

# 3rd International Conference on Molecular Simulation (ICMS2013)

The date of presentation is given at the first number.

Underlined is the presenter among more than one authors.

Candidates for student award is marked with ☆ in front of the poster title.

## Day 1 Nov. 18

9:00–9:10

**Opening Address by Conference Chair**

Shiaki Hyodo

**President of the Molecular Simulation Society of Japan**

Susumu Okazaki

*chair: Susumu Okazaki*

9:10 **PL1** **Structure, Dynamics, and Thermodynamic Stability of High Pressure Ices and Clathrate Hydrates**

Hideki Tanaka

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9 : 55 break

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10:10–11:40 **Oral**

*chair: Pavel Jungwirth*

10:10 **1O01** **Low-density liquid water is the mother of ice: on the relation between liquid structure, thermodynamics and ice crystallization**

Valeria Molinero

10:40 **1O02** **Molecular Simulation of Confined Water and Aqueous Solution**

Kenji Yasuoka

11:10 **1O03** **Structure and Dynamics of Nanoconfined Liquids**

Ward H. Thompson

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11 : 40 break

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11:55–12:55 **Oral**

*chair: Kenji Yasuoka*

11:55 **1O04** **Molecular Simulation for Liquid Interfaces**

Akihiro Morita

12:25 **1O05** **Motion of H<sup>+</sup> in H-bond Network: From Quantum Confinement to Fermi Resonance**

Jake Tan, Chen-Wei Tsai, Jer-Lai Kuo

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12 : 55 lunch

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13:45–14:45 **Poster**

14:45–16:15 **Oral**

*chair: Jeffery B. Klauda*

14:45 **1006** Interactions Between Ions and Proteins in Water: The Hofmeister Series

Pavel Jungwirth

15:15 **1007** Urea-induced denaturation of PreQ<sub>1</sub>-riboswitch

Changbong Hyeon

15:45 **1008** Biomolecular Simulations under Cellular Crowding Environment

Yuji Sugita, Ryuhei Harada, Isseki Yu, Takaharu Mori, Jaewoon Jung, Michael Feig

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16 : 15 break

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16:30–18:00 **Oral**

*chair: Wataru Shinoda*

16:30 **1009** Free-Energy Analysis of Water Affinity in Polymer Studied by Atomistic MD and the Energy-Representation Method

Tomonori Kawakami, Masahiro Kitabata, Isamu Shigemoto

17:00 **1010** Force Fields and Simulation Protocols in Predictions of Adsorption

Huai Sun

17:30 **1011** Free-Energy Analysis of Solvation and Membrane Effects on Protein Configurations

Nobuyuki Matubayasi

## Day 2 Nov. 19

*chair: Yuko Okamoto*

9:00 **PL2** Capturing pH mediated physics and chemistry in biological systems

Charles L. Brooks III

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9 : 45 break

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10:00–11:30 **Oral**

*chair: Mitsunori Ikeguchi*

10:00 **2001** Large-Scale and All-Atom Molecular Dynamics Simulation of Viruses Using the K Computer

Susumu Okazaki

10:30 **2002** Simulations of Biomembranes: Importance of Lipid Diversity and Structural Changes of Membrane Transport Proteins

Jeffery B. Klauda

11:00 **2003** Structure and function of the transmembrane domain of Amyloid Precursor Protein and its role in the amyloid  $\beta$  protein aggregation pathway

