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

*25P-001	The molecular strucutre of an axle-less F1-ATPase
	Emily Furlong, Ian Reiniger-Chatzigian, Yi Zeng, Simon Brown,
	Meghna Sobti, Alastair Stewart
	Molecular, Structural and Computational Biology Division, The Victor Chang Cardiac Research Institute, Darlinghurst, Australia/Division of Biomedical Science and Biochemistry, Research School of Biology, Australian National University, Acton, ACT, Australia
*25P-002	The role of charges in the enzymatic mechanism of acetoacetate
	decarboxylase
	Masato Ishizaka, Sören Rindfleisch, Florian Auer, Lukas Gingeleit,
	Tat Cheng, Michael Bielecki, Fabian Rabe von Pappenheim, Elke Penka, Ronald Kluger, Eri Sakata, Kai Tittmann
	Department of Molecular Enzymology, Georg-August University Göttingen, Göttingen, Germany./Max-Planck-Institute for Multidisciplinary Sciences, Göttingen, Germany.
*25P-003	Structural insights into the allosteric inhibition of P2X4 receptors
	<u>Cheng Shen</u> , Yuqing Zhang, Wenwen Cui, Yimeng Zhao, Danqi Sheng,
	Xinyu Teng, Miaoqing Shao, Muneyoshi Ichikawa, Jin Wang,
	Motoyuki Hattori
	Fudan University, Shanghai, China

June 25 [Tue]

Poster Sessions

*25P-004 Cryo-EM Structure of P-glycoprotein Bound by Three Elacridar P-gp-Inhibitor Molecules

<u>Norie Hamaguchi</u>, Naruhiko Adachi, Toshio Moriya, Masato Kawasaki, Satoshi Yasuda, Naohiko Anzai, Toshiya Senda, Satoshi Ogasawara, Takeshi Murata Graduate School of Medical and Pharmaceutical Sciences, Chiba University/Graduate School of Medicine, Chiba University/Graduate School of Science, Chiba University

*25P-005 Structural insights into the orthosteric inhibition of P2X receptors by classical non-ATP-analog antagonists

Danqi Sheng, Chenxi Yue Fudan University, Shanghai China

*25P-006 Ion selectivity mechanism of the MgtE channel for Mg2+ over Ca2+ Xinyu Teng, Danqi Sheng, Ye Yu, Jin Wang, Motoyuki Hattori Fudan University, Shanghai, China

*25P-007 Tracking the glucose/ xylose isomerase mechanism using freezing under high pressure

Agnieszka Klonecka, Joanna Slawek, Philippe Carpentier, Christoph Mueller-Dieckmann, Katarzyna Kurpiewska, Maciej Kozak SOLARIS National Synchrotron Radiation Centre, Kraków, Poland/Faculty of Physics, Astronomy and Applied Computer Science, Jagiellonian University, Kraków, Poland/ Doctoral School of Exact and Natural Science, Jagiellonian University, Kraków, Poland

*25P-008 Structure-activity relationship (SAR) study of hydrophobic moiety of nonsecosteroidal VDR ligands using diphenylsilane scaffold Narasinghe Mudiyanselage Hansaka Nirupama Thilakarathne, Takashi Misawa, Yosuke Demizu, Yuya Hanazono, Nobutoshi Ito, Hiroyuki Kagechika, Shinya Fujii Institute of Biomaterials and Bioengineering, Tokyo Medical and Dental University/ Medical Research Institute, Tokyo Medical and Dental University

*25P-009 Structural transformation of a dipeptide Gly-Phe by coffee-ring effect Ayaka Sako, Masaki Saito, Kazuo Eda, Atsuo Tamura Kobe University, Graduate School of Science

25P-010	Exploring the Continuous Conformational Variability of Glutamate Dehydrogenase Using Cryo-EM Single-particle Images and MD Simulations
	<u>Tingting Wang</u> , Osamu Miyashita, Hideki Shigematsu, Masaki Yamamoto, Florence Tama
	Computational Structural Biology Research Team, R-CCS, RIKEN, Japan
25P-011	Preliminary Cryo-EM study of Arabidopsis Magnesium Transporter MRS2-1
	<u>Hexin Xu</u> , Xinyu Teng, Cheng Shen, Yimeng Zhao, Xiaoyu Yang, Natsuko I. Kobayashi, Keitaro Tanoi, Motoyuki Hattori Fudan University, Shanghai, China
25P-012	Application of de novo protein design to structural analysis of non- canonical MgtE Mg2+ channel Zhixuan Zhao, Kimiho Omae, Ziyi Zhang, Xinyu Teng, Cheng Shen, Dangi Sheng, Wataru Iwasaki, Motoyuki Hattori
	Fudan University, Shanghai, China
25P-013	Native lipid NanoDisc application for structural determination of RND transporter
	<u>Kenta Tsutsumi</u> , Atsushi Nakagawa, Eiki Yamashita Institute for Protein Research, Osaka, Japan
25P-014	Preliminary cryo-EM study of the MgtE Mg2+ channel with the PRC- barrel domain
	Ziyi Zhang, Kimiho Omae, Cheng Shen, Wataru Iwasaki, Motoyuki Hattori

Fudan University, Shanghai, China

Protein: Structure & Function

*25P-015 Interpretation of Protein-Corona Formation and Inhibition of Fibrillation by Polyphenol Capped Gold Nanoparticles <u>Atanu Singha Roy</u>, Kakali Baruah, Ajit Kumar Singh, Anupam Nath Jha Department of Chemistry, National Institute of Technology Meghalaya, Shillong

793003, India

- *25P-016 CryoEM-sampling of metastable conformations appearing in cofactorligand association and catalysis of glutamate dehydrogenase Taiki Wakabayashi, Mao Oide, Masayoshi Nakasako Dept. Phys., Keio Univ., Kanagawa, Japan/RIKEN RSC, Hyogo, Japan
- *25P-017 Coarse-Grained Molecular Dynamics Simulations of Rotational Asymmetry in FOF1 ATPase Shintaroh Kubo, Yasushi Okada the University of Tokyo
- ***25P-018** Predicting enzyme function using an empirical approach with machine learning

<u>Suguru Fujita</u>, Tohru Terada Graduate School of Agricultural and Life Science, Faculty of Agriculture, The university of Tokyo.

*25P-019 Ca2+-induced formation of ice-like water network on the surface of type II antifreeze protein from Japanese smelt

<u>Tatsuya Arai</u>, Yue Yang, Sakae Tsuda, Kazuhiro Mio, Yuji Sasaki Graduate School of Frontier Sciences, The University of Tokyo/AIST-UTokyo Advanced Operando-Measurement Technology Open Innovation Laboratory (OPERANDO-OIL)

***25P-020** Predicting protein conformational motions with AlphaFold2 dictated by physical energy landscape

<u>Xingyue Guan</u>, Qianyuan Tang, Weitong Ren, Wenfei Li, Wei Wang Department of Physics, National Laboratory of Solid State Microstructure, Nanjing University, Nanjing 210093, China/Wenzhou Key Laboratory of Biophysics, Wenzhou Institute, University of Chinese Academy of Sciences, Wenzhou, Zhejiang 325000, China

***25P-021** Thermodynamic insights into the antiamyloid activity of lobeline on lysozyme fibrillation

<u>Vibeizonuo Rupreo</u>, Jhimli Bhattacharyya, Ria Saha, Rajib Kumar Mitra Department of Chemistry, National Institute of Technology Nagaland, Dimapur, Nagaland - 797103, India

*25P-022 Functional mechanism of a short wavelength absorbing cation channelrhodopsin, KnChR

Koki Natsume, Shoko Hososhima, Yuzhu Wang, Tatsuki Tanaka, Wataru Shihoya, Osamu Nureki, Hideki Kandori, Satoshi Tsunoda Nagoya Institute of Technology

*25P-023 Structural basis of inhibition and transport in Organic Cation Transporter 1

<u>Yi Cheng Zeng</u>, Meghna Sobti, Ada Quinn, Esther Kristianto, Simon Brown, Nicola Smith, Jamie Vandenberg, Renae Ryan, Alastair Stewart Molecular, Structural and Computational Biology Division, The Victor Chang Cardiac Research Institute, Darlinghurst, NSW, Australia/School of Clinical Medicine, Faculty of Medicine and Health, UNSW Sydney, Sydney, NSW, Australia

*25P-024 Rational Design of High-Affinity Protein Binders by Side Chain Dihedral Correlation Network

Yun-Jung Hsieh, Ta I Hung, Wei-Lin Lu, Chia-en Chang, Kuen-Phon Wu Institute of Biological Chemistry, Academia Sinica, Taipei, Taiwan/Institute of Biochemical Sciences, National Taiwan University, Taipei, Taiwan

*25P-025 Cytoplasmic domain of GtACR1 regulations the channel gating. <u>Hana Maruyama</u>, Shoko Hososhima, Satoshi Tsunoda, Yuya Ohki, Takashi Kikukawa, Takashi Tsukamoto, Hideki Kandori Graduate School of Engineering, Nagoya Institute of Technology

***25P-026** Beyond the Active site: The addition of a remote loop reveals a new complex biological function for chitinase enzymes

Dan Kozome, Adnan Sljoka, Paola Laurino Protein Engineering and Evolution Unit, Okinawa Institute of Science and Technology Graduate University, Okinawa, Japan

*25P-027 Towards the Cryo-EM Structures of Viral Annealase Proteins

Lucy Johanna Fitschen, Jodi Brewster, Jordan Nicholls, Stefan Mueller, Gökhan Tolun

School of Chemistry and Molecular Bioscience, and Molecular Horizons, University of Wollongong, Wollongong, Australia/ARC Industrial Transformation Training Centre for Cryo-electron Microscopy of Membrane Proteins (CCeMMP)

*25P-028 Targeting the oncoprotein GOLPH3 Anastasia Theodoropoulou, Luciano Abriata, Anita Nasrallah,

Francesco Talotta, Sarah Vacle, Fernando Meireles, Maria J. Marcaida, Giovanni D'Angelo, Matteo Dal Peraro Laboratory for Biomolecular Modeling, Institute of Bioengineering, EPFL, Switzerland

