Weik
	Univ. Grenoble Alpes, CEA, CNRS, Institut de Biologie Structurale, F-38044 Grenoble, France
*26P-017	Heterogeneity of Microtubule Lattices Revealed by Cryo-ET and Non-averaging Structural Analysis
	Hanjin Liu, Hiroshi Yamaguchi, Masahide Kikkawa, Tomohiro Shima
	Graduated School of Science, The University of Tokyo, Japan
*26P-018	Structural-dynamics insight of an alligator-derived antimicrobial peptide, AsCATH5, in interaction with membrane mimetics as revealed by solution NMR and MD simulation
	<u>Jeremia Oktavian Chrisnanto,</u> Kohei Kano, Mitsuki Shibagaki,
	Tefera Dessalegn Abeje, Hirai Fumi, Yasuhiro Kumaki, Hiroyuki Kumeta,
	Tomoyasu Aizawa
	Hokkaido University

Poster Sessions

June 26 [Wed]

*26P-019	Discovery and structural characterization of novel pore-forming toxins <u>Jana Susanne Anton</u> , Fernando Meireles, Juan F. Bada, Maria J. Marcaida, Matteo Dal Peraro Laboratory for Biomolecular Modeling, École polytechnique fédérale de Lausanne, Switzerland
*26P-020	Functional annotation of Cysteine Post-Translational Modifications based on protein sequences and structures and development of a consolidated Cysteine database (CysDBase) Devarakonda Himaja, Dr. Debashree Bandyopadhyay Birla Institute of Technology and Science, Hyderabad Campus , INDIA
*26P-021	Crystal structure and proton transporting mechanism of viral heliorhodopsin, V2HeR3 Ritsu Mizutori, Nipawan Nuemket, Shoko Hososhima, Sayaka Ohashi, Satoshi Tsunoda, Yuji Furutani, Oded Baja, Eriko Nango, Kota katayama, Hideki Kandori Nagoya Institute of Technology
*26P-022	Theoretical study on the photoactivation state of chloride pump NpHR using QM/MM Method Tomo Ejiri, Ryo Oyama, Shigehiko Hayashi Department of Chemistry, Graduate School of Science, Kyoto University, Kyoto, Japan
*26P-023	Investigating allosteric communication with ultrahigh-resolution X-ray crystallography <u>Caitlin Emma Hatton</u> , Pedram Mehrabi Institute for Nanostructure and Solid-State Physics, Universität Hamburg, Hamburg, Germany
*26P-024	Development of supramolecular micelles promoting oxidative protein folding under a crowded environment <u>Mai Kitamura</u> , Takahiro Muraoka Graduate School of Engineering, Tokyo University of Agriculture and Technology, Tokyo, Japan

*26P-025	Insights into the Cytochrome c Oxidase proton pumping mechanism from constant-pH MD simulations
	Ines Domingos Silva Pires, António M. Baptista, Miguel Machuqueiro BiolSI - Biosystems and Integrative Sciences Institute, Faculdade de Ciências, Universidade de Lisboa, Campo Grande, Ed. C8, Lisboa, Portugal
*26P-026	Mass photometry as a novel single molecule approach to study immunoglobulin binding protein (BiP) oligomerization and function Karina New, Miguel Lagos, Roi Asor, Zahra Alavi, Philipp Kukura, Christian Wilson Faculty of Chemical and Pharmaceutical Sciences, University of Chile, Santiago, Chile
*26P-027	Catalytic mechanism of the cytosolic θ type carbonic anhydrase from marine diatom Phaeodactylum tricornutum <u>Hiroto Negoro</u> , Hideaki Tanaka, Ginga Shimakawa, Hiroyasu Koteishi, Akihiro Kawamoto, Yusuke Matsuda, Genji Kurisu Institute for Protein Research, Osaka University/Department of Biotechnology, Graduate School of Engineering, Osaka University
*26P-028	Structure of nitric oxide reductase dimer revealed by single particle cryo-electron microscopy Ryohei Kawakami, Chai Gopalasingam, Hideki Shigematsu, Takehiko Tosha Grad. Sch. Sci., Univ. Hyogo, Hyogo, Japan/RIKEN SPring-8 Center, Hyogo, Japan
*26P-028 26P-029	cryo-electron microscopy Ryohei Kawakami, Chai Gopalasingam, Hideki Shigematsu, Takehiko Tosha

Poster Sessions

June 26 [Wed]

26P-031 Mechanism of Formate Oxidation by NAD-dependent Formate Dehydrogenase: Computational Studies on Near-Attack Conformations Jiri Kozelka, Yevgen Yurenko, Jan Novotný, Radek Marek Department of Condensed Matter Physics, Faculty of Science, Masaryk University, Brno, Czech Republic/Université Paris Cité, DSIMB UMR S U1134, Paris, France 26P-032 Imaging and inhibition analysis of human serum amyloid A aggregation using guantum dots Liangguan Shi, Tuya Gegen, Kuragano Masahiro, Tokuraku Kiyotaka Graduate School of Engineering, Muroran Institute of Technology 26P-033 Real-time Imaging and Inhibition Analysis of Amylin Aggregations Using Quantum Dot nanoprobes Xiaoyu Yin, Ziwei Liu, Tuya Gegen, Hayate Sawatari, Keiya Shimamori, Masahiro Kuragano, Kiyotaka Tokuraku Graduate School of Engineering, Muroran Institute of Technology 26P-034 Structural and functional analyses of YeeE/YeeD complex in thiosulfate uptake pathway Mai Ikei, Ryoji Miyazaki, Keigo Monden, Yusuke Naito, Azusa Takeuchi, Yutaro S. Takahashi, Yoshiki Tanaka, Keina Murata, Takaharu Mori, Muneyoshi Ichikawa, Tomoya Tsukazaki Fudan University 26P-035 TRANSPLANTATION OF ENZYMATIC FUNCTION BY EXCHANGE OF

P-035 TRANSPLANTATION OF ENZYMATIC FUNCTION BY EXCHANGE OF FUNCTION ELEMENTS

<u>Mikio Kataoka</u>, Yoichi Yamazaki, Yasushi Imamoto, Hironari Kamikubo Nara Institute of Science and Technology

Protein: Physical property

*26P-037 Thermal Boundary Conductance at the Protein–Water Interface <u>Futa Yoshimura</u>, Takahisa Yamato Graduate School of Science, Nagoya University, Nagoya, Japan

Structural Insights into Switching Mechanisms between Spontaneous Folding and Ligand-Induced Folding of Staphylococcal Nuclease Yujiro Mori, Issei Suzuki, Shogo Fukazawa, Kosuke Miki, Heinrich Roder, Kosuke Maki Grad. Schl. Sci., Nagoya Univ., Aichi, Japan/Fox Chase Cancer Ctr., Pennsylvania, United States 26P-039 α B-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 *26P-040 The function of multiple aggregates formed by the tumor suppressor protein p53 Emi Hibino, Reiji Hijikata, Takeshi Tenno, Hidekazu Hiroaki Grad. Scl. Pharm. Sci, Nagoya Univ. *26P-041 Spatiotemporal formation of a single liquid-like condensate of α -synuclein and subsequent aging by optical trapping

Keisuke Yuzu, Ching-Yang Lin, Po-Wei Yi, Chih-Hao Huang, Hiroshi Masuhara, Eri Chatani Graduate School of Science, Kobe University, Kobe, Japan/Department of Applied Chemistry, National Yang Ming Chiao Tung University, Hsinchu, Taiwan

26P-042 The oligomeric state is essential for fibroin nanofiber formation. Haruya Kajimoto, Kento Yonezawa, Takehiro Sato, Kok Sim Chan, Kiichi Hyashi, Takuya Sawai, Yusuke Okamoto, Rakuri Aiba, Yoichi Yamazaki, Sachiko Toma-Fukai, Hironari Kamikubo Division of Materials Science, Graduate School of Science and Technology, Nara Institute of Science and Technology, 8916-5, Takayama, Ikoma, Nara, Japan

Protein: Function

*26P-038

*26P-043 pH-gating mechanism of the bacterial inner membrane urea channel HpUrel of Helicobacter pylori Sahar Shojaei, Anna Stoib, Tobias Putz, Nađa Stević, Christine Siligan, Andreas Horner Institute of Biophysics, Johannes Kepler University Linz, Gruberstr. 40, 4020 Linz, Austria

*26P-044	Yeast complementation assays as a screening tool for urea, water, and ammonia permeability of membrane channels Anna Stoib, Sahar Shojaei, Felix Wolkenstein, Sandra Posch,
	Christine Siligan, Andreas Horner
	Institute of Biophysics, Johannes Kepler University Linz, Gruberstr. 40, 4020 Linz, Austria
*26P-045	Enhanced cell-membrane fluidity mediated by antifreeze proteins mitigates hypothermic injury to cells
	<u>Yue Yang</u> , Tatsuya Arai, Sakae Tsuda, Kazuhiro Mio, Yuji C. Sasaki
	Graduate School of Frontier Sciences, the University of Tokyo, Chiba, Japan
26P-046	Towards longer luminescence lifetime of the minimal luciferase picALuc
	Yuki Ohmuro-Matsuyama, Genta Kamiya, Kento Motoyama, Mitsuru Hattori, Ryogo Takai, Nobuo Kitada, Takeharu Nagai, Shojiro Maki, Hayato Matsui, Tadaomi Furuta
	Technology Research Laboratory, Shimadzu Corporation
26P-047	High-speed atomic force microscopy reveals functional dynamics of FnCas9
	Hideaki Tsukada