Laura Dominguez, Leigh Foster, D. Thirumalai, John E. Straub

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11 : 30 break

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**11:45–12:45 Oral**

*chair: Changbong Hyeon*

**11:45 2004 Molecular recognition by intrinsically disordered proteins – insights from molecular simulation**

David de Sancho, Michael Knott, Christopher M. Baker, Robert B. Best

**12:15 2005 Dynamics of Neuraminidase: from Hamiltonian REMD to novel inhibitor design**

Nanyu Han, Yuguang Mu

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12 : 45 lunch

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**13:45–14:45 Poster**

**14:45–16:15 Oral**

*chair: Yuji Sugita*

**14:45 2006 Combination of MD and SAXS to Explore Protein Dynamics**

Mitsunori Ikeguchi

**15:15 2007 Molecular simulation of protein folding, conformational change and self-assembly**

Peter G. Bolhuis

**15:45 2008 Generalized-ensemble algorithms for molecular simulations**

Yuko Okamoto

**16:30 Excursion Visit to K Computer**

**18:45 Banquet with Bay Cruise (Transportation bus departs at 18:00)**

**Day 3 Nov. 20**

*chair: Shiaki Hyodo*

**9:00 PL3 Chemical Reaction Rates from Ring Polymer Molecular Dynamics**

David E. Manolopoulos

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9 : 45 break

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**10:00–11:30 Oral**

*chair: Akihiro Morita*

**10:00 3001 Variational path integral molecular dynamics with applications to molecular systems**

Shinichi Miura

**10:30 3002 Learning chemistry from multiple first-principles simulations**

Giovanni La Penna

**11:00 3003 Molecular Statistical Mechanics of Chemical Reactions in Solution Phase**

Hirofumi Sato

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11 : 30 break

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**11:45–12:45 Oral**

*chair: Ward H. Thompson*

**11:45 3004 Crucial Role of Protein Flexibility in Enzymatic Catalysis**  
Shigehiko Hayashi

**12:15 3005 Massively Parallel Implementation of the QM/MM Approach Combined with a Theory of Solutions**  
Hideaki Takahashi, Yuji Miki, Akihiro Morita

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12 : 45 lunch

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**13:45–14:45 Poster**

**14:45–16:45 Oral**

*chair: Mitsuhiro Matsumoto*

**14:45 3006 Smoothed Profile Method for DNS of Particle Dispersions**  
Ryoichi Yamamoto, John J. Molina

**15:15 3007 Balance of Enthalpy and Entropy in Depletion Forces**  
Liel Sapir, Shahar Sukenik, Daniel Harries

**15:45 3008 Mechanical stability of proteins: from coarse-grained to all-atom simulations**  
Mai Suan Li, Maksim Kouza

**16:15 3009 Free Energy Analysis of Topological and Morphological Changes in Lipid Membranes: Molecular Dynamics Study**  
Wataru Shinoda, Shuhei Kawamoto

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16 : 45 break

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*chair: Nobuyuki Matubayasi*

**16:55 MSSJ Award Lecture: Effective sampling algorithms and analysis methods for protein simulations**  
Ayori Mitsutake

**17:30 Closing Remarks**  
Shiaki Hyodo

**17:35 Business Meeting of the Molecular Simulation Society of Japan (in Japanese)**

**Poster day 1**

**1P01 Coupled-Cluster Calculations on Electronic Circular Dichroism of Single and Double Helicenes**  
Ikenosako Mina, Takeharu Kusuki, Yoshito Nakai, Yoshihisa Inoue, Tadashi Mori

**1P02 Fluctuation Flooding Method (FFM) for enhancing conformational sampling of proteins**  
Ryuhei Harada, Yu Takano, Yasuteru Shigeta

**1P03 Multiscale enhanced sampling and its applications to large solvated proteins**  
Kei Moritsugu, Tohru Terada, Akinori Kidera