25P-029	IgG subclass oligomerization upon antigen binding – Full biophysical characterization of the missing link between antibody binding and complement activation Jürgen Strasser, Nikolaus Frischauf, Aran F. Labrijn, Frank J. Beurskens, Johannes Preiner University of Applied Sciences Upper Austria, Linz, Austria
25P-030	Regulation of enzyme structure and function by weak metal-ion binding <u>Masayuki Oda</u> , Yumi Kitagawa, Takuji Oyama, Kosuke Morikawa Kyoto Prefectural University
25P-031	Structure and function of stomatin-like protein FliL to assist flagellar motor stator PomAB in marine Vibrio Norihiro Takekawa, Tatsuro Nishikino, Ray Burton-Smith, Yuki Tajimi, Mitsuru Ikeda, Kazuyoshi Murata, Seiji Kojima, Takayuki Uchihashi, Katsumi Imada, <u>Michio Homma</u> Div Material Sci, Grad Sch Sci, Nagoya Univ
25P-032	Factors influencing pH-sensitive color changes in firefly bioluminescence were studied through computational analysis of hydrogen bond networks in close proximity to catalytic centers of luciferase and its mutants using QM/MM Kota Nosaka, <u>Naohisa Wada</u> kyoto Luminous Science Laboratory, Kyoto, Japan
25P-033	A double-edged sword: Bacteriophage PlyGRCS endolysin targeting MRSA Staphylococcus aureus isolates and serendipitous discovery of its interaction with a cold shock protein C (CspC) Padmanabhan Balasundaram Department of Biophysics, National Institute of Mental Health and Neuro Sciences (NIMHANS)
25P-034	Deciphering Protein Dynamics and Evolution: Insights from AlphaFold 2's Predicted Aligned Error Qian-Yuan Tang, Liangxu Xie, Xiangze Zeng Hong Kong Baptist University, Hong Kong, China

- 25P-035 Conformational Heterogeneity and Fluorescence Resonance Energy Transfer in the Calcium Indicator Yellow Cameleon YC3.60 Hiroki Tsubota, Yuna Kinoshita, Mamoru Shigeno, <u>Haruko Hosoi</u> Toho University
- 25P-036 Protein Dynamics and Mechanisms from Multiple Structures Robert L Jernigan, Mesih Kilinc, Kejue Jia, Weixia Deng, Pradeep Bk, Rthan Bush Department of Biochemistry, Biophysics and Molecular Biology, Iowa State University, Ames, IA, USA

Protein: Physical property

*25P-037	Role of aggregation-prone segments in fibril formation of the amyloidogenic apolipoprotein A-I variant
	<u>Norihiro Namba,</u> Takashi Ohgita, Hiroko Tamagaki-Asahina,
	Toshinori Shimanouchi, Takeshi Sato, Hiroyuki Saito
	Laboratory of Biophysical Chemistry, Kyoto Pharmaceutical University
*25P-038	Feasibility of immunoglobulin A purification using phosphate-modified zirconia particles
	<u>Shogo Kanoh</u> , Kentaro Shiraki, Katsuya Kato, Atsushi Hirano
	Nanomaterials Research Institute, National Institute of Advanced Industrial Science and Technology (AIST), Tsukuba, Ibaraki 305-8565, Japan/Faculty of Pure and Applied Sciences, University of Tsukuba, 1-1-1 Tennodai, Tsukuba, Ibaraki 305-8573, Japan
*25P-039	Development of a high-throughput data collecting system for
	antibody optimization: thermal stability and interaction kinetics Sae Ito, Ryo Matsunaga, Makoto Nakakido, Daisuke Komura, Hiroto Katoh, Shumpei Ishikawa, Kouhei Tsumoto
	antibody optimization: thermal stability and interaction kinetics Sae Ito, Ryo Matsunaga, Makoto Nakakido, Daisuke Komura, Hiroto Katoh,
*25P-040	antibody optimization: thermal stability and interaction kinetics Sae Ito, Ryo Matsunaga, Makoto Nakakido, Daisuke Komura, Hiroto Katoh, Shumpei Ishikawa, Kouhei Tsumoto
*25P-040	 antibody optimization: thermal stability and interaction kinetics Sae Ito, Ryo Matsunaga, Makoto Nakakido, Daisuke Komura, Hiroto Katoh, Shumpei Ishikawa, Kouhei Tsumoto Department of Bioengineering, School of Engineering, The University of Tokyo. Air/water-interface-induced self-assembly of biosurfactant protein
*25P-040	antibody optimization: thermal stability and interaction kinetics Sae Ito, Ryo Matsunaga, Makoto Nakakido, Daisuke Komura, Hiroto Katoh, Shumpei Ishikawa, Kouhei Tsumoto Department of Bioengineering, School of Engineering, The University of Tokyo. Air/water-interface-induced self-assembly of biosurfactant protein RolA from filamentous fungus Aspergillus oryzae Nao Takahashi, Yuki Terauchi, Takumi Tanaka, Akira Yoshimi, Hiroshi Yabu,

*25P-041 pKa, stretching vibrational frequencies, and nuclear magnetic resonance chemical shifts in H-bond networks of protein environments

Masaki Tsujimura, Keisuke Saito, Hiroshi Ishikita Graduate School of Engineering, The University of Tokyo, Tokyo, Japan

25P-042 The common feature of fibril formation mechanism of α-synuclein and apolipoprotein A-I Takashi Ohgita, Norihiro Namba, Hiroki Kono, Hiroyuki Saito Kyoto Pharmaceutical University

Protein: Function

*25P-043	Regulatory mechanism of HADH and its localization in cell organelles during temperature acclimation in Caenorhabditis elegans Yukina Mori, Misaki Okahata, Akihisa Fukumoto, Yohei Minakuchi, Atsushi Toyoda, Akane Ohta, Atushi Kuhara Faculty of Science and Engineering Konan University & Institute for Integrative Neurobiology, Kobe, Japan
*25P-045	Proposed design of kinetic parameters for agonist antibodies that induce OX40 clustering. Kan Ujiie, Aki Tanabe, Satoru Nagatoishi, Ryo Matsunaga, Kouhei Tsumoto Department of Bioengineering, School of Engineering, The University of Tokyo, Japan
*25P-046	In vitro assembly of a protein capsule and cargo molecules into virus-like particles. <u>Kenya Tajima</u> , Yusuke Sakai, Naohiro Terasaka Earth-Life Science Institute, Tokyo Institute of Technology, Tokyo, Japan
25P-047	FHL complex as a cell strategy to regulate proton motive force and survive under energy limited fermentative conditions Heghine Gevorgyan, Anna Poladyan, Anait Vassilian, <u>Karen Trchounian</u> Laboratory of Microbiology, Bioenergetics and Biotechnology, Research Institute of Biology, Yerevan State University
25P-048	Regulation Mechanism of Liquid-Liquid Phase Separation and Following Aggregation of Fused in Sarcoma by RNA Revealed by

Shinya Tahara, Uchu Matsuura, Shinji Kajimoto, Takakazu Nakabayashi Graduate School of Pharmaceutical Sciences, Tohoku University

Raman Microscopy

Protein: Measurement & Analysis

*25P-049	Platinum (II) stabilizes a molten-globule conformation of a small globular cytosolic protein Suman Tiwari, A.S.R. Koti Department of Chemical Sciences, TIFR, Mumbai, India.
*25P-050	Visualization and quantitative analysis of protein-protein interaction and cell fusion events using split Akaluc complementation in deep tissues Yiling Li, Genki Kawamura, Qiaojing Li, Takeaki Ozawa Department of Chemistry, School of Science, The University of Tokyo, Japan
*25P-051	Time-resolved study of the interaction mechanism between α1-acid glycoprotein and membrane by vacuum-ultraviolet circular-dichroism spectroscopy <u>Satoshi Hashimoto</u> , Koichi Matsuo Graduate School of Advanced Science and Engineering, Hiroshima University
25 P-0 52	Real-time Visualization of Structural Maintenance of Chromosomes Complexes by High-Speed Atomic Force Microscopy Kenichi Umeda, Yumiko Kurokawa, Yasuto Murayama, Noriyuki Kodera Nano Life Science Institute, Kanazawa University, Japan/PRESTO/JST, Japan
25P-053	Sensitivity of various occupancy estimation for synthetic data related to time-resolved serial femtosecond crystallography. <u>Sriram Srinivasa Raghavan</u> , Florence Tama, Osamu Miyashita RIKEN Center for Computational Science, Kobe, Japan.
25 P- 054	A state of partial Rb inactivation and intermediate E2F activation safeguards proliferation commitment Yumi Konagaya RIKEN Center for Biosystems Dynamics Research
25P-055	Real-time HS-AFM observation of EEA1-mediated vesicle fusion in the absence of canonical regulatorsTareg Omer Mohammed, Prem Babu, Shingo Fukuda, Toshio Ando Nano Life Science Institute, Kanazawa University, Kakuma-machi, Kanazawa 920- 1192, Japan

Protein: Design & Engineering

*25P-056	PMBiT: A Bioluminescent Probe for Large Antigen Detection <u>Cheng Qian</u> , Ayumu Ninomiya, Natsuki Shibukawa, Hiroshi Ueda, Takanobu Yasuda, Bo Zhu, Tetsuya Kitaguchi Graduate School of Life Science and Technology, Tokyo Institute of Technology, Kanagawa, Japan
*25P-057	Stabilizing Talin R3 in its Folded State: De Novo Design of a Peptide Binder as a Molecular Lock <u>Yuze Sun</u> , Jie Yan National university of singapore mechanobiology institute
*25 P- 058	Hibody: A Bioluminescent Immunosensor Based on "Trap & Release" of Luciferase-derived Peptide Fused to Antibody <u>Takanobu Yasuda</u> , Bo Zhu, Hiroshi Ueda, Tetsuya Kitaguchi Laboratory for Chemistry and Life Science, Institute of Innovative Research, Tokyo Institute of Technology, Kanagawa, Japan
*25P-059	Increased reaction efficiency by external stimuli-sensitive linker Momoka Takazawa, Koki Kamiya Graduate School of Science and Technology, Gunma University, Kiryu, Gunma, Japan
*25P-060	Design of Proteins that adopt interconvertible two distinct functional conformations <u>Toma Ikeda</u> , Tatsuya Nojima, Hideki Taguchi School of Life Science and Technology, Tokyo Institute of Technology, Japan
*25P-061	The symmetric SAKe protein scaffold Staf Wouters, Andreu Mor Maldonado, Hiroki Noguchi, Kenichi Kamata, Wim Maes, Karen Vanhoorelbeke, Jeremy Tame, Steven De Feyter, Arnout Voet Laboratory for Biomolecular Modeling and Design, KU Leuven, Heverlee, Belgium
*25P-062	Characterization of novel scFv×VHH format of biparatopic antibody against MtsA from Streptococcus pyogenes <u>Risa Asano</u> , Miyu Takeuchi, Makoto Nakakido, Chihiro Aikawa, Takeshi Yokoyama, Yoshikazu Tanaka, Ichiro Nakagawa, Kouhei Tsumoto Dept of Bioeng.Eng., Sch. of Eng., The Univ of Tokyo

25P-063	Functional protein complexes from symmetric designer proteins
	Arnout RD Voet, Staf Wouters, Bram Mylemans, Hiroki Noguchi
	KU Leuven, Belgium

25P-064 Development of a general methodology to design sensor proteins <u>Rie Tatsumi</u>, Nobuyasu Koga Institute for Protein Research (IPR), Osaka University, Osaka, Japan

25P-065A one-pot detection system using β-glucuronidase-based enzyme
switch and label-free antibody
Bo Zhu, Yukihiko Yamasaki, Takanobu Yasuda, Cheng Qian, Zhirou Qiu,
Hiroshi Ueda, Tetsuya Kitaguchi
Laboratory for Chemistry and Life Science, Institute of Innovative Research, Tokyo
Institute of Technology, Yokohama, Japan

Protein: Intrinsic disorder

*25P-066	The Relationship between Self-assembly and Local Dynamics of Intrinsically Disordered Proteins Ryoga Kobayashi, Takashi S. Kodama, Norio Yoshida, Hideki Nakamura, Yohei Miyanoiri, Hidehito Tochio, Naotaka Sekiyama Department of Biophysics, Graduate School of Science, Kyoto University, Kyoto, Japan
*25P-067	Interaction Mechanism of α-Synuclein with Synapsin in the Liquid Condensates Shunki Takaramoto, Keiichi Inoue The Institute for Solid State Physics, University of Tokyo, Kashiwa, Japan
25P-068	Global Analysis of Disordered Proteome in Cells Shouxiang Zhang, Tze Cin Owyong, <u>Yuning Hong</u> La Trobe University, Melbourne, Australia

Heme proteins

J

25P-069	Crystal structures of bovine heart cytochrome c oxidase with inhibitor
	complex
	<u>Tomohiro Ide</u> , Kyoko Shinzawa-Itoh, Kazumasa Muramoto

School of Science, University of Hyogo, Hyogo, Japan.