Mikihiro Shibata

Protein: Measurement & Analysis

- *26P-048 Development of Liquid Sample Observation Methods for Pulse-Electron Microscope Ryoya Katayama, Takeru Yamasaki, Tomoharu Matsumoto, Akihiro Narita Grad. Sch. of Sci., Nagoya Univ.
- *26P-049 Light-induced structural changes of heliorhodopsin 48C12 studied by using surface-enhanced infrared absorption spectroscopy <u>Tatsuya Sakamoto</u>, Soichiro Kato, Jingyi Tang, Insyeerah Jauhari, Yuji Furutani Graduate School of Engineering, Nagoya Institute of Technology, Aichi, Japan

*26P-050 Nanopore-based peptidome analysis based on the protein-protein interactions. Misa Yamaji, Ayaka Nakada, Kota Naito, Yoshikazu Tanaka, Ryuji Kawano Department of Biotechnology and Life Science, Tokyo University of Agriculture and Technology, Tokyo, Japan. 26P-051 Protein Acetylation Tracked using UV-Visible Absorption, Luminescence Spectroscopy and TD-DFT calculations Rajaram Swaminathan, Himanshi Devi, Simangka Borsaikia, Apoorva Badaya, Ravindra Venkatramani Department of Biosciences and Bioengineering, Indian Institute of Technology Guwahati, Assam, India 26P-052 Development of cysteine-specific modification technique for the quantitative analysis Arisa Suto, Yoshio Kodera, Takashi Matsui Grad. Sch. Sci., Kitasato Univ. 26P-053 Analysis of membrane translocation of Clostridioides difficile binary toxin using electrophysiologicaltechniques Yuki Mitani, Sotaro Takiguchi, Ryuji Kawano, Hideaki Tsuge Graduate School of Life Science, Kyoto Sangyo University, Kyoto, Japan. 26P-054 Simple and Efficient Detection Scheme of Two-Color Fluorescence Correlation Spectroscopy for Protein Dynamics Investigation from Nanoseconds to Milliseconds Yutaka Sano, Yuji Itoh, Atsuhito Fukasawa, Hiroyuki Oikawa, Satoshi Takahashi Tohoku University **Protein: Design & Engineering**

 *26P-055 Computational Design of engineered NT-193 antibody with broad activity against SARS-CoV-2 variant. Xu Pan, Hisham M. Dokainish, Katsumi Maenaka Faculty of Pharmaceutical Sciences, Hokkaido University
 *26P-056 In silico/in vitro evolution of peptide nanopore with β-barrel structure Mana Sato, Shoko Fujita, Tomoaki Matsuura, Ryuji Kawano Department of Biotechnology and Life Science, Tokyo University of Agriculture and Technology, Tokyo, Japan

*26P-057	The molecular basis through which Fv-supercharging affects the physicochemical properties of antibodies <u>Keisuke Kasahara</u> , Daisuke Kuroda, Jose Caaveiro, Satoru Nagatoishi, Kouhei Tsumoto Department of Bioengineering, School of Engineering, The University of Tokyo
*26P-058	Dynamics and Evolution of Uniform Substrate Binder <u>Yusran Abdillah Muthahari</u> , Paola Laurino Okinawa Institute of Science and Technology (OIST) Graduate School
*26P-059	Tandem artificial nucleocapsid to package longer RNA genome and expand protein architectures <u>Hualin Li</u> , Naohiro Terasaka Earth-Life Science Institute, Tokyo Institute of Technology, Tokyo, Japan
*26P-060	Exploring the design rules for artificial phase-separating peptides based on natural phase-separating protein sequences Joe Mori, Atsumi Hando, Satoshi Takahashi, Keisuke Ikeda, Kiyoto Kamagata Institute for Multidisciplinary Research for Advanced Materials, Tohoku University, Sendai, Japan/Department of Chemistry, Graduate School of Science, Tohoku University, Sendai, Japan
26P-061	Development of various fluorescence lifetime sensors using mTurquoise2 platform Chongxia Zhong, Satoshi Arai, Yasushi Okada 1. WPI Nano Life Science Institute, Kanazawa University, Kanazawa, Japan/2. Laboratory for Cell Polarity Regulation, RIKEN Center for Biosystems Dynamics Research, RIKEN, Osaka, Japan
26P-062	Darwinian ultrahigh-throughput evolution of biomolecules with in vitro compartmentalized gene amplification races <u>Taro Furubayashi</u> , Thibault Di Meo, Yoshihiro Minagawa, Hiroyuki Noji, Yannick Rondelez The University of Tokyo, Tokyo, Japan
26P-063	Magnetic bead-based protein display screening system for quantitative selection and evolution of functional proteins Shingo Ueno, Fumi Toshioka, Shoichi Tsuchiya, Takanori Ichiki Innovation Center of NanoMedicine (iCONM), Kawasaki Institute of Industrial Promotion

Protein: Intrinsic disorder

*26P-064

between ultrasonic cavitation and surfactants <u>Tomoki Ota</u>, Nakajima Kichitaro, Yamaguchi Keiichi, Goto Yuji, Ogi Hirotsugu Osaka University, Graduate School of Engineering *26P-065 The role of phase transitions of biopolymers in the formation and functioning of A-bodies <u>Anastasiia Gavrilova</u>, Yakov Mokin, Anna Fefilova, Aleksander Fonin Laboratory of structural dynamics, stability and folding of proteins, Institute of Cytology, Russian Academy of Sciences, St. Petersburg 194064, Russia

Improvement of detection ability for amyloid fibril seeds by interaction

*26P-066 High-Speed Atomic Force Microscopy Reveals Structural Dynamics of Microtubule-Associated Protein Tau Aggregation. <u>Tatsuya Kimura</u>, Kenjiro Ono, Ken-ichi Umeda, Daiki Muramatsu, Hiroki Konno, Noriyuki Kodera, Toshio Ando, Takahiro Nakayama WPI-Nano Life Science Institute, Kanazawa University, Kanazawa, Japan

26P-067 From the single-chain behavior to phase behavior of intrinsically disordered proteins Xiangze Zeng Hong Kong Baptist University

Heme proteins

*26P-068	Organ-specific probing of mitochondrial and lipid properties in Caenorhabditis elegans with Raman spectroscopy and imaging Evelina Nikelshparg, Mariela Pavan, Anat Ben-Zvi Ben-Gurion University Of The Negev, Faculty Of Natural Sciences, Life Sciences Department
26P-069	Purification and characterization of cholate-free cytochrome c oxidase from bovine heart Kyoko Shinzawa-Itoh, Kenta Tsutumi, Tomohiro Ide, Seishiro Mori, Eiki Yamashita, Kazumasa Muramoto Graduate School of Science, University of Hyogo

Membrane proteins

*26P-070	Time-resolved FTIR study of light-driven ion pump rhodopsin mutants converted from sodium to chloride pump. <u>Masahiro Yamamoto</u> , Hideki Kandori, Yuji Furutani Graduate School of Engineering, Nagoya Institute of Technology, Aichi, Japan
*26P-071	Detection of TEV protease by pore blockage in Outer Membrane Protein-G Nanopore mutation <u>Haruka Suzuki</u> , Toshiyuki Tosaka, Koki Kamiya Graduate School of Science and Engineering, Gunma University, Gunma, Japan
*26P-072	Construction and evaluation of the mutant β-barrel outer membrane protein nanopore <u>Toshiyuki Tosaka</u> , Koki Kamiya Graduate School of Science and Technology, Gunma University, Gunma, Japan
*26P-073	Detection of Polypeptide Related to Membrane Fusion through Nanopore MscL and Interaction based on Proteins Function Weibo Liang State Key Laboratory of Biotherapy, Sichuan University, Chengdu, China
*26P-074	Time-resolved infrared dual-comb spectroscopy using quantum cascade lasers reveals differences in conformational changes of two heliorhodopsins found from a bacterium and an archaeon <u>Toshiki Nakamura</u> , Soichiro Kato, Ryo Yamamoto, Manish Singh, Hideki Kandori, Yuji Furutani Department of Life Science and Applied Chemistry, Nagoya Institute of Technology, Nagoya, Japan,
*26P-075	Exploration of physical properties in streptomyces heliorhodopsin and the physiological function in the native cells <u>Koyo Yamada</u> , Rei Abe-Yoshizumi, Toshiki Nakamura, Yuji Furutani, Tatsuro Nishikino, Hideki Kandori Grad. Sch. of Eng., Nagoya Inst. of Tech., Aichi, Japan
26P-076	Rottlerin as an aquaporin-3 inhibitor for cancer therapy Inês Paccetti Alves, Marta Baptista, Catarina Pimpão, Inês V. da Silva, Bruno L. Victor, Graça Soveral Research Institute for Medicines (iMed.ULisboa), Faculty of Pharmacy, Universidade de Lisboa, Lisbon

Poster Sessions

June 26 [Wed]

26P-077 An attempt at high-resolution structural analysis of membrane proteins reconstituted into liposomes Atsuki Nakano, <u>Takaya Kawauchi</u>, Ren Kobayashi, Yuto Muto, Taichi Tsuyama, Ken Yokoyama Kyotosangyo University 26P-078 Reconstitution and nanoscale visualization of cadherin clusters on

F-078 Reconstitution and nanoscale visualization of cadnerin clusters on supported lipid bilayer in solution Shigetaka Nishiguchi, Tadaomi Furuta, Yui Kanaoka, Takayuki Uchihashi Exploratory Research Center on Life and Living Systems, National Institutes of Natural Sciences, Okazaki, Japan/Current affiliation: Department of Biotechnology, Osaka University, Suita, Japan