- 1P04 Drug export process of MATE multidrug efflux transporter by Molecular dynamics simulations**  
Wataru Nishima, Yoshiki Tanaka, Ryuichiro Ishitani, Osamu Nureki, Yuji Sugita
- 1P05 Effects of Initial Settings on Computational Protein-Ligand Docking Accuracies for several Docking Programs**  
Akifumi Oda, Noriyuki Yamaotsu, Shuichi Hirono, Yurie Watanabe, Shuichi Fukuyoshi, Ryoichi Nakagaki, Ohgi Takahashi
- 1P06 Logarithmic Mean-Force Dynamics (LogMFD): Application to a Model System of Protein-G**  
Tetsuya Morishita
- 1P07 Molecular Dynamics Simulation Underlying the Early Stage of the Channelrhodopsin**  
Mizuki Takemoto, Hideaki E. Kato, Michio Koyama, Jumpei Ito, Motoshi Kamiya, Shigehiko Hayashi, Andrés D. Maturana, Karl Deisseroth, Ryuichiro Ishitani, Osamu Nureki
- 1P08 Study on the C-terminal domain of eukaryotic RNA polymerase II by using a molecular dynamics simulation**  
Yasushige Yonezawa
- 1P09 Prediction of Protein-ligand Binding Affinities Using Molecular Simulations**  
Noriaki Okimoto, Atsushi Suenaga, Yoshinori Hirano, Makoto Taiji
- 1P10** ☆ **Conversion of CO<sub>2</sub>, CH<sub>4</sub> and C<sub>2</sub>H<sub>6</sub> to Organic Compound on Ir-Graphene Oxide Surface - A First-Principle Investigation**  
Chih-Chun Chen, Jia-Jen Ho
- 1P11** ☆ **Molecular Dynamics Study on Simplified Model Membranes for Normal Murine Thymocytes and Leukemic GRSL Cell Plasma Membranes**  
Soichiro Shibayama, Yoshimichi Andoh, Susumu Okazaki
- 1P12 Conformational analysis of PA-glycans by replica-exchange molecular dynamics simulation.**  
Shigehisa Watabe, Suyong Re, Wataru Nishima, Eiro Muneyuki, Yuji Sugita
- 1P13 Molecular Dynamics Simulation of Enzyme-Substrate Analogue Complex of Heterotetrameric Sarcosine Oxidase**  
Go Watanabe, Akinori Hiroshima, Haruo Suzuki, Shigetaka Yoneda
- 1P14** ☆ **Electrostatic steering mechanism in selective binding of KIF1A on microtubule**  
Yukinobu Mizuhara, Jun Ohnuki, Koji Umezawa, Takano Mitsunori
- 1P15** ☆ **Piezoelectricity of protein and its implication for function**  
Jun Ohnuki, Takato Sato, Koji Umezawa, Taro Q. P. Uyeda, Mitsunori Takano
- 1P16 Nature of Proton Transport in Hydrocarbon Membranes for Fuel Cell Applications**  
Yoong-Kee Choe, Eiji Tsuchida, Tamio Ikeshoji, Akihiro Ohira, Takahiro Ohkubo, Neil Henson, Yu Seung Kim
- 1P17 Non-equilibrium Molecular Dynamics Simulations for Amyloid Disruption by Hypersonic Wave**  
Hisashi Okumura, Satoru G. Itoh
- 1P18 Large scale molecular dynamics simulations of vapour-to-liquid nucleation**  
Jürg Diemand, Raymond Angelil, Kyoko K Tanaka, Hidekazu Tanaka