25P-070 Gold Nanoparticle Thin Film Electrode Enables Direct Electrochemical Control of Cytochrome P450 Reaction Yasuhiro Mie, Chitose Mikami, Yoshiaki Yasutake, Naoki Matsuda Bioproduction Research Institute, AIST

Membrane proteins

- *25P-071 Engineering cardiolipin binding to an artificial membrane protein reveals determinants for lipid-mediated stabilization <u>Mia Louis Abramsson</u>, Robin A Corey, Jan Škerle, Louise Persson, Olivia Andén, Abraham O Oluwole, Rebecca J Howard, Erik Lindahl, Carol V Robinson, David Drew, Kvido Strisovsky, Erik G Marklund, Phillip J Stansfeld, Michael Landreh Department of Microbiology, Tumor and Cell Biology, Karolinska Institutet, Stockholm, Sweden
- *25P-072 Identification of Novel Receptor for Polyphenolic Metabolites <u>Shota Nishikawa</u>, Yuki Masujima, Ryuji Ohue-Kitano, Ikuo Kimura Graduate School of Pharmaceutical Sciences, Kyoto University, Kyoto, Japan
- *25P-073 Mechanism of caffeine-induced functional recovery in RyR2 loss-offunction mutant

<u>Yuya Otori</u>, Raymond Burton-Smith, Nagomi Kurebayashi, Kazuyoshi Murata, Hiroaki Kato, Takashi Murayama, Haruo Ogawa Graduate School of Pharmaceutical Sciences, Kyoto University

- *25P-074 A proton-transfer mechanism in the malaria parasite lactate/H+ symporter suggests a transporter without conformational changes <u>Ciara J F Wallis</u>, Kasimir Gregory, Stephen Fairweather, Ruitao Jin, Sitong He, Giel van Dooren, Adele Lehane, Ben Corry Research School of Biology, The Australian National University, Canberra, Australia
- *25P-075 Ligand binding mechanism analysis of muscarinic acetylcholine receptors utilizing vibrational spectroscopy <u>Moeka Mizuno</u>, Yuya Sugiura, Ryoji Suno, Hideki Kandori, Kota Katayama Graduate School of Engineering, Nagoya Institute of Technology, Aichi, Japan

25P-076	Structural dynamics of potassium ion selective and cyclic nucleotide binding in a CNG channel SthK using ATR-FTIR <u>Tatsuro Nishikino</u> , Hiroto Fukuda, Koki Ogasawara, Yuji Furutani Grad. Sch. of Eng., Nagoya Inst. of Tech., Aichi, Japan.
25P-077	Where is the N-tail? A Computational Study of Intrinsically Disordered Regions of Human ATP-sensitive Potassium Channel <u>Katarzyna Walczewska-Szewc</u> , Wieslaw Nowak Institute of Physics, Faculty of Physics, Astronomy and Informatics, Nicolaus Copernicus University in Torun, ul. Grudziadzka 5, 87-100 Torun, Poland
25P-078	A Novel Gate Operation Pathway of Lipopolysaccharide Transport by Bacterial ABC Transporter MsbA and LptC <u>Kien Xuan Ngo</u> , Toshio Ando Nano Life Science Institute (WPI-NanoLSI), Kanazawa University, Kanazawa, Japan
25P-079	 Pulmonary Surfactant Protein C (SP-C): The role of palmitoyl chains on protein-protein interaction and oligomerization, from time-resolved fluorescence methodologies <u>Manuel Prieto</u>, Michelle Morán-Lalangui, Ana Coutinho, Jesús Pérez-Gil, Luís M. S. Loura, Begoña García-Álvarez 2 Univ Lisbon, IBB Inst Bioengn & Biosci, Inst Super Tecn, P-1049001 Lisbon, Portugal/3 Univ Lisbon, Associate Lab I4HB, Inst Hith & Bioecon, Inst Super Tecn, P-1049001 Lisbon, Portugal
DNA & DNA	binding proteins
*25P-080	Single molecule imaging of DNA higher-order structural formation by human transcription factor Yin Yang 1. Yan Xi, Takada Shoji, Terakawa Tsuyoshi Graduate School of science, Kyoto University, Kyoto, Japan
*25P-081	Differential dynamics specify MeCP2 function at nucleosomes and methylated DNA Gabriella N.L. Chua, John Watters, Paul Dominic Olinares, Joshua Luo, Brian Chait, Shixin Liu

Laboratory of Nanoscale Biophysics and Biochemistry, The Rockefeller University, New York, NY, USA/Tri-Institutional PhD Program in Chemical Biology, New York, NY, USA

Poster Sessions

25P-082 Mediator Mei5-Sae3 Stabilizes Dmc1 Recombinase Clusters for Efficient Assembly on RPA-Coated Single-Stranded DNA Hung-Wen Li, Chin-Dian Wei, Hao-Yen Chang, Chia-Hua Lu, Chih-Chun Chang, Asako Furukohri, Akira Shinohara, Peter Chi Department of Chemistry, National Taiwan University, Taiwan

RNA & RNA binding proteins

- *25P-083 Molecular mechanisms of interaction between RNase I and ribosomes <u>Atsushi Minami</u>, Takehito Tanzawa, Zhuohao Yang, Takashi Funatsu, Takayuki Kato, Tomohisa Kuzuyama, Hideji Yoshida, Tetsuhiro Ogawa Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan
- 25P-084 Nascent pre-ribosomal RNA acts as surfactants that suppress fusion of fibrillar centers in nucleolus <u>Tetsuya Yamamoto</u>, Tomohiro Yamazaki, Kensuke Ninomiya, Tetsuro Hirose Institute for Chemical Reaction Design and Discovery, Hokkaido University

DNA/RNA nanotechnology

*25P-085	Agent model for numerical simulation of the DNA active droplet Kei Goraku, Ryohei Furuichi, Masahiro Takinoue
	Department of Computer Science, Tokyo Institute of Technology, Tokyo, Japan
*25P-086	Construction of DNA droplets capable of autonomously moving by sensing nucleic acids
	<u>Kanta Takagi</u> , Tomoya Maruyama, Masahiro Takinoue
	Department of Computer Science, Tokyo Institute of Technology, Tokyo, Japan
*25P-087	Specific cell binding of functionalized DNA droplets
	Ryoya Hasegawa, Jing Gong, Shin-Ichiro M. Nomura, Masahiro Takinoue
	Department of Life Science and Technology, Tokyo Institute of Technology
*25P-088	DNA droplets based on self-assembled DNA nanostructure polymers
	with programmable multivalency
	Naoki Yoshida, Masahiro Takinoue
	School of Life Science and Technology, Tokyo Institute of Technology, Yokohama, Japan

*25P-089 DNA Nanostructure-based Chromatin-inspired Heterogeneous Fluid Gel Structures

<u>Hong Xuan Chai</u>, Masahiro Takinoue School of Life Science and Technology, Tokyo Institute of Technology, Japan

Nucleic acid: Others

*25P-090 Mechanical diversity and folding intermediates of parallel-stranded G-quadruplexes with a bulge

<u>Yashuo Zhang</u>, Huijuan You School of Pharmacy, Tongji Medical College, Huazhong University of Science and Technology, Wuhan 430030, China

***25P-091** Force propagation in dense DNA solution

<u>Miku Nakao</u>, Saki Matsuyama, Akinori Miyamoto, Yoshihiro Murayama Department of Biomedical Engineering, Tokyo University of Agriculture and Technology

Chromatin & Chromosomes

*25P-092	Molecular motor in a box: a model for chromatin remodelers <u>Sophie Klempahn</u> , Helmut Schiessel Cluster of Excellence Physics of Life, TUD Dresden University of Technology, 01307 Dresden, Germany
*25P-093	Coarse-grained Simulations for Unidirectional Translocation of Bacterial SMC Complex via DNA-segment Capture Masataka Yamauchi, Giovanni B. Brandani, Tsuyoshi Terakawa, Shoji Takada Dept. of Biophysics, Grad. of Sci., Kyoto Univ
25P-095	Theory of viscoelasticity of chromatin and its surrounding environment

Soya Sninkai, Shuichi Onami RIKEN Center for Biosystems Dynamics Research

Water & Hydration & Electrolyte

*25P-096 Effect of osmolytes on the activity of α-amylase Sachika Furukawa, Mafumi Hishida Department of Chemistry, Faculty of Science, Tokyo University of Science, Tokyo, Japan

25P-097 Investigations of hydration structures and dynamics around proteins and peptides with MD simulations <u>Takuya Takahashi</u>, Ryutaro Inou, Yui Nakamura, Simon Hikiri

College of Life Sciences, Ritsumeikan University, Kusatsu, Japan

Molecular genetics & Gene expression

*25P-098 Effects of transcription termination elements on in vitro genome transcription

Keisuke Saito, Yukino Matsui, Nobuhide Doi, Kei Fujiwara Dept. of Biosci. and Info., Keio University

Morphogenesis and Development

*25P-099 Three-dimensional Mechanical Cooperativity Optimises Epithelial Wound Healing Shu En Lim, Rob Tetley, Yanlan Mao University College London

Muscle

*25P-100	Myosin and tropomyosin-troponin complementarily regulate thermal activation of striated muscles
	<u>Shuya Ishii</u> , Kotaro Oyama, Fuyu Kobirumaki-Shimozawa,
	Tomohiro Nakanishi, Naoya Nakahara, Madoka Suzuki, Shin'ichi Ishiwata,
	Norio Fukuda
	QST, Gunma, Japan/Dept Cell Physiol, Sch Med, Jikei Univ, Tokyo, Japan
25P-101	Observation of power stroke coordination in DNA Origami based
	artificial myosin filaments
	<u>Hiroki Fukunaga,</u> Takumi Washio, Keisuke Fujita, Masashi Ohmachi,
	Hiroaki Takagi, Keigo Ikezaki, Toshio Yanagida, Mitsuhiro Iwaki
	Adv ICT Res Inst, NICT