26P-079Protonophoric function of the 2-oxoglutarate/malate carrier.Elena E Pohl, Kristina Žuna, Tatyana Tyschuk, Jürgen KreiterDepartment of Biomedical Sciences, University of Veterinary Medicine, 1210 Vienna,
Austria

DNA & DNA binding proteins

*26P-080 Dynamic interactions between DNA and a transcription factor, Photozipper, visualized by high-speed atomic force microscopy <u>Akihiro Tsuji</u>, Hayato Yamashita, Osamu Hisatomi, Masayuki Abe Graduate School of Engineering Science, Osaka University, Osaka, Japan

*26P-081 Mechanism of DNAs attraction mediated by low and high valence salts

Hongwei Zuo, Fujia TIAN, Liang DAI city university of Hong Kong

26P-082 Encapsulation of cell nucleolus by single-stranded DNA

Koichiro Maki, Jumpei Fukute, Taiji Adachi

Department of Medicine and Medical Science, Graduate School of Medicine, Kyoto University, Japan/Laboratory of Biomechanics, Institute for Life and Medical Sciences, Kyoto University, Japan/Department of Micro Engineering, Graduate School of Engineering, Kyoto University, Japan/Department of Mammalian Regulatory Network, Graduate School of Biostudies, Kyoto University, Japan

RNA & RNA binding proteins

*26P-083 Molecular simulations to investigate the protein-RNA assembly mechanism of Tetrahymena telomerase <u>Max Cutler</u>, Naoto Hori, Charles Laughton

Nottingham University

26P-084 Structural Ensembles of the 5'-UTR of Hepatitis C Virus RNA With and Without MicroRNA Using SIS-RNA Model Huong T Vu, Naoto Hori

School of Pharmacy, University of Nottingham, Nottingham, NG7 2RD, United Kingdom

DNA/RNA nanotechnology

*26P-085	Automation of DNA gel generation experiments using machine learning and pipetting robots Yuko Yoshida, Masahiro Takinoue
	Department of Computer Science, Tokyo Institute of Technology, Tokyo, Japan
*26P-086	Network formation of enzymes via DNA motif in two types of cascade reactions Aoi Mameuda, Koki Kamiya
	Graduate School of Science and Technology, Gunma University, Gunma, Japan
*26P-087	A Hydrogel Biosensor Combining Aptamer Recognition and DNA- Driven Hydrogels Satofumi Kato, Masahiro Takinoue, Hiroaki Onoe Keio University
*26P-088	Towards Rotary DNA Motor with Conformational Change <u>Akihiro Fukuda</u> , Yusuke Sato, Takeshi Yokoyama, Yoshikazu Tanaka, Shoichi Toyabe Tohoku University
*26 P- 089	Liquid-liquid phase separation of computational DNA droplets on the gold surface Koki Yoshida, Masahiro Takinoue Tokyo Institute of Technology

Nucleic acid: Others

*26P-090 Molecular Dynamics Simulations to Investigate Interactions Between Polymers and RNA in Polymer Nanoparticles

James Aaron Robins, Naoto Hori, Cameron Alexander, Keith Spriggs School of Pharmacy, University of Nottingham

*26P-091 Oligonucleotide Assembly enhanced by intrinsically disordered protein droplet

<u>Taiji Ueno</u>, Yoshihiro Minagawa, Hiroyuki Noji Department of Applied Chemistry, School of Engineering, The university of Tokyo.

Chromatin & Chromosomes

*26P-092	Mechanic Properties of Nucleosomes are Key Modulators of the Unwrapping Energy Landscape
	Maria Julia Maristany, Ignacio Perez Lopez, Stephen Farr, Jan Huertas,
	Rosana Collepardo-Guevara
	Department of Physics, University of Cambridge, UK
*26P-093	Protein search processes mediated by chromatin topology
	<u>Shuvadip Dutta</u> , Adarshkrishnan R., Ranjith Padinhateeri, Mithun K. Mitra
	Department of Physics, Indian Institute of Technology Bombay, Mumbai 400076, India
26P-094	Effect of the molecular crowding environment on the structure of
	poly-nucleosomes
	Tomoko Sunami, Amarjeet Kumar, Hidetoshi Kono
	iQLS, QST
26P-095	From Sequence to Structure: Refining Chromatin Models with PTM
	and Contact Frequency Data
	Justin Chan, Giovanni Brandani, Shoji Takada, Kono Hidetoshi
	Molecular Modeling and Simulation Team, iQLS, QST, Japan

Water & Hydration & Electrolyte

*26P-096 Effect of hydration state polymers on Liquid-Liquid Phase Separation. <u>Kengo Cho</u>, Mafumi Hishida Department of Chemistry, Faculty of Science, Tokyo University of Science, Tokyo, Japan

26P-097 Correlation analysis of co-solvation free energies in insulin dissociation Simon Hikiri, Nobuyuki Matubayasi

College of Life Sciences, Ritsumeikan University, Kusatsu, Japan

Molecular genetics & Gene expression

26P-098 PML protein localization and bioinformatic interactome analysis in ageing related diseases.

<u>Eugene Smirnov</u>, Sergey Silonov, Aleksandra Nozdracheva, Konstantin Turoverov, Alexander Fonin Laboratory of Structural Dynamics, Stability and Folding of Proteins, Russian Academy of Sciences, St. Petersburg 194064, Russia

Morphogenesis and Development

***26P-099** Nuclear softening triggers a transcriptional burst during early embryogenesis

<u>Masahito Tanaka</u>, Rin Sakanoue, Atsushi Takasu, Yasuki Miyagawa, Naoko Watanabe, Kei Miyamoto, Yuta Shimamoto Laboratory of Physics and Cell Biology, National Institute of Genetics, Shizuoka, Japan.

Muscle	
*26 P -100	Cardiac cycle-dependent alterations in redox states revealed by cryo-Raman spectral analysis
	<u>WenJin Ho</u> , Yoshinori Harada, Kentaro Mochizuki, Yasuaki Kumamoto, Masahito Yamanaka, Katsumasa Fujita, Hideo Tanaka
	Department of Pathology and Cell Regulation, Kyoto Prefectural University of Medicine
26P-101	Effects of Near-Infrared Laser Irradiation on Circular Cardiomyocyte Network
	<u>Momo Akada</u> , Kentaro Kito, Masahito Hayashi, Tomoyuki Kaneko LaRC, FB, Grad. Sch. Sci. & Eng., Hosei Univ., Tokyo, Japan

Molecular motor

***26P-102** Directionality on kinesin-1 motility can be determined depending on the anchor points

<u>Rieko Sumiyoshi</u>, Masahiko Yamagishi, Junichiro Yajima Department of Life Sciences Graduate School of Arts and Sciences, The University of Tokyo, Tokyo, Japan/JSPS Research Fellow

*26P-103 Three-dimensional motility of myosin IC bound to lipid membrane <u>Yusei Sato</u>, Kohei Yoshimura, Kyohei Matsuda, Takeshi Haraguchi, Akisato Marumo, Masahiko Yamagishi, Suguru Sato, Kohji Ito, Junichiro Yajima Department of Life Sciences, Graduate School of Arts and Sciences, The University of Tokyo

*26P-104 Energetics of engineered FoF1-ATP synthase with high H+/ATP ratio Kiyoto Yasuda, <u>Riku Marui</u>, Hiroshi Ueno, Hiroyuki Noji The University of Tokyo, Graduate school of Engineering, Department of Bioengineering, Noji Lab.

*26P-105 Biomolecular motors use two asymmetries to generate unidirectional movement

Shintaro Nakayama, Akane Furuta, Maki Yoshio, Misako Amino, Kazuhiro Oiwa, Ken'ya Furuta Advanced ICT Research Institute, National Institute of Information and Communications Technology, Hyogo, Japan/Department of Life Science, Graduate School of Science, University of Hyogo, Hyogo, Japan

26P-107 Nuclear Spin Catalysis in Living Cells and Biomolecular Motors Vitaly K Koltover

Federal Research Center of Problems of Chemical Physics and Medical Chemistry, Russian Academy of Sciences, Chernogolovka, Moscow Region, 142432, Russia

26P-108 Mechanical force measurement of F1-ATPase using accurate revolution control by an optical-vortex tweezers. Yu Hashimoto, Tomoko Otsu-Hyodo, Taro Ando, Yoshiyuk Ohtake,

Sayaka Kazami, Yuji Kimura, Yu Takiguchi, Hiroyasu Itoh Central Research Laboratory, Hamamatsu Photonics K.K., Hamamatsu, Japan

26P-109 Flagellar rotation-speed difference observed in the same bacterial cells

<u>Tsubasa Ishihara</u>, Shuichi Nakamura Grad. Sch. Eng., Tohoku Univ.