- 1P19** Cation- $\pi$  vs. Water- $\pi$  : Theoretical analysis of structural stability of the active center of TI lipase employing a novel accurate and efficient description of effective potential  
Atsushi Nakamura, Jiyoung kang, Masaru Tateno
- 1P20** A reaction analysis for deprotonation of sulfonic group of perfluorosulfonic acid molecule in the low hydration level by density functional theory  
Hironori Sakai, Takashi Tokumasu
- 1P21** Multiple length and time scales of dynamic heterogeneities in fragile and strong glass-forming liquids  
Kang Kim, Shinji Saito
- 1P22** Molecular Dynamics Simulation for Phase Behavior of Bolaamphiphilic Solution  
Susumu Fujiwara, Takumi Miyata, Takashi Itoh, Masato Hashimoto
- 1P23** Theoretical studies on the stability of clathrate hydrates  
Takato Nakayama, Masakazu Matsumoto, Hideki Tanaka
- 1P24** Wetting Transition of Ethanol-Water Droplet on Smooth and Textured Surfaces  
Atanu K. Metya, Jayant K Singh
- 1P25** Phase Transitions of Water Confined in Slit Pores  
Toshihiro Kaneko, Jaeil Bai, Kenji Yasuoka, Ayori Mitsutake, Xiao Cheng Zeng
- 1P26** The channel size dependence of the friction coefficient between the water droplet and the solid wall  
Akinori Fukushima, Toshiki Mima, Ikuya Kinefuchi, Takashi Tokmasu
- 1P27** ☆Molecular Dynamics Simulation for Melting Process of Various types of Core-Shell Nanoparticles  
Yuki Tamura, Noriyoshi Arai
- 1P28** Molecular Simulation Study of Structure and Dynamical Properties of Nitrate Anion in Sodium Chloride Aqueous Solution  
Shigeki Matsunaga
- 1P29** ☆Application of a Liquid Theory to Colloidal Dispersion Systems  
Masao Inoue, Akira Yoshimori
- 1P30** Dynamic Simulations of Ion Atmosphere of Complex Electrolytes Solutions  
Hitoshi Washizu, Tomoyuki Kinjo, Hiroaki Yoshida
- 1P31** A VT Replica Exchange Molecular Dynamics Study in Ultra Supercooled TIP4P/2005 Water  
Liang-Chun Liu, Jer-Lai Kuo
- 1P32** Molecular dynamics simulation of vapour-liquid nucleation of water: determining formation energy of a cluster  
Akio Kawano, Kyoko K. Tanaka, Hidekazu Tanaka
- 1P33** ☆All-atom Analysis of Electrical Conductivity in Concentrated Electrolyte Solutions  
Kai-Min Tu, Nobuyuki Matubayasi
- 1P34** ☆Development of Ab-initio Mean-force Dynamics Using a Logarithmic Energy Landscape for Constructing Free-Energy Profiles  
Makoto Nakamura, Masao Obata, Tetsuya Morishita, Tatsuki Oda
- 1P35** Hybrid quantum mechanics (QM) / molecular mechanics (MM) molecular dynamics (MD) simulation of catalytic mechanism by complex of Ras and Ras-GAP: Full description of catalytic mechanisms  
Masaru Tateno, Tetsuhiko Itagaki, Jiyoung Kang

- 1P36** ☆ **A variational path integral molecular dynamics study of molecules using a fourth order propagator**  
Yuki Kamibayashi, Shin-ichi Miura
- 1P37** **Analyses of Coarse-Grained Equation of Motion**  
 Shiaki Hyodo
- 1P38** ☆ **Molecular Dynamics Simulations of Surfactant Interfaces Using Coarse-Grained Force Field**  
Chunwei Yang, Huai Sun
- 1P39** ☆ **Nuclear quantum effect on  $\text{OH}^-(\text{H}_2\text{O})_2$  with *ab initio* path integral molecular dynamics**  
Yudai Ogata, Yukio Kawashima, Kaito Takahashi, Masanori Tachikawa
- 1P40** **A Theory of Phase Transition for Two-Minimum Potential Systems**  
 Ayumi Suematsu, Akira Yoshimori, Masafumi Saiki, Jun Matsui, Takashi Odagaki
- 1P41** **A Boundary Condition for Multilevel Simulations for Diffusive Systems**  
Motoyuki Shiga, Marco Masia
- 1P42** **Direct Formation of “Janus”-Particles via Molecular Dynamics Simulations**  
Jan-Hubert Antonio Wittmann, Reinhard Strey
- 1P43** ☆ **Molecular Dynamics Simulation on Complexation of a Charged Dendrimer with a Linear Polyelectrolyte**  
Satoshi Isono, Akira Kazuno, Susumu Fujiwara, Takashi Itoh, Masato Hashimoto
- 1P44** **Molecular dynamics simulation for shape change of water-in-oil droplets**  
Naohito Urakami, Akio Takaki, Masayuki Imai, Takashi Yamamoto
- 1P45** ☆ **Gelation of tetrahedral monomers studied by molecular dynamics simulation**  
Shizuka Ogawa, Takeshi Miyakawa, Ryota Morikawa, Masako Takasu, Takamasa Sakai, Ung-il Chung
- 1P46** **Monte Carlo Simulations of Structure and Entanglements in Polymer Melts**  
Krzysztof Moorthi, Kazunori Kamio, Javier Ramos, Doros N Theodorou
- 1P47** **Band Gap Design of Thiophene Polymer Based on The Density Functional Theory**  
Patricia Lubis, Mineo Saito
- 1P48** **A numerical simulation of Brownian-particle suspensions using the lattice Boltzmann method**  
Hiroaki Yoshida, Tomoyuki Kinjo, Hitoshi Washizu
- 1P49** **Structural Analysis of Telechelic Polymer Solutions -A Dissipative Particle Dynamics Study-**  
 Noriyoshi Arai
- 1P50** **Tetratic phase of bilayer systems under geometrical confinement: Monte Carlo simulation**  
 Takamichi Terao
- 1P51** ☆ **Thermal conductivity in binary and ternary mixed system of molten alkali halides**  
Yoshiki Ishii, Keisuke Sato, Norikazu Ohtori, Mathieu Salanne, Paul Anthony Madden
- 1P52** ☆ **The dependence of ordering process on Coulombic interaction treatment 5CB system**  
Takuma Nozawa, Kazuaki Takahashi, Tetsu Narumi, Kenji Yasuoka
- 1P53** **First principles calculation of the PCBM on the  $\text{TiO}_2$  anatase (001) surface**  
Kiyotaka Fujino, Mineo Saito