Molecular motor

*25P-102	Spontaneous γ subunit rotation upon conformational changes of the α, β subunits in F1-ATPase Masahiro Motohashi, Mao Oide, Chigusa Kobayashi, Jaewoon Jung, Eiro Muneyuki, Yuji Sugita Fac. Sci. Engineering, Chuo Univ./RIKEN CPR
*25P-103	Characterization of the motility of tetrahymena kinesin 9A and 9B <u>Hiroto Ishii</u> , Masahiko Yamagishi, Junichiro Yajima Graduate School of Arts and Science, The University of Tokyo, Tokyo, Japan
*25P-104	Modeling the motion of heterodimeric kinesins reveals head- head coordination in a KIF1A dimer <u>Tomoki Kita</u> , Kazuo Sasaki, Shinsuke Niwa Tohoku University
*25P-105	Structural analysis of ATP synthases embedded in a lipid bilayer under proton motive force by cryoEM <u>Atsuki Nakano</u> , Jun-ichi Kishikawa, Kaoru Mitsuoka, Ken Yokoyama Fac. of Life Sci., Kyoto Sangyo Univ
25P-106	Application of information theory to understand cooperative force generation between skeletal myosin molecules <u>Motoshi Kaya</u> , Arun Kasimchetty, Hideo Higuchi Department of Physics, University of Tokyo
25P-107	Extreme-Value Analysis of Intracellular Cargo Transport by Motor Proteins Takuma Naoi, Yuki Kagawa, Kimiko Nagino, Shinsuke Niwa, Kumiko Hayashi Institute for Solid State Physics, The University of Tokyo/Department of Applied Physics, Graduate School of Engineering, Tohoku University
25P-108	Comparative analysis of cilia force production in effective and recovery strokes of isolated Volvox carteri cells Ryuta Yamaguchi, Katsuya Shimabukuro National Institute of Technology, Ube College

25P-109 Cryo-EM structure of mammalian V-ATPase. <u>Yui Nishida</u>, Atsuko Nakanishi, Atsuki Nakano, Fuka Ueda, Kaoru Mitsuoka, Ken Yokoyama

Kyoto Sangyo Univ, Kyoto, Japan

Single Molecule Biophysics

*25P-110 Regulation of anticalin-CTLA4 binding mechano-stability by altering protein pulling geometry Yang Sun

Department of Chemistry, University of Basel/Department of Biosystems Science and Engineering, ETH Zurich

*25P-111 Versatile peptide probes for labeling cell-surface GPCR

<u>Toshiki Yoda</u>, Yasushi Sako, Asuka Inoue, Masataka Yanagawa Molecular and Cellular Biochemistry, Graduate School of Pharmaceutical Sciences, Tohoku University, Miyagi, Japan

*25P-112 The role of von Willebrand factor-like Domains in Mucin Adhesion Rebecca Schlatterer, Oliver Lieleg, Bizan N. Balzer

Institute of Physical Chemistry, University of Freiburg, Freiburg, Germany

*25P-113 Self-fueled Peptide Assembly Investigated via AFM-based Imaging Christiane Wenzel, Mahesh Pol, Kun Dai, Charalampos Pappas.

Bizan N. Balzer, Thorsten Hugel Cluster of Excellence livMatS @ FIT – Freiburg Center for Interactive Materials and Bioinspired Technologies, University of Freiburg, Georges-Köhler-Allee 105, D-79110 Freiburg, Germany /Institute of Physical Chemistry, University of Freiburg, Albertstraße 21, D-79104 Freiburg, Germany

*25P-114 Force-Dependent Structural Changes of Filamin C Rod Domains Regulated by Filamin C Dimer Yunxin Deng, Jie Yan

Mechanobiology Institute, National University of Singapore, Singapore 117411

25P-115 State-of-the-art high-speed atomic force microscopy for filming faster biomolecular dynamics

<u>Shingo Fukuda</u>, Akihiro Otomo, Ryota lino, Toshio Ando WPI NanoLSI, Kanazawa Univ.

- 25P-116 In-cell single-molecule FRET measurement of cytosolic RAF proteins Kenji Okamoto, Yasushi Sako RIKEN CPR
- *25P-117 Extent of stochasticity in folding dynamics determines the forcetolerance and longevity of mechanosensing proteins Pritam Saha, Vishavdeep Vashisht, Ojas Singh, Gaurav Kumar Bhati, Surbhi Garg, Dr. Sabyasachi Rakshit Department of chemical sciences, Indian Institute of Science Education and research Mohali, India

Cell biology: Adhesion

25P-118 A Novel Semi-Automatic Software Tool for Focal Adhesion Analysis Joanna Hajduk, Patrycja Twardawa, Zenon Rajfur Doctoral School of Exact and Natural Sciences, Jagiellonian University, Łojasiewicza 11, 30-348 Cracow, Poland/Faculty of Physics, Astronomy and Applied Computer

Science, Jagiellonian University, Łojasiewicza 11, 30-348 Cracow, Poland

Cell biology: Motility

*25P-119	Effect of Substrate Elasticity on Adhesion and Motility of Cancer Cells Shotaro Yamamoto, Tomoko Oyama, Kotaro Oyama, Mitsumasa Taguchi, Hiromi Miyoshi Tokyo Metropolitan University
*25P-120	Decoding Antidote Access: Binding/Unbinding Pathways toward Organophosphate-Inhibited HuAChE Kowit Hengphasatporn, Nalinee Kongkaew, Thanyada Rungrotmongkol, Yasuteru Shigeta, Ryuhei Harada Center for Computational Sciences, University of Tsukuba, Tsukuba, Ibaraki, Japan
*25 P-121	β-arrestin acts as an inhibitor of trimeric G protein signaling in eukaryotic chemotaxis <u>Masaki Muromoto</u> , Satomi Matsuoka, Masahiro Ueda Graduate School of Frontier Biosciences, Osaka University

*25P-122	Conversion from Linear Contraction to Rotation of Stress Fibers in Migrating Keratocytes <u>Chika Okimura</u> , Shu Akiyama, Yukinori Nishigami, Tatsunari Sakurai, Yoshiaki Iwadate Department of Biology, Yamaguchi University
*25P-123	Rebirth of Fish Epidermal Keratocyte Sheets Norihiko Nishimura, Chika Okimura, Yoshiaki Iwadate Department of Biology, Yamaguchi University
*25P-124	Integration between Epidermal Keratocyte Sheets Accompanied by Rapid Disassembly of Actomyosin Cables <u>Kazuma Shimizu</u> , Chika Okimura, Yoshiaki Iwadate Department of Biology, Yamaguchi University
*25 P -125	Side-by-side interaction of adjacent cells dominates the collaborative dynamics and ordering of collective cells <u>Mitsuru Sentoku</u> , Miki Takei, Masahru Endo, Kenji Yasuda Department of Pure and Applied Physics, Graduate School of Advanced Science and Engineering, Waseda University, Tokyo, Japan
*25P-126	Mimicking dynamics of human gastrulation: microprint culture of two types of cells derived from human iPS cells Ryo Kojima, Hazuki Tuboi, Miyu Mori, Chihiro Takeuchi, Kiyoshi Ohnuma Department of Bioengineering, Nagaoka University of Technology
25P-127	Structural and Functional Insights into Drosophila melanogaster Sperm Flagella: A Focus on Axonemal Architecture and Beating Patterns Sho Tamai, Kosei Sato, <u>Kazuhiro Oiwa</u> Graduate School of Science, University of Hyogo/ National Institute of Information and Communications Technology
25P-128	Structural Changes of Beating Comb Plates of Ctenophores during Effective and Recovery Strokes as Probed by Time-resolved X-ray Diffraction Recording <u>Hiroyuki Iwamoto</u> , Mio Kosaka, Ryo Yokoya, Kei Jokura, Kazuhiro Oiwa, Kazuo Inaba

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Poster Sessions

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- 25P-129 Identifying direct and indirect interactions among collectively moving individuals using pairwise information flow metric <u>M. Mohiuddin</u>, Sulimon Sattari, Udoy S. Basak, Tamiki Komatsuzaki Graduate School of Chemical Sciences and Engineering, Hokkaido University, Japan/ Comilla University, Cumilla-3506, Bangladesh
- 25P-130 Structural and functional analyses of the C-terminal cytoplasmic domain of a flagellar export gate protein, FlhB <u>Miki Kinoshita</u>, Tomoko Miyata, Keiichi Namba, Tohru Minamino 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

*25P-131	Crosstalk of two bacterial actins composed of the force generation unit of Spiroplasma swimming
	<u>Daichi Takahashi,</u> Makoto Miyata, Ikuko Fujiwara
	Research Institute for Interdisciplinary Science, Okayama University, Japan/Graduate School of Science, Osaka Metropolitan University, Japan
*25P-132	Microtubule Fatigue Under Repetitive Mechanical Stress
	<u>Syeda Rubaiya Nasrin,</u> Akira Kakugo, Neda M. Bassir Kazeruni, Masatoshi Ichikawa Kyoto University
25P-133	Dominant negative mutations in γ-tubulin cause partial loss of protofilaments in centriole triplet microtubules Yuki Nakazawa, Mao Horii, Akira Noga, Ken-ichi Wakabayashi, Masafumi Hirono
	Dep. Frontier Biosci., Hosei Univ., Tokyo, Japan/STG, OIST, Okinawa, Japan
25P-134	Actin fluctuations regulate cofilin binding Akihiro Narita Nagoya Univ.
25P-135	Signaling Mechanisms to Regulate Activation of Actin Depolymerization Factor Cofilin in Mast Cells
	<u>Ruriko Suzuki</u> , Satoru Yokawa, Tadahide Furuno, Naohide Hirashima Grad. Sch. Pharm. Sci., Nagoya City Univ., Nagoya, Japan

Poster Sessions

June 25 [Tue]

Cell biology: Signal transduction & Cell membrane

***25P-136** The maximum phagocytic limit of macrophages is determined by the maximum expansion ability of the local cell membrane surrounding antigens.

Dan Horonushi, Sota Suzuki, Maiha Ando, Haruka Yuki, Kenji Yasuda Department of Pure and Applied Physics, Graduate School of Advanced Science and Engineering, Waseda University, Tokyo, Japan.