Single Molecule Biophysics

*26P-110 Enhanced Interpretation of STED-FCS Diffusion Law Plot Dependencies

Barbora Svobodova, Radek Sachl

Faculty of Mathematics and Physics, Charles University, Prague, Czech Republic/J. Heyrovsky' Institute of Physical Chemistry of the Czech Academy of Sciences, Prague, Czech Republic

*26P-111 High sensitivity detection of HBV RNA based on 3D-DNA nanomachine and biological nanopore sensing technology

Shixin Yan, Jia Geng

Department of Laboratory Medicine, State Key Laboratory of Biotherapy and Cancer Center, Med-X Center for Manufacturing, West China Hospital, Sichuan University, 610041, Chengdu, China

*26P-112 NANOSPACER: Optical analysis of biomolecules and nanoparticles in solution using nanofluidic devices

Oliver Vanderpoorten UiT The Arctic University of Norway, Tromsoe, Norway

*26P-113 Single-molecule manipulation of genome integrity guardians <u>María Ortiz</u>, Roberto Galleto, Borja Ibarra IMDEA Nanociencia, Faraday 9, 28049, Madrid, Spain

*26P-114 Interdomain linkers regulate the mechanotransduction in proteins <u>Tanuja Joshi</u>, Pritam Saha, Sabyasachi Rakshit Department of Chemical Sciences, Indian Institute of Science Education and Research Mohali, Punjab, India-140306,

26P-115 Single-molecule sensing with aerolysin pore-forming toxins Matteo Dal Peraro Institute of Bioengineering, School of Life Sciences, Ecole Polytechnique Fédérale de Lausanne (EPFL), Lausanne, Switzerland

26P-116 Live-cell Single-molecule Imaging and Mapping of Human SWI/SNF Chromatin Remodelers Reveal Bromodomain-mediated and Cancermutants-specific Landscape of Multi-modal DNA-binding Dynamics Wilfried Engl, Hendrik Sielaff, Aliz Kunstar-Thomas, Siyi Chen, Woei Shyuan Ng, Ziqing Winston Zhao National University of Singapore

Cell biology: Adhesion

26P-117 Dynamics and function of adhesion GPCR Rinshi Kasai, Shigetaka Nishiguchi, Takayuki Uchihashi National Cancer Center Research Institute

Cell biology: Motility

*26P-118	Sheet-like structure of bacterial actin MreBs driving helicity switching
	by cryo electron tomography
	<u>Haruka Yuasa,</u> Yuya Sasajima, Hana Kiyama, Daichi Takahashi,
	Takuma Toyonaga, Tomoko Miyata, Fumiaki Makino, Keiichi Namba,
	Makoto Miyata
	Graduate School of Science, Osaka Metropolitan University, Osaka, Japan
*26P-119	Rotation of bacterial cell-bodies in different species capable and
	incapable of flagellar wrapping
	<u>Naoki Ogura,</u> Aoba Yoshioka, Daisuke Nakane, Hirofumi Wada
	Department of Biomedical Engineering, Tokyo University of Agriculture and Technology
*26P-120	Viscoelasticity dependence of ciliary beating and the resulting flow
	<u>Saki Tamura,</u> Misako Otaki, Yoshihiro Murayama
	Department of Biomedical Engineering, Tokyo University of Agriculture and Technology
*26P-121	Water flow navigates the long journey of surface-associated bacteria
	living in hot springs
	<u>Naoki Uemura</u> , Naoya Chiba, Masatada Tamakoshi, Daisuke Nakane
	Department of Engineering Science, The University of Electro-Communications,
	Tokyo, Japan

*26P-122 Analysis of Fluctuations in Measurement Data of Bacterial Flagellar Motors

Kenta Takemori, Yusuke V. Morimoto Graduate School of Computer Science and Systems Engineering, Kyushu Institute of Technology, Fukuoka, Japan

*26P-123 Quantitative model of vascular cell motility in angiogenesis Hayate Segawa, Shigetomo Fukuhara, Kazushi Ikeda, Yuichi Sakumura Nara Institute of Science and Technology

*26P-124 Suppressing Bacterial Surface Colonization and Motility with Biosurfactants

Li Xiaojie, Andrew Utada Grad. Sch. Of Sci. and Tech. Univ. of Tsukuba, Tsukuba, Japan

*26P-125 Cell Type-Dependent Coordinated Regulation of Rho GTPases in Cell Motility Yufei Wu

Nara Institute of Science and Technology

- 26P-126 A highly conserved Arg-391 residue of FlhA is involved in export switching of the flagellar type III secretion system in Salmonella <u>Tohru Minamino</u>, Miki Kinoshita, Motoshi Sakai, Takayuki Uchihashi, Norihiro Takekawa, Katsumi Imada, Keiichi Namba Graduate School of Frontier Biosciences, Osaka University
- 26P-128 Cell size variation affects bacterial swimming speed <u>Riu Osanai</u>, Shuichi Nakamura Dept.appl.Phys.,Grad.Sch.Eng.,Tohoku Univ.

26P-129 Atomic model comparison of the L- and R-type straight bacterial flagellar filaments for understanding the supercoiling mechanism Fumiaki Makino, Kasim Waraichi, Miki Kinoshita, Tomoko Miyata, Tohru Minamino, Keiichi Namba Graduate School of Frontier Biosciences, Osaka University, Suita, Osaka, Japan/JEOL YOKOGUSHI Research Alliance Laboratories, Osaka University, Suita, Osaka, Japan

Cell biology: Cytoskeleton & Membrane skeleton

- *26P-130 Deformability cytometry of Jurkat cells for cell immunotherapy Lija Fajdiga, Nina Bernat, Lara Betocchi, Špela Zemljič Jokhadar, Jure Derganc Institute of Biophysics, Faculty of Medicine, University of Ljubljana, Slovenia
- ***26P-131** Development and application of an optogenetic tool to control the actin polymerization

Kei Yamamoto, Yosuke Yamazaki, Kazuhiro Aoki, Makito Miyazaki RIKEN Center for Biosystems Dynamics Research, Yokohama, Japan

 26P-132
 Polymerization of PEG-attached actin

 Masaya Sagara, Hiroto Narita, Kuniyuki Hatori
 Department of mechanical systems engineering, Faculty of engineering/Graduate school of science and engineering, Yamagata university

26P-133 Measurement of intracellular forces using centrifuge polarizing microscope (CPM)

Makoto Goda, Michael Shribak, Zenki Ikeda, Naobumi Okada, Tomomi Tani, Gohta Goshima, Rudolf Oldenbourg, <u>Akatsuki Kimura</u> Marine Biological Laboratory, Woods Hole, Massachusetts, USA/National Institute of Genetics, Mishima, Japan/Sokendai, Mishima, Japan

26P-134 Construction of a mechanical model for C. elegans gastrulation <u>Tokitaka Katayama</u>, Akatsuki Kimura National Institute of Genetic (NIG)/Genetics Course, SOKENDAI

Cell biology: Signal transduction & Cell membrane

- *26P-135 Cell size feedback mechanism for propagating cell-cell signals in Dictyostelium discoideum Yukihisa Hayashida, Chikoo Oosawa, Takuo Yasunaga, Yusuke V Morimoto Grad. Sch. Comp. Sci. and Sys. Eng., Kyushu Inst. Tech., Fukuoka, Japan
- *26P-136 Cellular Guardianship Symphony by the Dynamic Duo of LL37 and HNP-1

<u>Jing Zhang</u>, Kaori Sugihara The University of Tokyo

26P-137 Slow diffusion and signal amplification on membranes regulated by phospholipase D Gen Honda, Satoshi Sawai, Miho Yanagisawa Department of Basic Science, Graduate School of Arts and Sciences, University of Tokyo, Tokyo, Japan

26P-138 Small extracellular vesicles trigger integrin-mediated adhesion signal in the recipient cells Koichiro M Hirosawa, Yusuke Sato, Eriko Yamaguchi, Naoko Komura, Hiromune Ando, Yasunari Yokota, Kenichi G.N. Suzuki iGCORE, Gifu Univ., Japan

Biological & Artificial membrane: Structure & Property

*26P-139	Membrane tension and its effect on a membrane structure Zuzana Johanovská, David Šťastný, Radek Šachl, Martin Hof Heyrovský Institute of Physical Chemistry, Czech Academy of Sciences, Prague, Czech Republic/Charles University, Faculty of Mathematics and Physics, Prague, Czech Republic
*26P-141	Elucidating the Molecular Mechanism of the Dual Cooperative Effect Between antimicrobials LL37 and HNP1: A Study of Peptide-Lipid and Peptide-Peptide Interactions Yuge Hou, Kaori Sugihara The institute of industrial Science, The University of Tokyo
26P-142	Dimerization of transmembrane peptides synergistically enhances the lipid scrambling activities <u>Hiroyuki Nakao</u> , Toshiki Tsujii, Hiroaki Saito, Keisuke Ikeda, Minoru Nakano Faculty of Pharmaceutical Sciences, University of Toyama, Toyama, Japan
26P-143	Mechanism of action and lipid-mediated synergistic interactions of antimicrobial peptides: New regulatory mechanisms also for membrane proteins? Burkhard Bechinger University of Strasbourg / CNRS, Chemistry UMR7177/Institut Universitaire de France IUF

Poster Sessions

June 26 [Wed]

26P-144 Impact of Acetonitrile Molecules on Miscibility Transition Temperature of Multicomponent Lipid Vesicles Shota Matsuzawa, <u>Kazunari Yoshida</u> Graduate School of Science and Engineering, Yamagata University/Faculty of Engineering, Yamagata University

Biological & Artificial membrane: Dynamics

- *26P-145 Effect of actin encapsulation on the behavior of lipid bilayers under osmotic stress Ken Bessho, Mahito Kikumoto, Yuki Mizutani, Moka Ito, Kingo Takiguchi
- ***26P-146** PORE-FORMING ACTIVITIES OF β -HAIRPIN ANTIMICROBIAL PEPTIDES EVALUATED BY LIPID BILAYER SYSTEM

Nagoya Univ., Grad. Sch. Sci., Dept. Bio. Sci.