- 1P54** **Molecular Photovoltaic Devices and Density-functional Theory Simulation**  
Shozo Yanagida, Kazuhiro Manseki
- 1P55** **First-Principles Calculation of Surface State in Topological Insulators Bi<sub>2</sub>Te<sub>3</sub> and Related Materials: Film Thickness Dependence**  
Takehiro Kato, Hiroki Kotaka, Fumiyuki Ishii
- 1P56** ☆ **Nitrogen Adsorption and Dissociation on Double-Icosahedral Ru<sub>19</sub> nanocluster - A First-Principle Investigation**  
Chen-Hao Yeh, Jia-Jen Ho
- 1P57** **The Highly Effective Catalytic Behavior of a Metal Nanocluster Ru<sub>79</sub> on the Dissociation of a N<sub>2</sub> Molecule - A Quantum-chemical Calculation**  
Shiuan-Yau Wu, Chen-Hao Yeh, Jia-Jen Ho
- 1P58** **Magnetic Dipole-Dipole Interaction and Magnetic Structure in Solid Oxygen**  
Dewi Mulyati, Masao Obata, Makoto Nakamura, Suprijadi Haryono, Tatsuki Oda

## Poster day 2

- 2P01** ☆ **Study of Water-Methanol Mixture Flowing into Carbon Nanotubes with Molecular Dynamics Simulation: Effect of Electric Field**  
Winarto, Daisuke Takaiwa, Eiji Yamamoto, Takuma Akimoto, Kenji Yasuoka
- 2P02** ☆ **Study of aqueous NaOH solution surface via computational analysis of sum frequency spectra**  
Takako Imamura, Tatsuya Ishiyama, Akihiro Morita
- 2P03** **Quantum simulation for exotic molecules: Quantum Monte Carlo and Path Integral approach**  
Katsuhiko Koyanagi, Yukiumi Kita, Kenta Yamada, Yukio Kawashima, Masanori Tachikawa
- 2P04** **Vibrational Spectroscopic Response at Water/Vapor and Ice/Vapor Interfaces : Effect of Charge Transfer**  
Tatsuya Ishiyama, Hideaki Takahashi, Akihiro Morita
- 2P05** ☆ **Solubility of hydrophobic molecules at the liquid-vapor interfaces of water and simple liquids**  
Kiharu Abe, Kenichiro Koga
- 2P06** **Free energy analysis of CO<sub>2</sub> dissolved in room temperature ionic liquids**  
Ryosuke Ishizuka, Nobuyuki Matubayasi
- 2P07** ☆ **Pore Formation Dynamics in Phospholipid/Cholesterol Bilayer under Stretching: System Size Dependence**  
Taiki Shigematsu, Kenichiro Koshiyama, Shigeo Wada
- 2P08** **System size dependence of vapor/liquid coexistence of water in hydrophobic nanocylinder**  
Toshiki Mima, Ikuya Kinefuchi, Akinori Fukushima, Takashi Tokumasu, Shu Takagi, Yoichiro Matsumoto
- 2P09** **Molecular Dynamics Simulation of L-J Fluids near a Heating Surface**  
Kazuhiro Itoh, Hiroshige Kumamaru, Yuji Shimogonya
- 2P10** **The observation of the longitudinal wave velocity in a model supercooled liquid**  
Tadashi Muranaka, Jun Matsui, Yasuaki Hiwatari