*25P-137 Lipid domains in the inner leaflet of cell plasma membranes serve as a signaling platform for K-Ras <u>Toshiki Mori</u>, Koichiro M. Hirosawa, Rinshi S. Kasai, Tomohiko Taguchi, Yasunari Yokota, Kenichi G.N. Suzuki

UGSAS, Gifu Univ., Japan

25P-138 ERK-mediated STAT3 inhibition causes dynamic heterogeneity in IL-6 signaling

Keisuke Fujita, Masahiro Ueda Laboratory for Cell Signaling Dynamics, RIKEN BDR, Osaka, Japan

25P-139 Intracelluar information flow in RAS-MAPK signaling Nobuhisa Umeki, Yoshiyuki Kabashima, <u>Yasushi Sako</u> Cellular Informatics Laboratory, RIKEN, CPR , Wako, Japan

Biological & Artificial membrane: Structure & Property

*25P-140 Protein accumulation on amphiphilic protein-phospholipid hybrid leaflet <u>Masato Suzuki</u>, Koki Kamiya

Graduate School of Science and Technology, Gunma University

*25P-141 Creation of Asymmetric Membrane Vesicles with a Protein Inner Membrane Mixed with Phospholipids Yuki Nagai, Koki Kamiya Graduate School of Science and Technology, Gunma Univ., Gunma, Japan

*25P-142 Triglyceride-Tethered Membrane Lipase Sensor Samara Elizabeth Bridge, Upeksha Mirissa Lankage, Bruce Cornell, Stephen Holt, Matt Padula, Charles Cranfield School of Life Sciences, University of Technology Sydney, Ultimo, NSW 2007, Australia

25P-143	Characterization of Lipid Vesicles Adsorbed on Bovine Serum Albumin: Adhesion and Large Nano-indentation
	<u>Eman Ramadan Sarsour,</u> Tomohiro Hayashi
	Department of Physics, Faculty of Science, Helwan University, Cairo, Egypt
25P-144	Decoding functional oligomeric states of membrane-associated
	protein oligomers forming membrane pores
	Radek Šachl, Vandana Singh, Sabína Čujová, Petra Riegerová, Martin Hof,
	Julia P. Steringer, Walter Nickel
	Department of Biophysical Chemistry, J. Heyrovsky' Institute of Physical Chemistry of the Academy of Sciences of the Czech Republic, Prague, 182 23, Czech Republic

25P-145 Solid-Supported-Membrane-Based Electrophysiology: Application to the Analysis of Membrane Binding Ronald J Clarke, Francesco Tadini-Buoninsegni School of Chemistry, University of Sydney, Sydney NSW, Australia

Biological & Artificial membrane: Dynamics

*25P-146	Unraveling of the mechanisms of hierarchical mesoscale domain organization in cell plasma membranes by super-resolution microscopy and single-molecule tracking. <u>Toui Kawai</u> , Rinshi S. Kasai, Koichiro M. Hirosawa, Yasunari Yokota, Takahiro K. Fujiwara, Akihiro Kusumi, Kenichi G. N. Suzuki United Grad. Sch, Agri, Sci, Gifu Univ., Gifu, Japan
*25P-147	Impact of peptides on the solubility of Amphotericin B and its sterol- specific membrane activity Lissy M. Hartmann, Stephen A. Holt, Robert A. Russell, Anton P. Le Brun, Anneka C. Pereira Schmidt, Chandra H. Chavali, Evelyne Deplazes, Charles G. Cranfield School of Life Sciences, University of Technology Sydney, 15 Broadway, Ultimo, NSW 2007, Australia
*25 P-148	Acceleration of lipid exchange reaction between human cells and supported lipid bilayers <u>Asahi Gono</u> , Takashi Okuno Faculty of Science, Yamagata Uniy.

25P-149 Antibacterial activity of C-terminal fragments of NEMURI Moynul Hasan Institute for Genetic Medicine, Hokkaido University, Japan.

Biological & Artificial membrane: Excitation & Channels

- *25P-150 Photocaged amino acid method elucidates the potency of individual positively-charged residues in PIP2-dependency of the Kir2.1 inwardly rectifying potassium channel <u>Junxian Zhou</u>, Natsuki Mizutani, Kohei Yamamoto, Yoshifumi Okochi, Yasushi Okamura Graduate School of Medicine, Osaka University
- **25P-151** ATP directly regulates the voltage-gated proton channel <u>Akira Kawanabe</u>, Kohei Takeshita, Maki Takata, Yuichiro Fujiwara Faculty of Medicine, Kagawa University

Membraneless Organella, autophage, Liquid-liquid phase separation

*25P-153	Coalescence of liquid or gel-like DNA-encapsulating micro-droplets
	Takashi Nishio, Helmut Schiessel
	Cluster of Excellence Physics of Life, TUD Dresden University of Technology/ Biomedical Research Institute, National Institute of Advanced Industrial Science and Technology (AIST)
*25P-154	Aberrant phase transition of stress granules in living cells observed by Raman/Brillouin microscopy and machine learning
	<u>Ren Shibuya</u> , Shinji Kajimoto, Hideyuki Yaginuma, Tetsuro Ariyoshi, Yasushi Okada, Takakazu Nakabayashi
	Graduate School of Pharmaceutical Sciences, Tohoku University, Japan
*25P-155	Molecular weight polydispersity initiates nucleation of polymer blends
	around the phase separation boundary
	<u>Akari Kamo</u> , Arash Nikoubashman, Miho Yanagisawa
	Department of Physics, Graduate School of Science, The University of Tokyo, Japan
*25P-156	Hyperphosphorylation of nucleolar protein Nopp140 drives mitotic
	nucleolar disassembly.
	Hisashi Shimamura, Yuki Norizoe, Takahiro Sakaue, Shige H. Yoshimura
	Faculty of Integrated Human Studies, Kyoto University

25P-157 Molecular dynamics of autophagosomal lipid transfer Yuji Sakai, Kazuaki Matoba, Nobuo Noda, Yuji Sugita

YUJI Sakai, Kazuaki Matoba, Nobuo Noda, YUJI Sugita Institute for Life and Medical Sciences, Kyoto University/RIKEN iTHEMS/Graduate School of Medicine, The University of Tokyo

25P-158 Designer coacervates for protein sequestration

Akihiro Kishimura, Biplab K C, Ryoma Omae, Hiroshi Kamizawa, Gakuto Takeda, Takumi Yamada, Hinano Nakamoto, Teruki Nii, Takeshi Mori, Yoshiki Katayama Department of Applied Chemistry, Faculty of Engineering, Kyushu University, Fukuoka, Japan/Graduate School of System Life Sciences, Kyushu University, Fukuoka, Japan/ Center for Molecular Systems (CMS), Kyushu University, Fukuoka, Japan/Center for Future Chemistry, Kyushu University, Fukuoka, Japan

25P-159 Highly Charged Proteins and Their Repulsive Interactions in Regulation of Biomolecular Condensation

<u>Cheng Tan</u>, Jaewoon Jung, Yuji Sugita RIKEN Center for Computational Science

25P-160 A Key Role of Less Bulky-Hydrophobic Amphipathic α-helix in Autophagy

Taki Nishimura, Gianmarco Lazzeri, Noboru Mizushima, Roberto Covino, Sharon A Tooze

PRESTO, Japan Science and Technology Agency, Tokyo, Japan/Department of Biochemistry and Molecular Biology, Graduate School of Medicine, The University of Tokyo, Tokyo, Japan/Molecular Cell Biology of Autophagy Laboratory, The Francis Crick Institute, London, UK

Chemoreception

*25P-161 Ligand recognition of the Vibrio cholerae chemoreceptor for two distinct attractants, pyruvate and serine

<u>Fuga Omori</u>, Hirotaka Tajima, Ikuro Kawagishi Grad. Sch. Sci. and Engin., Hosei Univ

Neuroscience & Sensory systems

*25P-162 HS-AFM reveals the structural role of CaMKII in synaptic structural plasticity

<u>Taisei Suzuki</u>, Takashi Sumikama, Hideji Murakoshi, Mikihiro Shibata Grad. Sch. NanoLS., Kanazawa University, Ishikawa, Japan

- *25P-163 Physics of transport through capillaries and the blood-brain barrier: comparative study of hydrogel phantom and living mouse models <u>Anastasia S Vanina</u>, Alexander Sychev, Ivan Proskurkin, Anastasia Lavrova, Eugene Postnikov Kursk State University
- 25P-164 Spontaneous depolarization wave in the embryonic CNS: optical imaging with a voltage-sensitive dye Yoko Momose-Sato, Katsushige Sato Kanto-Gakuin University, College of Nutrition
- 25P-165 Oscillations in the embryonic chick olfactory bulb: optical imaging with a voltage-sensitive dye <u>Katsushige Sato</u>, Yoko Momose-Sato Komazawa Women's University, Faculty of Human Health

Neuronal circuit & Information processing

*25P-166	Environmental oxygen information generates temperature response
	diversity in C. elegans
	Misaki Okahata, Taichiro Iki, Sawako Yoshina, Yohei Minakuchi,
	Shohei Mitani, Toshie Kai, Toru Miura, Atsushi Toyoda, Akane Ohta,
	Atsushi Kuhara
	Inst. for Integrative Neurobio., Konan Univ, Japan/Graduate School of Frontier
	Biosciences Osaka Univ., Japan
25P-167	Construction of Single-Cell Level Linear Neural Network with Agarose
	Micro Fabrication Technology
	<u>Shion Sakamoto</u> , Kentaro Kito, Masahito Hayashi, Tomoyuki Kaneko
	Hosei university/LaRC/FB

Behavior

25P-168 Anticipation Behavior of the Physarum Plasmodia to Periodic Light Stimulus

Kazuki Moriguchi

Graduate School of Systems Information Science, Future University Hakodate, Hokkaido, Japan

Photobiology: Vision & Photoreception

*25P-169 The mechanism regulating the binding properties of retinal isomers in opsins

<u>Chihiro Fujiyabu</u>, Takahiro Yamashita Kyoto University, Kyoto, Japan

*25P-170 Light-induced structural changes of a rhodopsin domain in a rhodopsin-bestrophin giant ion channel complex studied by time-resolved infrared spectroscopy

Honda Nastuki, Rei Yoshizumi, Kandori Hideki, Furutani Yuji Graduate School of Engineering, Nagoya Institute of Technology

*25P-171 Exploration of the Diversity of Absorption Spectra in Vertebrate Retinal Photo-isomerase, RGR

> <u>Chunyangguang Li</u>, Takashi Nagata, Naoya Morimoto, Keiichi Inoue The Institute for Solid State Physics, The University of Tokyo, Kashiwa, Japan

*25P-172 Exploring the spectral tuning mechanism of bestrhodopsin from Phaeocystis antarctica

> <u>Yifan Liu</u>, Masae Konno, Inoue Keiichi, Ariel Chazan, Andrey Rozenberg, Oded Béjà The Institute for Solid State Physics, The University of Tokyo, Japan

25P-173 In Vitro Analysis of the Effect of Narrowband and Broadband Light in Visible Range on Lens Epithelial Cell Migration <u>Hiromi Miyoshi</u>, Aki Nishida, Masafumi Otomo, Takuto Suzuki, Yuki Tani Department of Mechanical Systems Engineering, Tokyo Metropolitan University

25P-174 Insights into light-driven chloride ion pump mechanism of NM-R3 and NpHR by molecular dynamics simulation Masahiko Taguchi, Akiya Moriuchi, Hinano Ogawa, Osamu Miyashita, Eriko Nango IMRAM Tohoku Univ./Grad. Sch. Sci. Tohoku Univ.