Yuki Hagiri, Wakana Hashimoto, Ryuji Kawano Department of Biotechnology and Life Science, Tokyo University of Agriculture and Technology, Tokyo, Japan

- *26P-147 Synthesis of fluorescence-derivative of DHA-containing phospholipids and its behavior in lipid bilayers. Kotaro Shimizu, Masanao Kinoshita, Nobuaki Matsumori Kyushu University
- *26P-148 Morphological change in liposomes that encapsulating F-actins with the adjusted length distribution <u>Yuki Mizutani</u>, Moka Ito, Mahito Kikumoto, Masahito Hayashi, Kingo Takiguchi Nagoya Univ., Grad. Sch. Sci., Dept. Bio Sci.
- 26P-149
 Membrane viscosity of phase-separated ternary GUVs having Lo domains as the basis for lipid rafts

 Julia Tanaka, Yuka Sakuma
 Department of Physics, Tohoku University

Biological & Artificial membrane: Excitation & Channels

26P-150 Development of a high-throughput device for recording channel currents using agarose gel beads <u>Mami Asakura</u>, Wang Shuyan, Minako Hirano, Toru Ide Dept. of Comp. Tech. Soln., Okayama Univ.

Biological & Artificial membrane: Transport & Signal transduction

*26P-151 The fusion site of cell penetrating peptide sequences affects the cytoplasmic transport Akari Miwa, Koki Kamiya Graduate School of Science and Technology, Gunma University, Gunma, Japan

Membraneless Organella, autophage, Liquid-liquid phase separation

*26P-152	Phase separation and phase transition of protein mixture on chemically modified glass surfaces
	Toya Yoshida, Tomohiro Nobeyama, Kentaro Shiraki
	Institute of Pure and Applied Sciences, University of Tsukuba
*26P-153	Quantitative analysis of lipophagy by a small molecule fluorescent reporter
	Siyang Ding, Jesse Rudd-Schmidt, Oana Sanislav, Jinyun Zou, Tze Cin Owyong, Ebony Monson, Karla Helbig, Paul Fisher, Ian Ganley, Kazuhide Shaun Okuda, Ilia Voskoboinik, Yuning Hong Department of Biochemistry and Chemistry, La Trobe Institute for Molecular Science, La Trobe University, Melbourne, VIC, Australia
*26 P-1 54	Quantitative Raman analyses and photo-regulation of nucleic acid- peptide droplets formed by liquid–liquid phase separation <u>Kohei Yokosawa</u> , Shinya Tahara, Shinji Kajimoto, Takakazu Nakabayashi Graduate School of Pharmaceutical Sciences, Tohoku University, Japan
*26P-155	A new method for structural switching of multiphase coacervates based on rational design of charged polypeptides
	<u>Hinano Nakamoto</u> , Hiroshi Kamizawa, Takumi Yamada, Biplab K C,

Teruki Nii, Takeshi Mori, Yoshiki Katayama, Akihiro Kishimura Graduate School of Systems Life Sciences, Kyushu University

26P-156 Tunable Wetting Properties in Multicomponent Protein Condensates <u>Ding Wang</u>, Lei-Han Tang Department of Physics, Hong Kong Baptist University, Hong Kong SAR, China/ Department of Materials Science and Engineering, Southern University of Science and Technology, Shenzhen, China 26P-157 Super enhancer-derived IncRNA CCAT1-L regulates the condensation of human Mediator Junho Lee Korea Advanced Institute of Science and Technology (KAIST)

26P-158 Metastable phase-separated droplet generation and long-time DNA enrichment by laser-induced Soret effect Mika Kobayashi, Yoshihiro Minagawa, Hiroyuki Noji

University of Tokyo/Tokyo University of Agriculture and Technology

Neuroscience & Sensory systems

*26P-159	High-speed AFM reveals activity-dependent stable complexes of kinase domains in CaMKIIβ Keisuke Matsushima, Hideji Murakoshi, Mikihiro Shibata Graduate School of Natural Science and Technology, Kanazawa University, Ishikawa, Japan
*26P-160	Molecular mechanism of classical conditioning in earthworm Sukehiro Kabayama, Yoshiichiro Kitamura Kanto-Gakuin University
26P-161	Ion-channel-based complete synchronization between neurons Seido Nagano Department of Bioinformatics, Ritsumeikan University
26P-162	Models of complex structure-related diffusion anomalies of transport in the brain's extracellular space Eugene B Postnikov, Igor Sokolov, Anastasia Lavrova, Dmitry Postnov Kursk State University

Neuronal circuit & Information processing

- *26P-163 Outgrowth order in breaking symmetry of immature neurites is another regulation factor of neuronal polarity formation <u>Soya Hagiwara</u>, Ryohei Yamazaki, Nanami Abe, Naoya Takada, Kenji Yasuda Dept. Phys., Sch. Adv. Sci. & Eng., Waseda Univ., Japan
- 26P-164 Large-scale voltage-sensitive dye imaging of mouse prefrontal cortex: Biophysical mapping of intra- and inter-hemispheric connections <u>Takashi Tominaga</u>, Pooja Gusain, Makiko Taketoshi, Yoko Tominaga Institute of Neuroscience, Tokushima Bunri University, Sanuki, Japan/Kagawa School of Pharmaceutical Sciences, Tokushima Bunri University, Sanuki, Japan

Behavior

*26P-165 Swimming ciliate, Stentor selects anchoring sites accompanied by extracellular geometries Syun Echigoya, Katsuhiko Sato, Toshiyuki Nakagaki, Yukinori Nishigami RIES Hokkaido University, Sapporo, Japan

Photobiology: Vision & Photoreception

- *26P-166 Electrophysiological Study of the Effect of Weak Organic Acids on the Transport Activity of Proton Pumping Rhodopsin of Rhizobacteria Zikun Lyu, Shunki Takaramoto, María del Carmen Marín, Hiromu Yawo, Keiichi Inoue The Institute for Solid State Physics, University of Tokyo, Kashiwa, Japan
- *26P-167 Structural basis for early proton transfer reaction on a primate bluesensitive pigment

<u>Yosuke Mizuno</u>, Hiroo Imai, Hideki Kandori, Kota Katayama Graduate school of Engineering, Nagoya Institute of Technology, Aichi, Japan

*26P-168	Time-resolved crystallography uncovering cryptochrome signal transduction mechanism Yuhei Hosokawa, Po-Hsun Wang, Mai Nakamura, Nicolas Caramello, Sylvain Engilberge, Antoine Royant, Lars-Oliver Essen, Ming-Daw Tsai, Junpei Yamamoto, Manuel Maestre-Reyna National Taiwan University, Taipei, Taiwan/Academia Sinica, Taipei, Taiwan/Osaka University, Osaka, Japan
*26P-169	Light-induced difference FTIR analysis of xenorhodopsin from Nanosalina at 77 K Yuma Ito, Tatsuro Nishikino, Hideki Kandori, Yuji Furutani Department of Life Science and Applied Chemistry, Nagoya Institute of Technology
26P-170	Spectroscopic and functional characterization of novel viral rhodopsins <u>Takashi Nagata</u> , Shunki Takaramoto, Andrey Rozenberg, Oded Béjà, Keiichi Inoue The Institute for Solid State Physics, The University of Tokyo, Japan
26P-171	Modulation of intracellular calcium responses using photocyclic vertebrate visual pigments <u>Kazumi Sakai</u> , Shion Aoki, Takahiro Yamashita Kyoto University

Photobiology: Photosynthesis

*26P-172	Carotenoids binding effect of the photoreaction processes on Xanthorhodopsin
	<u>Shota Itakura</u> , Yosuke Mizuno, Kota Katayama, Rei Abe-Yoshizumi,
	Ariel Chazan, Oded Béjà, Hideki Kandori
	Graduate School of Engineering, Nagoya Institute of Technology, Aichi, Japan
*26P-173	Mobility of protein complexes in plant thylakoid membranes analy

 *26P-173
 Mobility of protein complexes in plant thylakoid membranes analyzed by high-speed atomic force microscopy

 Yudai Nishitani, Daisuke Yamamoto

 Fac. Sci., Fukuoka Univ., Fukuoka, Japan

*26P-174	Excitation energy transfer dynamics among antenna pigments in the ΔpshX-reaction center from Heliomicrobium modesticaldum <u>Risa Kojima</u> , Kevin E Redding, Daisuke Kosumi, Hirozo Oh-oka College of Life Science, Ritsumeikan University, Shiga, Japan
26P-175	Structural Basis for Enabling Photosynthesis with Extremely Low- energy Near-infrared Light in the LH1–RC Complex of a Thermophilic Purple Nonsulfur Bacterium Yukihiro Kimura, Ryo Kanno, Kaisei Mori, Ryuta Seto, Yoshiki Matsuda, Shinji Takenaka, Hiroyuki Mino, Malgorzata Hall, Endang R. Purba, Akira Mizoguchi, Bruno M. Humbel, Michael T. Madigan, Zheng-Yu Wang-Otomo, Kazutoshi Tani Graduate School of Agriculture, Kobe University, Nada, Kobe, Japan
26P-176	Characterization of an LH1–RC photocomplex from a novel Japanese hot spring purple sulfur bacterium, Caldichromatium japonicum Akane Minamino, Mohit. K. Saini, Endang R. Purba, Malgorzata Hall, Shinji Takenaka, Vera Thiel, Bruno M. Humbel, Michael T. Madigan, Zheng-Yu Wang-Otomo, Kazutoshi Tani, Yukihiro Kimura Department of Agrobioscience, Graduate School of Agriculture, Kobe University, Nada, Kobe, Japan
Photobiology	r: Optogenetics & Optical control
*26P-177	Channel gating mechanism of K+ selective channelrhodopsin, KCR <u>Ryotaro Shimamura</u> , Shoko Hososhima, Hideki Kandori, Satoshi Tsunoda Department of Life Science and Applied Chemistry, Nagoya Institute of Technology
*26P-178	In vivo single-cell 3D optogenetics technology with light-field

microscopy

<u>Tomoyoshi Inoue</u>, Ryuki Imamura, Naoya Kataoka, Akihiro Fukushima, Shin Usuki, Takuma Sugi Program of Biomedical Science, Graduate School of Integrated Sciences for Life, Hiroshima University, Hiroshima, Japan