- 2P11** Distance Constraint Networks in Glass-Forming Liquids  
Yasushi Takeuchi
- 2P12** Hydrophobic Cavity in the Potassium Channels: Anomalous Nanospace Controlling Ion Conduction  
Takashi Sumikama, Shinji Saito, Shigetoshi Oiki
- 2P13** Diffusion of Ions in the Salt Solutions between Ice Walls  
Yuta Nohara, Yoshimichi Hagiwara
- 2P14** Glass Transition in One-Component Soft-Core System  
Junko Habasaki, Akira Ueda
- 2P15** ☆ Guest-host hydrogen bond in sII and sH amyloid-alcohol hydrate from molecular dynamics simulation  
Masaki Hiratsuka, Ryo Ohmura, Amadeu K Sum, Kenji Yasuoka
- 2P16** A new scaling relation for nucleation rates  
Kyoko K. Tanaka, Hidekazu Tanaka, Juerg Diemand, Raymond Angelil
- 2P17** Relaxation Mode Analysis for Biomolecules  
Ayori Mitsutake, Yuta Koizumi, Toshiki Nagai, Hiroshi Takano
- 2P18** Evaluation of Protein-Protein Complex Model Using Molecular Dynamics Simulation with the Solution Theory in the Energy Representation  
Kazuhiro Takemura, Nobuyuki Matubayasi, Akio Kitao
- 2P19** Free energy calculation of protein conformational changes using parallel cascade selection molecular dynamics simulation and Markov state model  
Yasutaka Nishihara, Ryuhei Harada, Akio Kitao
- 2P20** Free energy analysis of flip-flop motion of protein in phospholipid bilayer  
Tomoko Mizuguchi, Nobuyuki Matubayasi
- 2P21** Hybrid QM/MM simulation of bovine cytochrome c oxidase (I) : Double-stratified architecture for electron transfer  
Jiyoung Kang, Masaru Tateno
- 2P22** The Interaction Between Transmembrane Region of Amyloid Precursor Protein (APP) and Cholesterols.  
Naoyuki Miyashita, Fumiko Ogushi, Yuji Sugita
- 2P23** Difficulty of Observation of Attractive Part in Effective Interaction between Like-Charged Proteins: a Theoretical Study with Simple Models  
Ryo Akiyama, Takumi Yamashita
- 2P24** A Molecular Dynamics Study on the Antigen-Antibody Interactions  
Takefumi Yamashita, Hideaki Fujitani
- 2P25** A model-potential-free analysis of protein-protein interactions using small-angle X-ray scattering  
Tomonari Sumi, Hiroshi Imamura, Takeshi Morita, Yasuhiro Isogai, Keiko Nishikawa
- 2P26** Computational study of protein-ligand interactions: Molecular screening for therapeutic drugs  
Yosinori Hirano, Noriaki Okimoto, Makoto Taiji
- 2P27** ☆ Theoretical Study for Understanding the Mechanism of Amyloid Fibril Formations  
Naohiro Nishikawa, Phuong Nguyen, Philippe Derreumaux, Yuko Okamoto