Photobiology: Photosynthesis

*25P-175 Structure of S2 High-Spin State Manganese Cluster of Photosystem II by Multi-frequency Electron Paramagnetic Resonance (EPR) Spectroscopy

Kosaki Shinya, Nakajima Yoshiki, Shen Jian-Ren, Mino Hiroyuki Grad. Sch. Sci., Nagoya Univ., Aichi, Japan

*25P-176 Oxygen-evolving photosystem II structures during S1–S2–S3 transitions

Hongjie Li, Yoshiki Nakajima, Michihiro Suga, Jian-Ren Shen Research Institute for Interdisciplinary Science and Graduate School of Natural Science and Technology, Okayama University, Okayama, Japan

25P-177 Theoretical analysis of the light-harvesting process in C. thermophilum type-I reaction center that binds three different species of chlorophyll molecules

> Wataru Shimooka, Hirotaka Kitoh-Nishioka, Shigeru Itoh, <u>Akihiro Kimura</u> Department of Physics, Graduate School of Science, Nagoya University

25P-178 Electron Transfer Reactions in the Photosynthetic Reaction Center Complex lacking Iron-Sulfur Cluster Fx of Green Sulfur Bacterium Chlorobaculum tepidum

> Tomomi Inagaki, Yukie Kojima, Kazuki Terauchi, Chihiro Azai Graduate School of Life Sciences, Ritsumeikan University, Shiga, Japan

Photobiology: Optogenetics & Optical control

*25P-179 Antitumor effects of photo-induced cell death using an outward proton pump rhodopsin Shin Nakao, Keiichi Kojima, Naoya Kenmotsu, Yosuke Togashi, Yuki Sudo Grad. Sch., Med. Dent. and Pharm. Sci., Okayama Univ., Okayama, Japan.

25P-180 Microbial Rhodopsin Engineering through Machine Learning and Automated Experiments

Keiichi Inoue, Takashi Nagata, Masae Konno, Masayuki Karasuyama, Yu Inatsu, Kazuhito V. Tabata, Ichiro Takeuchi The Institute for Solid State Physics, The University of Tokyo/RIKEN Center for Advanced Intelligence Project

Radiobiology & Active oxygen

25P-181 Analysis of Radiation-induced Stem Cell Competition and Bystander Response Using Titanium Characteristic X-ray Microbeam <u>Masanori Tomita</u>, Yuki Fujimichi, Atsushi Ito Sustainable Sys. Res. Lab., CRIEPI, Chiba, Japan

Origin of life & Evolution

*25P-182	The role of non-biological membraneless polyester microdroplets as protocells at the origins of life <u>Tony Z Jia</u> , Kuhan Chandru Earth-Life Science Institute, Tokyo Institute of Technology, 2-12-1-IE-1 Ookayama, Meguro-ku, Tokyo 152-8550, Japan/Blue Marble Space Institute of Science, 600 1st Ave, Floor 1, Seattle, WA 98104, USA,
*25P-183	Creation of a Membraneless Protocell with Earth-abundant Transition Metal Catalysts Chen Chen, Tony Z. Jia, Ryuhei Nakamura Biofunctional Catalyst Research Team, RIKEN Center for Sustainable Resource Science (CSRS), Wako, Japan
*25P-184	Primordial Evolution by Linking Sequence Information and Vesicle Reproduction <u>Akiko Baba</u> , Keidai Sato, Shuna Asanuma, Ivo Henkys, Tomoko Kawahata, Ulf Olsson, Anna Wang, Masayuki Imai Grad. Sch. Sci., Tohoku Univ.
25P-185	Genetic properties influencing transcriptional variability Saburo Tsuru, Chikara Furusawa Universal Biology Institute, Graduate School of Science, The University of Tokyo, Tokyo, Japan

Synthetic biology & Artificial cells

*25P-186 Optimizing the in vitro expression profile of central dogma-related proteins <u>Chisato Nishizawa</u>, Shunsuke Aburaya, Yuishin Kosaka, Kenji Sugase, Wataru Aoki Graduate School of Agriculture, Kyoto University, Kyoto, Japan

*25P-187 Characterization of ribosome biogenesis in vitro <u>Yuishin Kosaka</u>, Yumi Miyawaki, Megumi Mori, Shunsuke Aburaya, Chisato Nishizawa, Takeshi Chujo, Tatsuya Niwa, Takumi Miyazaki, Takashi Sugita, Hideki Taguchi, Kazuhito Tomizawa, Kenji Sugase, Mitsuyoshi Ueda, Wataru Aoki Kyoto University *25P-188 Light-Induced Control of Directional Movement in Chlamydomonas-

25P-188 Light-Induced Control of Directional Movement in Chlamydomonas-Encapsulated Liposomes

<u>Hiromasa Shiraiwa</u>, Koichiro Akiyama, Shunsuke Shiomi, Masahito Hayashi, Tomoyuki Kaneko LaRC, FB, Grad. Sch. Sci. & Eng., Hosei Univ., Tokyo, Japan

*25P-189 Construction of asymmetric lipid-protein membrane tension sensing system by using mechanosensitive channels <u>Kotaro Baba</u>, Koki Kamiya Graduate School of Science and Technology, Gunma University, Gunma, Japan

25P-190 Designing a reproduction cycle of vesicles coupled with artificial metabolic pathways

<u>Minoru Kurisu</u>, Peter Walde, Masayuki Imai Department of Physics, Graduate School of Science, Tohoku University, Sendai, Japan

25P-191 Dynamic Instability of Totally-synthetic Supramolecular Dipeptide Fibers upon Hybridization of Surfactant Micelles Ryou Kubota, Shogo Torigoe, Kazutoshi Nagao, Yuya Hamanaka, Itaru Hamachi Graduate School of Engineering, Kyoto University

Genome biology

*25P-192 Heterogeneity of Genomic Sequence within Population in Single Plaque of Influenza Virus Revealed by Revio analysis Kenji Tamao, Masayuki Su'etsugu, Hiroyuki Noji, Kazuhito Tabata Appl.Chem., Grad.Sch.Eng., Univ. Tokyo

Computational biology: Bioinformatics

*25P-193 Improving Protein Complex Prediction through the Generation of Multiple Decoy Structures using Docking Software and Aggressive Refinement by AlphaFold2 Seiya Tanaka, Masaki Koyama, Hiroki Onoda, Leonard Chavas, George Chikenji Nagoya University

Computational biology: Molecular simulation

*25P-195	Elucidating the Binding Pathway of 'Abltide' to Abl Kinase through Enhanced 2D Replica Exchange Molecular Dynamics Simulations Yichao Wu Osaka University, WPI Premium Research Institute for Human Metaverse Medicine (WPI-PRIMe)
*25 P-196	Exploring Protein-Lipid Interactions in Membranes: A Coarse-Grained Perspective with Implicit Solvent Modeling <u>Diego Ugarte</u> , Shoji Takada, Yuji Sugita RIKEN R-CCS, Kobe, Japan/RIKEN BDR, Kobe, Japan/RIKEN CPR, Saitama, Japan
*25 P-197	Investigating TDP43 Condensation and Contributions of Ions: A Multiscale Comparative Analysis of Coarse-Grained Models Yangyang Zhang, Cheng Tan, Yuji Sugita RIKEN Center for Computational Science, Kobe, Japan
*25P-198	Virtual alanine scan for entire sequence of SARS-CoV-2 main protease complexed with ensitrelvir <u>Ayato Mizuno</u> , Tomoki Nakayoshi, Koichi Kato, Eiji Kurimoto, Akifumi Oda Faculty of Pharmacy, Meijo University, Aichi, Japan

*25P-199 Simulation of Lipid Membranes and Their Interaction with Polystyrene Nora Kremer, Thorsten Koslowski University of Freiburg *25P-200 Enhancing protein conformation sampling with coevolution Antoni Marciniak, Darko Mitrovic, Lucie Delemotte KTH Royal Institute of Technology/SciLifeLab *25P-201 Charge-Charge Interactions in Molecular Dynamics Simulations of Glycans, Glycosaminoglycans, and Lipopolysaccharides Denys Biriukov, Miguel Riopedre-Fernández, Hector Martinez-Seara National Centre for Biomolecular Research, Faculty of Science, Masaryk University, Brno, Czech Republic/Central European Institute of Technology, Masaryk University, Brno, Czech Republic/Institute of Organic Chemistry and Biochemistry, Czech Academy of Sciences, Prague, Czech Republic *25P-202 Quantifying chromosome structural dynamical pathways during cell fate decision making process Xiakun Chu Advanced Materials Thrust, Function Hub, The Hong Kong University of Science and Technology (Guangzhou), Nansha, Guangzhou, Guangdong 511400, China *25P-203 Physical determinants of multiphase organisation in protein/RNA condensates Pin Yu Chew, Jerelle A. Joseph, Rosana Collepardo-Guevara, Aleks Reinhardt Yusuf Hamied Department of Chemistry, University of Cambridge, Cambridge, United Kingdom *25P-204 Molecular determinants of lipid selectivity of VPS13 lipid transport protein Thanchanok Chanachanvong Thanchanok Chanachanvong, Puey Ounjai, Tanadet Pipatpolkai *25P-205 Ligand dependent conformational plasticity that guides substrate transport cycle of ABC transporters Sungho Bosco Han, Jim Warwicker, Hao Fan, Stephen Prince The University of Manchester, Manchester, United Kingdom/Agency for Science, Technology and Research (A*STAR), Singapore

25P-206	Exploring 3D cell spreading in supramolecular hydrogels and dynamics-induced hydrogel surface reconfiguration through molecular simulations Tianjie Li, Chun Hon Lau, <u>Yi Wang</u> Department of Physics, The Chinese University of Hong Kong, Shatin, N.T., Hong Kong, China
25P-207	The role of computational approaches in uncovering mechanisms of ferroptotic cell death signal Karolina Mikulska-Ruminska Faculty of Physics, Astronomy and Informatics, Nicolaus Copernicus University in Torun, Poland
25P-208	Dimerization of APP-C99 using BE-ABMD simulations Shingo Ito, Yuji Sugita RIKEN, Cluster for Pioneering Research
25P-209	Computer aided engineering of nonstandard biotechnological enzyme – nitrile hydratase case Lukasz Peplowski Department of Biophysics, Nicolaus Copernicus University, Torun, Poland
25P-210	Why bestatin inhibitor prefers human carnosinase II (CN2) to human carnosinase I (CN1): Simulation study. Borvornwat Toviwek Department of Chemistry, Faculty of Science, Kasetsart University, Chatuchak, Bangkok, 10900, Thailand
25P-211	Computational and Biochemical Studies on the Molecular Interactions Between Melanopsin and its AntagonistsC Ruisi Zou, Kohei Obayashi, Hisao Tsukamoto, Toshifumi Mori Interdisciplinary Graduate School of Engineering Sciences, Kyushu University
25P-212	Atom Filtering Algorithm and GPU-Accelerated Calculation of Simulation Atomic Force Microscopy Images Romain Amyot, Noriyuki Kodera, Holger Flechsig WPI-NanoLSI, Kanazawa University, Kanazawa, Japan/JSPS International Research Fellow