26P-179 Identification of the important region for photoactivity in photoactivated adenylyl cyclase Minako Hirano, Masumi Takebe, Hinase Kondo, Mami Asakura, Toru Ide Grad. Sch. Health Sys., Okayama University

Radiobiology & Active oxygen

26P-180 Verification of the dosimetry techniques using GAFCHROMIC films for the study of the mechanism of the FLASH effect with synchrotron radiation

Munetoshi Maeda, Ryoichi Hirayama, Ayumi Shiro, Mika Maeda, Masanori Tomita The Wakasa Wan Energy Research Center, Fukui, Japan

Origin of life & Evolution

*26P-181 Observation of Fatty Acid Vesicle Condensation on the Surface of Simulated Hydrothermal Vent Minerals

Shunto Harada, Hye-Eun Lee, Yamei Li, Ryuhei Nakamura Earth-Life Science Institute (ELSI), Tokyo Institute of Technology/RIKEN Center for Sustainable Resource Science

***26P-182** Genome reduction increases parasite sensitivity and promotes the evolution of endosymbiotic mutualism

Yuki Kanai, Chikara Furusawa Department of Biological Sciences, Graduate School of Science, The University of Tokyo

*26P-183 Proofreading inherited by template-directed ligation <u>Hiroyuki Aoyanagi</u>, Yasuhiro Magi, Shoichi Toyabe Department of Applied Physics, Graduate School of Engineering, Tohoku University, Japan

26P-184 The Relation Between Biology and Physics: Origins of Life Research and its Philosophical Implications

Julieta Macome Cambridge University

Synthetic biology & Artificial cells

***26P-185** Control of lipid membrane composition in vesicles by external stimulations

Sumin Lee, Koki Kamiya Graduate School of Science and Engineering, Gunma University, Gunma, Japan

*26P-186 Construction of enzymatically assembled gel-in-liposome as artificial Cell model Wancheng Zhang, Aileen Cooney, Lorenzo Di Michele, Yuval Elani,

<u>Wancheng Zhang</u>, Alleen Cooney, Lorenzo Di Michele, Yuvai Ela Tomoaki Matsuura Earth-life Science Institute, Tokyo Institute of Technology

*26P-187 Creation of cell-sized droplets entrapping DNA or living cells through phase separation under one-dimensional confinement Mayu Shono Department of Chemical Engineering and Materials Science, Doshisha University,

*26P-188 TOWARD THE CONSTRUCTION OF A SELF-REPLICATING SYSTEM DRIVEN BY GENE EXPRESSION OF REPLICATION CYCLE REACTION (RCR)

Yuya Yamahishi, Sonoyama Naoki, Kawakami Naoki, Hasebe Tomonori, Su'etsugu Masayuki College of Science, Rikkyo University, Tokyo, Japan

26P-189 Analysis of biochemical reaction in liposomes after terahertz wave irradiation

<u>Gakushi Tsuji</u>, Masaya Oki, Yuusuke Yamaguchi Department of Applied Chemistry and Biotechnology, Graduate School of Engineering, University of Fukui/Life Science innovation center, University of Fukui, Fukui, Japan

26P-190 Cation-selective pores from POSS-decorated amphiphilic diblock copolymers

<u>Denis G Knyazev</u>, Nora Hagleitner-Ertuğrul, Asad Ullah, Nikolaus Goessweiner-Mohr, Hazrat Hussain, Peter Pohl Institute of Biophysics, Johannes Kepler University Linz, Gruberstr. 40, 4020, Linz, Austria

Computational biology: Bioinformatics

Kyoto, Japan

- *26P-191 Learning What AlphaFold2 Learned
 - Nozaki Kosei

Department of Applied Physics, Graduate School of Engineering, Nagoya University.

- *26P-192 Theoretical Insights into Conformational Changes in SOD1 Involved in the Pathological Mechanism of Amyotrophic Lateral Sclerosis: Residue Interaction Network Analysis Shun Fujimaki, Norifumi Yamamoto Chiba Tech
- 26P-193 Analysis of protein-protein interaction search space by ensemble docking <u>Nobuyuki Uchikoga</u>, Yuri Matsuzaki Dept Network Design, Sch Interdiscip Math Sci, Meiji Univ.

Computational biology: Molecular simulation

*26P-194	Experimental and Computational Predictions of the Intrinsic Reactivity of Small Molecules with Lipid Membranes Yeshurun Amarasingham Tanna, John Sanderson, Mark Wilson Department of Chemistry, University of Durham (UK)
*26P-195	Multiphase condensate formation of postsynaptic density: a comparative study of protein assembly in 3D and 2D systems <u>Risa Yamada</u> , Shoji Takada Department of Biophysics, Graduate School of Science, Kyoto University, Kyoto, Japan
*26P-196	Molecular simulations of TMEM16A channel blockers Tanadet Pipatpolkai School of Physical and Mathematical Science, Department of Physics and Applied Physics, Nanyang Technological University, Singapore, 637371
*26P-197	Different behavior of dissociation pathways of glutamine and glutamate in complex with Medaka Taste Receptor T1R Duy Phuoc Tran, <u>Vi Toan Lam</u> , Hao Thai Nguyen, Akio Kitao, Atsuko Yamashita Life Science & Technology, Tokyo Institute of Technology, Japan
*26P-198	Decoding Peptide Solvation Dynamics: Uncovering the Influence of Hydrophobic Forces and Addressing Solvation Complexity via Multiway Statistical Analysis Techniques. Monika Phougat, Narinder Singh Sahni, Devapriya Choudhury School of Computational & Integrative Sciences, Jawaharlal Nehru University, New Delhi, India

*26P-199	Ubiquinone/ubiquinol exchange pathway in the photosynthetic RC– LH1 supercompex Yosuke Teshirogi, Yoshitaka Moriwaki, Tohru Terada Dept. of Biotechnol., Grad. Sch. of Agri and Life Science., The Univ. of Tokyo
*26P-200	Spectrum analyses on the non-linear response of a red blood cell model <u>Tetsuya Yamamoto</u> , Hiroshi Watanabe Keio University
*26P-201	Lipid-protein interaction fingerprints for the Kv7 ion channels Ali Saad Kusay, Lucie Delemotte, Sara Liin Division of Neurobiology, Department of Biomedical and Clinical Sciences, Linköping University, Linköping, Sweden
*26P-202	A drug design strategy based on in vitro and in silico studies applied to the development of inhibitors against alpha-glucosidase and alpha- amylase receptors of diabetics from selected Metformin derivatives. Nor Akmalyati Sulong Department of Chemistry, Quantum Information Science and Technology (QIST), Faculty of Science, University Malaya, 50603 Kuala Lumpur, Malaysia /Institute for Molecular Science, Myodaiji, Okazaki 444-8787, Japan
*26P-203	Molecular Dynamics Study on the Solvent Influence and Stability of the Aspirin-Hydroxypropyl-β-Cyclodextrin Complex Structure Helmia Jayyinunnisya, Dedy Rendrawan, Lince Meriko, Kazutomo Kawaguchi, Hidemi Nagao Graduate School of Mathematical and Physical Science, Kanazawa University, Kanazawa, Japan
*26P-204	Molecular dynamics calculations of peptides self-assembling on boron nitride surfaces <u>Hiroki Maeda</u> , Chishu Homma, Eiji Yamamoto, Yuhei Hayamizu Tokyo Tech
*26P-205	Docking and Molecular Dynamics Simulation Study of BAK1 and BRI1 Proteins in Arabidopsis thaliana Plant Lince - Meriko, Helmia Jayyinunnisya, Dedy Rendrawan, Kazutomo Kawaguchi, Hidemi Nagao Kanazawa University

26P-206	Hydrogen Bond Energies in Helical Secondary Structures Dissected by Negative Fragmentation Approach and Density Functional Theory <u>Hiroko X Kondo</u> , Haruki Nakamura, Yu Takano Kitami Institute of Technology
26P-207	Insight into structural propagating mechanism of photoactivated adenylate cyclase OaPAC by microsecond molecular simulation. <u>Akiya Fukuda</u> , Masahiko Taguchi, Shun Sakuraba, Justin Chan, Eriko Nango, Hidetoshi Kono Graduate School of Science, Tohoku University
26P-208	Integration of AlphaFold with Molecular Dynamics for Sampling Conformational States of Transporter Proteins Jun Ohnuki, Kei-ichi Okazaki Institute for Molecular Science, National Institutes of Natural Sciences, Okazaki, Japan
26P-209	Molecular simulations of cholesterol recognition by SREBP cleavage- activating protein Charal Khiewdee, Puey Ounjai, Tanadet Pipatpolkai Department of Biology, Faculty of Science, Mahidol University, Bangkok, Thailand 10400
26P-210	 Development and Application of a Protocol for Predicting Membrane Permeability of Cyclic Peptides Based on Molecular Dynamics Simulations <u>Masatake Sugita</u>, Takuya Fujie, Keisuke Yanagisawa, Masahito Ohue, Yutaka Akiyama Department of Computer Science, School of Computing, Tokyo Institute of Technology, W3-34, 2-12-1, Ookayama, Meguro-ku, Tokyo, Japan/Middle Molecule IT-based Drug Discovery Laboratory (MIDL), Tokyo Institute of Technology, W8-76, 2-12-1, Ookayama, Meguro-ku, Tokyo, Japan
26P-211	Impact of glycan shielding on antibody epitopes on viral envelope proteins revealed by molecular dynamics simulations Suyong Re, Kenji Mizuguchi Artificial Intelligence Center for Health and Biomedical Research, National Institutes of Biomedical Innovation, Health, and Nutrition, Osaka, Japan