- 2P28** Large-scale and All-atom Molecular Dynamics Simulation of Viruses using the K-computer 2. Development of a Highly Parallelized General-Purpose Molecular Dynamics Simulation Program, MODYLAS  
Yoshimichi Andoh, Noriyuki Yoshii, Kazushi Fujimoto, Keisuke Mizutani, Hidekazu Kojima, Atsushi Yamada, Susumu Okazaki, Kazutomo Kawaguchi, Hidemi Nagao, Kensuke Iwahashi, Fumiyasu Mizutani, Kazuo Minami
- 2P29** ☆ Large-scale and all-atom molecular dynamics simulation of viruses using the K-computer. 3. Equilibration of the system and the stable structure of poliovirus capsid in solution  
Hidekazu Kojima, Noriyuki Yoshii, Atsushi Yamada, Yoshimichi Andoh, Kazushi Fujimoto, Keisuke Mizutani, Atsushi Nakagawa, Akio Nomoto, Susumu Okazaki
- 2P30** ☆ Inhomogeneous dielectric environment of F<sub>0</sub> motor using dipole-dipole correlation  
Asahi Konno, Koji Umezawa, Mitsunori Takano
- 2P31** The elucidation of the phase transition mechanism for biomechanical material application of Ti-Ni-Cu alloy  
Shuichi Ikeda, Shiaki Hyodo
- 2P32** ☆ MD simulations of ATP/ADP bound form of Ca<sup>2+</sup>-ATPase using improved polyphosphate parameters of the CHARMM force field  
Yasuaki Komuro, Suyong Re, Chigusa Kobayashi, Eiro Muneyuki, Yuji Sugita
- 2P33** New generalized ensemble algorithm without the detailed balance condition  
Satoru G. Itoh, Hisashi Okumura
- 2P34** ☆ Analysis of the structural transition of a chignolin molecule using the 3D-RISM/MD hybrid simulation method  
Shinnosuke Gyoubu, Shinichi Miura
- 2P35** An efficiently extendable and fine-grain parallelized molecular dynamics simulation program: Mid  
Takahiro Koishi, Hironori Takeich
- 2P36** A simulation method for protein systems by replica-exchange molecular dynamics using genetic crossover  
Yoshitake Sakae, Tomoyuki Hiroyasu, Mitsunori Miki, Katsuya Ishii, Yuko Okamoto
- 2P37** A 3D-RISM integral equation study of hydrated peptides  
Hiroshi Iwasaki, Shinichi Miura
- 2P38** Generalized centroid molecular dynamics  
Atsushi Horikoshi
- 2P39** ☆ Theoretical studies of redox potential for metalloprotein by combining MD and QM/MM simulations  
Masashi Iwayama, Hiroaki Saito, Kimikazu Sugimori, Kazutomo Kawaguchi, Hidemi Nagao
- 2P40** ☆ Development of a Simultaneous Simulation Method for Phonons and Electrons  
Yusuke Masao, Honami Imanishi, Mitsuhiro Matsumoto
- 2P41** ☆ Spontaneous self-assembly of diblock copolymers in nanoconfined geometries by dissipative particle dynamics  
Koji Takahashi, Takahiro Koishi
- 2P42** Bottom-up construction of interaction models of non-Markovian dissipative particle dynamics  
Yuta Yoshimoto, Ikuya Kinefuchi, Toshiki Mima, Akinori Fukushima, Takashi Tokumasu, Shu Takagi