25P-213 Small GTPase Ran: exploring nucleotide-specific conformations Erika Balog, Janka Czigleczki, Balint Dudas, Pedro Tulio de Resende Lara, David Perahia, Hyunbum Jang, Ruth Nussinov Department of Biophysics and Radiation Biology, Semmelweis University, Budapest, Hungary

- 25P-214 Large-scale coarse-grained MD simulations for heterogeneous biomolecular systems by efficient parallelization Jaewoon Jung, Cheng Tan, Yuji Sugita RIKEN R-CCS/RIKEN CPR
- 25P-215
 Molecular dynamics simulation of amyloid-β aggregates

 Hisashi Okumura, Satoru G. Itoh
 Exploratory Research Center on Life and Living Systems/Institute for Molecular

 Science/Graduate Institute for Advanced Studies
 Studies

Computational biology: Biological modeling and simulation

- *25P-216 Mathematical model of glioma cell migration and deformation dependent on adhesion dynamics to extracellular matrix Haruna Tagawa, Daisuke Kanematsu, Asako Katsuma, Naoyuki Inagaki, Yonehiro Kanemura, Yuichi Sakumura Nara Institute of Science and Technology, Nara, Japan
- *25P-217 Reassessing the Exon-Foldon correspondence using Frustration Analysis

Ezequiel Alejandro Galpern, Hana Jaafari, Carlos Bueno, Peter G. Wolynes, Diego U. Ferreiro Protein Physiology Lab, Instituto de Química Biológica de la Facultad de Ciencias

Exactas y Naturales, CONICET - Universidad de Buenos Aires, C1428EGA, Buenos Aires, Argentina

- *25P-218 Computational study of the agonism/antagonism effect of small molecules to toll-like receptor (TLR) 7 Ruitao Jin, Sitong He, Ben Corry Australian National University
- *25P-219 Torsion Angles to Map Protein Conformational Changes <u>Katie Blaze O'Flynn</u>, Helen Ginn Deutsches Elektronen-Synchrotron DESY, Hamburg, Germany

*25P-220	Simulating three dimensional epithelial monolayer tissue deformation using cell center model <u>Tomohiro Mimura</u> , Yasuhiro Inoue Department of Micro Engineering, Graduate School of Engineering, Kyoto University, Kyoto, Japan
*25P-221	Development and validation of novel anticancer drugs against protein kinase D2 Ahmed Shemy, Olivia Appelmans, Lauren Voets, Johan Van Lint, Wim De Borggraeve, Arnout Voet KU Leuven, Department of Chemistry, Biochemistry, Molecular and Structural Biology, Celestijnenlaan 200G - box 2403, B-3001 Leuven, Belgium
*25P-222	An investigation of the molecular mechanisms underpinning the aggregation of POR-BT isomers within membranes using molecular dynamics simulations <u>Manish Kesherwani</u> , Qian Wu, Masayasu Taki, Yoshiki Tanaka, Quan Manh Phung, Sawako Enoki, Yasushi Okada, Shigehiro Yamaguchi, Florence TAMA Institute of Transformative Bio-Molecules, Nagoya University, Nagoya, Japan
25P-223	Membrane morphology of Clathrin-Mediated Endocytosis <u>Suguru Ushioda</u> , Masashi Tachikawa Tachikawa Lab, Faculty of Science, Yokohama City Univ.
25P-224	Novel Dengue Vaccine Development – A Multiscale Simulation Study Jan K. Marzinek, Raghuvamsi Palur, Peter Bond Bioinformatics Institute (A*STAR), Singapore
25P-225	Navigating Bio-Systems Through A Deep Learnt Lens-Scape of Multiscale Analytics Haibin Su The Hong Kong University of Science and Technology
25P-226	Exploring intermediate states along binding of inhibitors to protein kinases using large-scale molecular dynamics simulations <u>Ai Shinobu</u> , Suyong Re, Hiraku Oshima, Yuji Sugita WPI Premium Research Institute for Human Metaverse Medicine, Osaka University, Japan /RIKEN Center for Biosystems Dynamics Research, Japan

Poster Sessions

June 25 [Tue]

Computational biology: machine learning for molecules or cell systems

*25P-227 Prediction of Olfactory Perception From Learned Molecular Representation

Zi Hui Lau, Tetsuya J. Kobayashi Department of Electrical Engineering and Information Systems (EEIS), Graduate School of Engineering, The University of Tokyo, Bunkyo-ku, Tokyo, 113-8656, Japan

*25P-228 Label-free detection of senescent cells using Raman imaging and machine learning

<u>Hiroko Kodama</u>, Ren Shibuya, Hiroaki Takahashi, Shinji Kajimoto, Takakazu Nakabayashi Faculty of Pharmaceutical Science, Tohoku Univ.

Mathematical & Theoretical biology

*25P-229	Theory for Optimal Estimation and Control with Resource Limitations in Biological Information Processing
	Takehiro Tottori, Tetsuya Kobayashi
	Laboratory for Neural Computation and Adaptation, RIKEN Center for Brain Science/ Institute of Industrial Science, The University of Tokyo
*25P-230	Framework for efficient drug selection using machine learning Shunta Nonaga, Koji Tabata, Tamiki Komatsuzaki
	Graduate School of Chemical Sciences and Engineering, Hokkaido University, Sapporo, Japan
*25P-231	THEORETICAL STUDY ON PARTIAL AND TOTAL ADAPTATION OF MULTIPLE TISSUES UNDER FORCE INTERACTION Ryunosuke Suzuki, Taiji Adachi
	Kyoto University, Kyoto, Japan
25P-232	Stoichiometric constraints alter thermodynamic fates of growing
	systems
	<u>Atsushi Kamimura,</u> Yuki Sughiyama, Tetsuya J. Kobayashi
	The University of Tokyo

25P-233 Information Geometry of Equilibrium and Nonequilibrium Chemical Reaction Networks

<u>Tetsuya J. Kobayashi</u>, Dimitri Loutchko, Atsushi Kamimura, Shuhei Horiguchi, Yuki Sughiyama Institute of Industrial Science, The University of Tokyo, Japan/Department of Mathematical Informatics, Graduate School of Information Science and Technology, The University of Tokyo, Japan/Universal Biology Institute, The University of Tokyo, Japan

Ecology & Environment

*25P-234 Mutual Reinforcement Between Spatial Structure and Species Coexistence in a Living Soil Model

<u>Riz Fernando Noronha</u>, Kim Sneppen, Kunihiko Kaneko Niels Bohr Institute, Copenhagen, Denmark

Nonequilibrium state & Biological rhythm

*25P-235 Mechanism of scaling behavior of an intracellular reaction-diffusion wave in cell-size space Sakura Takada, Shunshi Kohyama, Natsuhiko Yoshinaga, Nobuhide Doi, Kei Fujiwara

Dept. of Biosci. and Info., Keio Univ., Yokohama, Japan

25P-236 Chaotic Oscillations of Sarcomeres within Cardiomyocytes Induced by Calcium Fluctuations: Identification and Physiological Significance of 'S4C'

Seine A. Shintani

Department of Biomedical Sciences, College of Life and Health Sciences, Chubu University/Center for Mathematical Science and Artificial Intelligence, Chubu University/Institute for Advanced Research, Nagoya University, Nagoya

Measurements

*25P-237 Quantitative correspondence between drug-response curves in the REMA test measured fluoromerically and colourimetrically <u>Alexander V. Sychev</u>, Anastasia Lavrova, Eugene Postnikov Kursk State University

*25P-238 Development of a dual-luciferase indicator for 'Mix-and-read' detection of Cu2+

<u>Ti Wu</u>, Mitsuru Hattori, Takeharu Nagai SANKEN, Osaka University, Japan/Graduate school of Pharmaceutical Scinences, Osaka University, Japan

*25P-239 Combined analysis of static and dynamic cell-mechanics with unbiased transcriptomics for thousands of single cells <u>Akifumi Shiomi</u>, Taikopaul Kaneko, Kaori Nishikawa, Dino Di Carlo, Hirofumi Shintaku

Cluster for Pioneering Research, RIKEN, Japan/Department of Bioengineering, University of California, USA

25P-240 Nanoendoscopy-AFM measurement of nuclear stiffness in living different metastatic cancer cells

<u>Takehiko Ichikawa</u>, Kundan Sivashanmugan, Takeshi Shimi, Kojiro Ishibashi, Takeshi Yoshida, Akiko Kudo, Eishu Hirata, Rikinari Hanayama, Hiroshi Kimura, Takeshi Fukuma Nano Life Science Institute (WPI-NanoLSI)

25P-241 Enzyme reaction measurement using graphene biosensors and its application to SARS-CoV-2 detection

Takao Ono, Yohei Watanabe, Shin-ichi Nakakita, Yasushi Kanai, Naruto Miyakawa, Ayumi Shinagawa, Shota Ushiba, Shinsuke Tani, Yasuo Suzuki, Masahiko Kimura, Daichi Chiba, Kazuhiko Matsumoto SANKEN, Osaka Univ., Osaka, Japan

25P-242 Construction 4 channels polarization-dependent fluorescence correlation spectroscopy for detection of protein interaction. <u>Masataka Kinjo</u>, Riku Ando, Akira Kitamura Hokkaido University, Sapporo, Japan.

Bioimaging

*25P-243 Development of luminescent glucose sensor and its application <u>Tanaka Rikuto</u>, Sugiura Kazunori, Hattori Mitsuru, Nagai Takeharu Graduate School of Frontier Biosciences, Osaka University

*25P-244	Quantitative chemical and physical imaging of heterochromatin in a living cell using Raman-Brillouin microscopy <u>Masato Machida</u> , Atsushi Shibata, Kentaro Fujii, Shinji Kajimoto, Takakazu Nakabayashi Graduate School of Pharmaceutical Sciences, Tohoku University
*25P-245	Investigating the Mechanical Properties and Dynamics of Focal Adhesions in Living Cells by Nanoendoscopy-AFM Technique Alam Mohammad Shahidul, Tetsuya Shirokawa, Takehiko Ichikawa, Clemens M. Franz, Takeshi Fukuma Nano Life Sicence Institute, Kanazawa University
*25P-246	Native molecular properties of full-length SARS-CoV-2 Open Reading Frame 6 (ORF6) protein observed using HS-AFM Goro Nishide, Keesiang Lim, Maiki Tamura, Akiko Kobayashi, Qingci Zhao, Masaharu Hazawa, Toshio Ando, Noritaka Nishida, Richard W. Wong Division of Nano Life Science in the Graduate School of Frontier Science Initiative, WISE Program for Nano-Precision Medicine, Science, and Technology Kanazawa University, Kanazawa, Japan
*25P-247	Shannon entropy and complexity in describing and visualizing the chemical diversity of surrounding cells by mass spectrometry imaging techniques Lili Xu, Manabu Machida, Tomoaki Kahyo, Mitsutoshi Setou Hamamatsu University School of Medicine, Hamamatsu, Japan
*25P-248	Label-free detection of supersulfides ina living cell using Raman microscopy <u>Keisuke Koga</u> , Shinji Kajimoto, Shinya Tahara, Tomohiro Konno, Takakazu Nakabayashi Graduate School of Pharmaceutical Sciences, Tohoku University
*25P-249	Mechanical properties of human platelets in biochemical confinement Vincent Gidlund, Jan Seifert, Johanna Rodriguez, Carmela Rianna, Tilman E. Schäffer Institute of Applied Physics, University of Tübingen, Tübingen, Germany