26P-212	Hybrid of Manifold Learning and Molecular Simulation to Reconstruct
	the Protein Conformational Change using Cryo-Electron Microscopy
	Experiment

Takashi Yoshidome Department of Applied Physics, Graduate School of Engineering, Tohoku University, Japan

26P-213 Elucidating the Importance of Water Models in Protein-ATP Interactions in High ATP Concentrations Toshifumi Mori, Norio Yoshida

> Institute for Materials Chemistry and Engineering, Kyushu University/Interdisciplinary Graduate School of Engineering Sciences, Kyushu University

- 26P-214 Fast Computational Method for the Hydration Free Energy Taichi Ito, Takashi Yoshidome Department of Applied Physics, Graduate School of Engineering, Tohoku University, Japan
- 26P-215 Molecular dynamics method for studying a flow on lipid bilayer <u>Masaki Otawa</u>, Satoru G. Itoh, Hisashi Okumura The Graduate University for Advanced Studies/Institute for Molecular Science

Computational biology: Biological modeling and simulation

*26P-216 Metabolome Analysis in Mice Liver Using Thermodynamics <u>Takumi Abekawa</u>, Satoshi Ohno, Shinya Kuroda Department of Biological Sciences, School of Science, University of Tokyo, Tokyo, Japan

*26P-217 INVESTIGATION OF FACTORS AFFECTING THE QUALITY OFSINGLE PHOTON EMISSION COMPUTED TOMOGRAPHY (SPECT) IMAGES: A MONTE CARLO SIMULATION STUDY

> Mercy Amondi, Jared Ombiro, Sankara Aluko, Duke oeba School of Pure, Applied and Health Sciences, Maasai Mara University. P.O. Box 861-20500, Narok, Kenya. mercyamondi161@gmail.com

*26P-218 Elucidating the adaptive mechanical behaviors of actomyosin bundles in cells

<u>Shihang Ding</u>, Taeyoon Kim, Shinji Deguchi Graduate School of Engineering Science, Osaka University, Osaka, Japan

*26P-219	Development of a Protein Language Model-Based Thermal Stability Prediction Model for Nanobodies
	Yuki Hashidate, Yasuhiro Matsunaga
	Graduate School of Science and Engineering, Saitama University, Saitama, Japan
*26P-221	Prediction of the mutation effects of Indonesian Protein Nsp3 SARS- Unique domain (SUD)-pyridostatin as a ligand interaction via end- point binding free energy calculations Dedy Rendrawan, Kazutomo Kawaguchi, Hidemi Nagao Faculty of Mathematics and Physics, Institute of Science and Engineering, Kanazawa University, Japan,
*26P-222	Adhesive Active Brownian Particle Model for Cell Populations
	<u>Sota Shimamura,</u> Nen Saito, Shuji Ishihara
	Graduate School of Arts and Sciences, The University of Tokyo, Tokyo, Japan
26P-223	Estimation of biochemical reaction parameters using a mathematical model of the cell-free translation system Shunnosuke Ban, Yusuke Himeoka, Tomoaki Matsuura, Yoshihiro Shimizu, Chikara Furusawa
	Department of Physics, Graduate School of Science, The University of Tokyo, Tokyo, Japan
26P-224	Coevolution of functionality and foldability of lattice proteins Norifumi Maruyama, <u>Macoto Kikuchi</u> Cybermedia center, Osaka University/Department of Physics, Osaka University
26P-225	Live imaging-based inference of mechanical potential of cell-cell interaction in 3D-multicellular systems <u>Hiroshi Koyama</u> , Toshihiko Fujimori Div. Embryology, National Institute for Basic Biology, Japan/SOKENDAI (Grad. Univ. Advanced Studies)
26P-226	Visualization of protein conformational ensembles using refinement in the information content space: Application to SANXS data <u>Tomotaka Oroguchi</u> , Rintaro Inoue, Ken Morishima, Masaaki Sugiyama Department of Physics, Faculty of Science and Technology, Keio University/RIKEN SPring-8 Center

Poster Sessions

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Computational biology: machine learning for molecules or cell systems

*26P-227 Examining Cell Division Dynamics in Cyanidioschyzon merolae Through High-Resolution 3D Imaging Jin Kousaka, Atsuko H. Iwane, Yuichi Togashi

Ritsumeikan University

- *26P-228 Structure changes in the ABC transporter cycle using nonlinear morphing method, MOVE-DM. Shota Shimogochi, Kazuhi Harai, Ryota Kiyooka, Naoyuki Miyashita Grad. Sch. BOST, KINDAI Univ.
- 26P-229 Predicting Novel PLP-binding Proteins Using Transfer Learning of Graph Neural Network-based Ligand-binding Site Prediction Masafumi Shionyu, Momoka Nakamoto, Atsushi Hijikata, Takashi Nakamura, Yukio Mukai Graduate School of Bioscience, Nagahama Institute of Bio-Science and Technology, Shiga, Japan

Mathematical & Theoretical biology

*26P-230	Population dynamics of generalist/specialist strategies in the feast- famine cycle Rintaro Niimi, Chikara Furusawa, Yusuke Himeoka
	Graduate School of Science, the University of Tokyo
*26P-231	Mathematically deriving loop mobility for single protein structures Virginia Apostolopoulou, Nicholas Pearce, Helen Ginn Hamburg Centre for Ultrafast Imaging, Universität Hamburg, Luruper Chaussee 149, DE-22761, Hamburg, Germany
*26P-232	Appearance and adaptive properties of bow-tie structures from simple metabolic networks Yudai lyoda, Chikara Furusawa, Yusuke Himeoka Graduate School of Science, The University of Tokyo, Japan
26P-233	Reinforcement learning is a common principle for biological control of complex adaptive systems Tomoyuki Yamaguchi Research Institute, Nozaki Tokushukai Hospital, Osaka, Japan

26P-234 Evaluation of different machine learning models applied to diagnostics and treatment success at the pulmonary tuberculosis <u>Anastasia Lavrova</u>, Diljara Esmedlyaeva, Artem Veselsky, Pavel Gavrilov, Eugene Postnikov Saint-Petersburg State Research Institute of Phthisiopulmonology, Russia

Ecology & Environment

*26P-235 Analysis of Alcanivorax borkumensis Biofilm on Binary Oil-mixtures using Microfluidic Devices Rei Shimizu Graduate School of Sci. and Tech., Univ. of Tsukuba, Tsukuba, Japan

Nonequilibrium state & Biological rhythm

26P-236 Spiral wave and homogeneous cyclic modes on membrane <u>Hiroshi Noguchi</u>, Frédéric van Wijland, Jean-Baptiste Fournier University of Tokyo

Measurements

- *26P-237 Hydrogel Colloidosomes from Aqueous Two-Phase Emulsion as New Microreactors for Digital Nucleic Acid Detection Bicheng Zhang, Kanji Tomohara, Hiroyuki Noji Graduate School of Engineering, The University of Tokyo
- *26P-238 Development for the detection of salivary biomarkers by solid-state nanopore

<u>Eiji Kato</u>, Ryo Akita, Sotaro Uemura Department of Biological Sciences, Graduate School of Science, The University of Tokyo, Tokyo, Japan

26P-239 3D structural determination of biological ensembles using high-order spatial correlations in single-particle X-ray scattering Wenyang Zhao, Osamu Miyashita, Miki Nakano, Florence Tama RIKEN Center for Computational Science, Kobe, Japan

- 26P-240IR measurement of flavin mononucleotide aqueous solution by
resonance IR method
Minori Yamakawa, Hirona Takahashi, Konoka Mifune, Makoto Sakai
Okayama University of Science
- 26P-241 Selective IR measurement of fluorescent protein chromophores by resonance IR spectroscopy Konoka Mifune, Hirona Takahashi, Minori Yamakawa, Makoto Sakai Okayama University of Science

Bioimaging

*26P-242	Molecular mechanisms of selective binding of extracellular vesicles to cells as revealed by single particle tracking and super resolution microscopy <u>Tatsuki Isogai</u> , Koichiro M. Hirosawa, Miki Kanno, Yasunari Yokota, Kenichi G. N. Suzuki UGSAS, Gifu Univ. Gifu, Japan
*26P-243	Development of Fluorescent Thermometers Based on Carbon Quantum Dots with Various Detection Modes Yuki S. Kato, Shingo Sotoma, Yukiho Shimazaki, Shunsuke Chuma, Kohki Okabe, Madoka Suzuki, Yoshie Harada Department of Biological Sciences, School of Science, Osaka University, Osaka, Japan
*26P-244	Nanoendoscopy-AFM Measurements of Live Cells: Impact on Proliferation and Stress Response Mohammad Mubarak Hosain, Takehiko Ichikawa, Takeshi Fukuma Nano Life Science, Kanazawa University
*26P-245	Miniaturization of the laser spot for cantilever deflection detection to realize ultra-high-speed AFM <u>Karen Kamoshita</u> , Kenichi Umeda, Noriyuki Kodera Grad. Sch. Math. & Phys., Kanazawa Univ.

*26P-246 Live-cell imaging defines a threshold in CDK activity at the G2/M transition Hironori Sugiyama, Yuhei Goto, Yohei Kondo, Damien Coudreuse, Kazuhiro Aoki The Exploratory Research Center on Life and Living Systems *26P-247 Elucidation of various IL-1 α/β release control mechanisms in cellular inflammatory responses by live-cell imaging of secretion activity. Yu Peng, Zhuohao Yang, Masahiro Nagata Laboratory for Nanobiology, Institute for Protein Research, Osaka University *26P-248 Cryogenic X-ray Ptychographic Imaging of Cltured Cells toward Visualization of Hierarchical Structures of Chromatin Yuta Kinami, Kurumi Nishimagi, Kosei Harada, Masayoshi Nakasako, Yukako Oma, Masahiko Harata, Yuki Takayama Graduate School of Agricultural Science, Tohoku University, Sendai, Japan/RIKEN SPring-8 Center, Hyogo, Japan *26P-249 X-ray diffraction imaging tomography at cryogenic temperature is powerful to reveal 3D structures of biological specimens Kosei Harada, Yuki Takayama, Masayoshi Nakasako Faculty of Science and Technology, Keio University, Yokohama, Japan/RIKEN SPring-8 Center, Hyogo, Japan *26P-250 Development of nano-endoscopic high-speed atomic force microscopy Hikaru Ichida, Kenichi Umeda, Mohammad Shahidul Alam, Risa Omura, Kudo Makiko, Takehiko Ichikawa, Takeshi Fukuma, Takahiro Nakayama, Mikihiro Shibata, Norivuki Kodera Grad. Sch. NanoLS., Kanazawa Univ. 26P-251 CTCF-mediated Chromatin Looping is Coupled to the Formation of Phase-separated Transcriptional Condensate Ryangguen Lee, Moo-Koo Kang, Yong-Jin Kim, Bobae Yang, Hwanyong Shim Department of Biological Sciences, Korea Advanced Institute of Science and Technology (KAIST)

26P-252 Mechanical properties of hiPSC derived-cardiomyocytes: investigating beating and non-beating cell's nucleus via Nanoendoscopy-AFM Alexis S. Borowiak, T. Ichikawa, M. Ito, T. Shimi, Y. Kono, R. Nitta, T. Fukuma

Nano Life Science Institute, Kanazawa University, Kakuma-machi, Kanazawa 920-1192, Japan

26P-253 New mode of intercellular communication: direct vesicle delivery to neighboring cells

<u>Tomohiro Minakawa</u>, Fumiyoshi Ishidate, Takahiro K. Fujiwara, Jun K. Yamashita Department of Cellular and Tissue Communication, Graduate School of Medicine, The University of Tokyo, Tokyo, Japan

26P-254 Photonic chips: a versatile platform for super-resolution microscopy Jean-Claude Tinguely, Vishesh Dubey, Luis Enrique Villegas-Hernandéz, Krishna Agarwal, Balpreet Singh Ahluwalia Uit The Arctic University of Norway

Bioengineering

*26P-255 Direct and continuous monitoring of multi-component antibiotic gentamicin in blood

<u>Chen Chen</u>, Changjian Zhao, Yu Wang, Jia Geng Department of Laboratory Medicine, State Key Laboratory of Biotherapy and Cancer Center, West China Hospital, Sichuan University and Collaborative Innovation Center, Chengdu 610041 China./Tianfu Jincheng Laboratory, City of Future Medicine, Chengdu 610500, China.

*26P-256 Microalgae- and Cyanobacteria-Mediated Fabrication of Functionalized Gold Nanoparticles for Photothermal Applications Reham Samir Hamida, Shingo Sotoma, Madoka Suzuki, Yoshie Harada

Institute for Protein Research, Osaka University, Osaka, Japan

*26P-257 Iono-chromic control of G-protein Ras by fused with M13 peptide Zhang Ziyun, Sabek Yassine, Nobuyuki Nishibe, Shinsaku Maruta Department of Biosciences, Graduate School of Science and Engineering Soka University, Hachioji, Tokyo JAPAN

*26P-258 Redox-Active Liquid-Liquid Phase Separation Materials Promote Oxidative Protein Folding <u>Yukino Yamashita</u>, Munehiro Kumashiro, Tomohide Saio, Takahiro Muraoka Department of Applied Chemistry, Graduate School of Engineering, Tokyo University of Agriculture and Technology 26P-259 SeeDB-Live: minimally invasive optical clearing media for

fluorescence imaging of live tissue ex vivo and in vivo Shigenori Inagaki, Nao Tamagawa, Yuki Kambe, Rei Yagasaki, Aki Teranishi, Misato Miyagawa, Hikari Takeshima, Shunki Tamura, Satoshi Fujimoto, Yuki Naito, Keisuke Ito, Hideki Enomoto, Katsuhiko Hayashi, Takashi Sato, Yoshiaki Tagawa, Satoru Okuda, Tatsuo Sato, Takeshi Imai Kyushu University

Crystal growth & Crystallization technique

*26P-260	Emergence of order from proteins under nucleation
	Dimitrios Triantafyllidis, Suraksha Smitha, Felix Lehmkühler,
	Arwen R. Pearson
	Hamburg Centre for Ultrafast Imaging, Institute for Nanostructure and Solid State
	Physics, Universität Hamburg, Luruper Chaussee 149, 22761 Hamburg, Germany

Virus structure, function, SARS-CoV-2

*26P-261	Complementation of Influenza A virus genome segments by cellular coinfection
	<u>Yuu Kawahara,</u> Hiroyuki Noji, Kazuhito Tabata
	Department of Applied Chemistry, School of Engineering, University of Tokyo
*26P-262	Observation of the compaction process ribonucleoprotein complex
	formed by SARS-CoV-2 genome RNA and N protein by using
	fluorescence correlation spectroscopy.
	<u>Takuya Katayama,</u> Yuji Itoh, Naoya Kaneda, Satoshi Takahashi
	IMRAM, Tohoku Univ./Grad. Sch. Life Sci., Tohoku Univ.
26P-263	Pseudo-luciferase activity of the SARS-CoV-2 spike protein
	<u>Ryo Nishihara,</u> Hisham M Dokainish, Yoshiki Kihara, Yuji Sugita, Ryoji Kurita
	Health and Medical Research Institute, National Institute of Advanced Industrial
	Science and Technology/Japan Science and Technology Agency, PRESTO

Mechanosensing and Mechanobiology, Biological Temperature

*26P-264 Tracking of adipogenesis process of mesenchymal stem cells cultured in the spherical microwells <u>You-Hsuan Liu</u>, Karen G. Rosal, Chon-In Cheong, Feng-Chiao Tsai, Keng-Hui Lin

Institute of Physics, Academia Sinica, Taipei, Taiwan

- *26P-265 Phenotypic Heterogeneity and Cell Orientation in Extracellular Matrix Production of Escherichia coli <u>Fumiaki Yokoyama</u>, Kazumasa Takeuchi Department of Physics, The University of Tokyo, Tokyo, Japan
- 26P-266 "DIY Statistical Mechanics" to understand the concepts of Boltzmann distribution and local temperature Kiyoshi Ohnuma, Masayo Inoue, Noritaka Masaki, Masako Ohtaki, Taro Toyota Nagaoka University of Technology

Biophysics of disease

*26P-267	Tissue Repair in Colorectal Cancer Organoids Jimmy Van Hear, Yanlan Mao, Chris Tape Laboratory for Molecular Cell Biology, University College London
*26P-268	Clustering of Progerin Induces Nuclear Deformation By Disrupting Heterochromatin Organization YigJi Lee, Won-Ki Cho Department of Biological Sciences, Korea Advanced Institute of Science and Technology, Daejeon, 34141, Republic of Korea
26P-269	Fibrinogen-erythrocyte binding and erythrocyte-erythrocyte adhesion as determinants for cardiovascular risk Catarina Lopes, Ryan Gouveia e Melo, Luís Mendes Pedro, Filomena Carvalho, <u>Nuno C. Santos</u> Instituto de Medicina Molecular, Faculdade de Medicina, Universidade de Lisboa, Lisbon, Portugal

Miscellaneous topics

*26P-270 Immobilization of Liposomes within Porous Aluminum Oxide and Intact Release: Basic Study and Applications to Single Particle Analysis for Exosomes

Masahiro Okada, Yusuke Sato, Tetsuji itoh, Seiichi Nishizawa Graduate School of Science, Tohoku University

*26P-271 Enhancing Tomato Shelf Life through Nanoparticle-Based Preservation Techniques

Kummari Swathi, CH.Shanthi Devi Research Scholar, Andhra University, Visakhapatnam/Assistant Professor, St. Francis College for Women, Hyderabad

26P-272 NOVEL ANTIBACTERIAL AGENTS IN TREATING MULTIDRUG RESISTANT BACTERIA CAUSING WOUND INFECTIONS IN DIABETIC PATIENTS

<u>Shailaja Raj Marla</u>, Mithali Raj Marla, Maria Shajan St. Francis College For Women