 *25P-250 Oblique Line Scan Illumination Enables Expansive, Accurate and Sensitive Single Protein Measurements in Solution and in Living Cells <u>Amine Driouchi</u>, Mason Bretan, Brynmor Davis, Alec Heckert, Markus Seeger, Maité Bradley Silva, William Forrest, Jessica Hsiung, Jiongyi Tan, Hongli Yang, Eric Betzig, Xavier Darzacq, Russ Berman, Daniel Anderson Eikon Therapeutics
 *25P-251 Characterization of a novel membrane voltage sensor in the bacterial flagellar type III export apparatus

Sakata Kai, Minamino Tohru, Morimoto Yusuke Grad. Sch. Comp. Sci. and Sys. Eng., Kyushu Inst. Tech., Fukuoka, Japan

- 25P-252 Atomic force microscopy (AFM)-based nanoindentation of the RSJ2 Ralstonia phage Udom Sae-Ueng, Chooseel Bunsuwansakul, Namthip Phironrit, Christian Nehls National Science and Technology Development Agency, Pathum Thani, Thailand
- 25P-254
 Scanning-free functional Fluorescence Microscopy Imaging Toward Spatial Mapping of Biomolecular Information in Live Cell

 Sho Oasa, Aleksandar Krmpot, Stanko Nikolic, Andrew Clayton, Igor Tsigelny, Jean-Pierre Changeux, Lars Terenius, Milivoj Belic, Rudolf Rigler, Vladana Vukojevic

 Department of Clinical Neuroscience (CNS), Karolinska Institutet, Stockholm, Sweden
- 25P-255 Characteristics of extracellular collagen in cartilage revealed by polarization-resolved second harmonic generation imaging Ming-Xin Lee Institute of Translational Medicine and New Drug Development

Bioengineering

*25P-256 Exploring Biological Changes in Whole and Serum Blood of Healthy and Diabetic Patients Using Drying Droplets Anusuya Pal, Amalesh Gope, Miho Yanagisawa Graduate School of Arts and Sciences, The University of Tokyo, Tokyo, Japan

*25P-257 Fabrication of a Nanobody-based Ratiometric Bioluminescent Immunosensor for Point-of-care Testing <u>Yinghui Yang</u>, Akihito Inoue, Takanobu Yasuda, Hiroshi Ueda, Bo Zhu, Tetsuya Kitaguchi Graduate School of Life Science and Technology, Tokyo Institute of Technology, Kanagawa, Japan

- *25P-258 High-throughput nano/micro biological particle analyzer with unsupervised denoising for enhanced sensitivity Yuichiro Iwamoto, Benjamin Salmon, Yusuke Yoshioka, Bin Xu, Ryosuke Kojima, Alexander Krull, Sadao Ota The University of Tokyo, Tokyo, Japan
- *25P-259 Multicolor autonomous bioluminescence imaging based on bacterial bioluminescence system Subhan Hadi Kusuma, Mitsuru Hattori, Takeharu Nagai Graduate School of Frontier Bioscience, Osaka University, Japan/SANKEN, Osaka University, Japan
- 25P-260 Isolation of novel fluorogenic RNA aptamers via affinity- and fluorescence-based in vitro selection

<u>Ryo lizuka</u>, Keisuke Ito, Tomotaka Tayama, Sotaro Uemura Department of Biological Sciences, Graduate School of Science, The University of Tokyo

Crystal growth & Crystallization technique

- *25P-261 Protein calixarene crystal engineering <u>Niamh Maria Mockler</u>, Kiefer Ramberg, Colin Raston, Peter Crowley School of Biological and Chemical Sciences, University of Galway, H91 TK33, Galway, Ireland
- *25P-262 Investigation of crystallization of crystallized protein expressed using E. coil Yume Kosuge, Koki Kamiya

Graduate School of Science and Technology, Gunma University, Kiryu, Gunma, Japan

Virus structure, function, SARS-CoV-2

*25P-263 Study of the binding site dynamics, druggability and cryptic pocket formation in different human coronaviruses' main protease (Mpro) Ahmed Adel Ezat Biophysics Department, Faculty of Science, Cairo University, 21613 Giza, Egypt

*25P-264 Cryo-EM structure of the Borna disease virus 1 RNA-free nucleoprotein complex

Shinya Goto, Yuya Hirai, Keizo Tomonaga, Takeshi Noda, Masayuki Horie, Yukihiko Sugita Laboratory of Ultrastructural Virology, Institute for Life and Medical Sciences, Kyoto University/Laboratory of Ultrastructural Virology, Graduate School of Biostudy, Kyoto University

25P-266 Unraveling the Dynamics of SARS-CoV-2 Spike: From Glycosylation States to Cryptic Pockets and Antibody Binding

<u>Mohd Firdaus Samsudin</u>, Lorena Zuzic, Palur Raghuvamsi, Aishwary Shivgan, Nikhil Tulsian, Himanshi Chawla, Joel Allen, Max Crispin, Paul MacAry, Ganesh Anand, Peter Bond Bioinformatics Institute, A*STAR, Singapore

Mechanosensing and Mechanobiology, Biological Temperature

*25P-267 Development of a high-frequency focused ultrasound system for applying noninvasively localized mechanical stimulation to cells in culture

 Natsumi Fujiwara, Shao Ying Tan, Akira Nagakubo, Masahiro Kino-oka, Hirotsugu Ogi
 Graduate School of Engineering, Osaka University, Japan

 25P-268 Force transmission by retrograde actin flow-induced dynamic stretching of Talin

 Sawako Yamashiro, David Rutkowski, Kelli Ann Lynch, Ying Liu,

Dimitrios Vavylonis, Naoki Watanabe Laboratory of Single-Molecule Cell Biology, Kyoto University Graduate School of Biostudies, Kyoto, Japan/Department of Pharmacology, Kyoto University Graduate School of Medicine, Kyoto, Japan

25P-269 Modulating E-Cadherin Engagement to Alter Cell Junctional Tension in Spheroids

<u>Seongho Kim</u>, Isaac T.S. Li Department of Chemistry, The University of British Columbia, Canada

Biophysics of disease

*25P-270 Opportunities of Raman spectroscopy in pulmonary arterial hypertension

Elvin Suleyman oglu Allakhverdiev, Olga Slatinskaya, Oleg Rodnenkov, Tamila Martynyuk, Georgy Maksimov National Medical Research Center of Cardiology named after academician E.I. Chazov of Ministry of Health of the Russian Federation

*25P-271 Exploring Dapagliflozin Therapy Effects on Nanomechanics and Morphology of Red Blood Cells in Type I Diabetes Mellitus

Patrycja Lidia Twardawa, Bartłomiej Matejko, Agata Kubisiak, Katarzyna Cyranka, Tomasz Klupa, Marta Targosz-Korecka Jagiellonian University, Faculty of Physics, Astronomy and Applied Computer Science, M. Smoluchowski Institute of Physics, Kraków, Poland/Jagiellonian University, Doctoral School of Exact and Natural Sciences, Kraków, Poland

25P-272 Opposite effects of extracellular chloride and pH on closely related CIC-6 and CIC-7 transporters suggest non-overlapping function in endo-lysosomes

Maria Antonietta Coppola, Paola Gavazzo, Ilaria Zanardi, Abraham Tettey-Matey, Antonella Liantonio, Paola Imbrici, Peying Fong, <u>Michael Pusch</u> Institute of Biophysics, CNR, Genoa, Italy

Miscellaneous topics

*25P-273 Structural and Magneto Absorption Study of Hard and Soft Ferrite Usha Praveena V J Department of Physics, St. Francis College for Women, Hyderabad-500 016, Telangana, India

25P-274 FUNCTIONALIZED CNT AND ACTIVATED CARBON

Shikha Chander, Meenu Mangal

St. Francis Degree & Postgraduate Women's College, Begumpet 500016, Hyderabad, Telangana, India

Single Molecule Biophysics

25P-275 Quantifying ligand binding kinetics in G-quadruplex DNA with fluorescence lifetime correlation analyses

<u>Chao-Han Cheng</u>, Chih-Chieh Ko, Yong-Zhan Hong, Chung-Chieh Wu Department of Applied Chemistry, National Pingtung University, Pingtung, Taiwan

Cell biology: Motility

25P-276 Inference of cellular traction forces using temporal information Kazuko Hamaoka, Hirokazu Tanimoto Grad. Sch. Nanobioscience, Yokohama City Univ.

Computational biology: Molecular simulation

25P-277 Secondary Proton Transfer in the Qo Site of Cytochrome bc1 <u>Guilherme M. Arantes</u>, Sofia Camilo Department of Biochemistry, Instituto de Quimica, Universidade de Sao Paulo, Brazil

Computational biology: Biological modeling and simulation

- 25P-278 Phase-field model of Dictyostelium fruiting body formation Seiya Nishikawa, Satoshi Kuwana, Gen Honda, Satoshi Sawai, Shuji Ishihara Graduate School of Arts and Sciences, University of Tokyo
 25P-279 Topography-mediated cell communication
- Aleksandra Ardaševa, Varun Venkatesh, Daiki Matsunaga, Shinji Deguchi, Amin Doostmohammadi Niels Bohr Institute, University of Copenhagen, Denmark

Mathematical & Theoretical biology

25P-280 Latecomer Killing: Elaborate Response in Yeast Communities <u>Tetsuhiro S. Hatakeyama</u>, Kunihiko Kaneko, Kunihiro Ohta, Miki Tamura, Arisa Oda Earth Life Science Institute (ELSI), Tokyo Institute of Technology

Biophysics of disease

25P-281 Reinstating heart rate variability improves cardiac output in heart failure - novel insights from proteomics

David Crossman, George Guo, Julia Shanks, Jizhong Bai, Martin Middleditch, Gus Grey, Julian Paton, Rohit Ramchandra Manaaki Manawa—The Centre for Heart Research, Department of Physiology, University of Auckland, 85 Park Road, Grafton, Auckland, 1023, New Zealand

25P-282 Development of the two-fingered microhand and micro fluidic system for measuring the mechanical properties of cell

Masaru Kojima, Masahiro Totani, Masahiro Kawakami, Toshihiko Ogura, Tatsuo Arai

Osaka University

25P-283 Targeting Retinal Angiogenesis: Potential of AT11-L0 Aptamer for Drug Delivery

David Moreira, Jessica Lopes-Nunes, Fátima Santos, Maria Oliveira, António Paulo, Maria Campello, Carla Cruz, <u>Cândida Ascensão Teixeira Tomaz</u>

Departamento de Química, Universidade da Beira Interior, Covilhã, Portugal/CICS-UBI-Health Sciences Research Centre, Universidade da Beira Interior, Covilhã, Portugal