- 2P43** Multiscale simulation for thermo-hydrodynamic lubrication of a polymeric liquid between parallel plates  
Shugo Yasuda, Ryoichi Yamamoto
- 2P44** Molecular Dynamics Simulation Study on the High Pressure Behavior of a Peptide  
Yoshiharu Mori, Hisashi Okumura
- 2P45** ☆ Heterogeneous Dynamics of Supercooled Polymer Melts: Relaxation Mode Analysis Study  
Nobuyuki Iwaoka, Hiroshi Takano
- 2P46** Thermal Stability of Gramicidin A in Lipid Bilayer: A Molecular Dynamics Study  
Hiroaki Saito, Kawaguchi Kazutomo, Hidemi Nagao
- 2P47** Coarse-grained Molecular Simulation for Viscoelastic Properties of Polyurethane Shape Memory Polymer  
Shimpei Matsuda, Masaaki Nishikawa, Masaki Hojo
- 2P48** Interaction of the nucleotide with a phospholipid bilayer: a molecular dynamics simulation study  
Daisuke Takaiwa, Kenji Yasuoka, Toshikazu Ebisuzaki
- 2P49** Relaxation modulus connection between Atomistic and Coarse-Grained Molecular Simulation of Polymer Melts using the Rouse Parameters  
Nobuyoshi Yamato, Iori Yonekawa, Kazuaki Z Takahashi, Kenji Yasuoka, Yuichi Masubuchi
- 2P50** ☆ Phase Transition between Ice VII and Plastic Ice  
Kazuhiro Himoto, Masakazu Matsumoto, Hideki Tanaka
- 2P51** First-Principles Study of Artificial Superlattice  $(\text{LaMnO}_3)_n/(\text{SrTiO}_3)_m$   
Miho Nishida, Fumiyuki Ishii, Hiroki Kotaka, Mineo Saito
- 2P52** Uncovering a Dynamically Formed Substrate Access Tunnel in Carbon Monoxide Dehydrogenase/Acetyl-CoA Synthase  
Po-hung Wang, Maurizio Bruschi, Luca De Gioia, Jochen Blumberger
- 2P53** ☆ Reactive Force Field (ReaxFF) Developed for Aluminophosphate Zeolite  
Liang Xin, Huai Sun
- 2P54** Magnetism and electronic structure calculations of Pd-TM alloys and hydrogen system  
Norio Nunomura, Masanori Hara, Satoshi Akamaru
- 2P55** Analysis for Crystal Growth of Methane Hydrate from Molecular Dynamics Simulation  
Daisuke Yuhara, Daisuke Takaiwa, Kenji Yasuoka
- 2P56** Comparative ab initio Calculation Studies of Hydration Structures on Cs and other Alkaline metals  
Masahiko Machida, Kazuhiro Sakuramoto, Masahiko Okumura, Hiroki Nakamura, Yuya Sato, Keiko Akutsu, Motoyuki Shiga
- 2P57** ☆ Molecular Dynamics Study on Carbon Diffusion Mechanism of Pearlitic Steel in Wire Drawing  
Yohei Sameshima, Ken-ichi Saitoh, Masanori Takuma, Yoshimasa Takahashi
- 2P58** ☆ DFT Study of Negatively Charged Oxygen Vacancy on (110)  $\text{TiO}_2$  Surface  
Taizo Shibuya, Kenji Yasuoka, Susanne Mirbt, Biplab Sanyal

- 2P59** Evidence of Phonon-Assisted Auger Recombination and Multiple Exciton Generation in Semiconductor Quantum Dots Revealed by Temperature-Dependent Phonon Dynamics  
Hyeon-Deuk Kim

**Poster day 3**

- 3P01** ☆Dual effect of crowders on fibrillation kinetics of polypeptide chains revealed by lattice models  
Nguyen Truong Co, Chin-Kun Hu, Mai Suan Li
- 3P02** ☆A Molecular Simulation Study to Investigate Actin Filament Elongation Mechanism  
Nobuhiko Wakai, Yasutaka Nishihara, Kazuhiro Takemura, Takashi Fujii, Keiichi Namba, Akio Kitao
- 3P03** ☆Replica exchange MD and *ab initio* fragment MO calculations on effects of solvating water molecules on stable conformations of amyloid- $\beta$  dimer  
Akisumi Okamoto, Atsushi Yano, Kazuya Nomura, Shin'ichi Higai, Noriyuki Kurita
- 3P04** MHC I Recognition Mechanism of MIR2 from Kaposi's Sarcoma-Associated Herpesvirus  
Pai-Chi Li, Mizuho Kajikawa, Naoyuki Miyashita, Satoshi Ishido, Yuji Sugita
- 3P05** Multiple molecular dynamics simulations of human adult haemoglobin: Novel intra-molecular signal propagation pathways for nonlinear response in ligand-binding  
Tetsuhiko Itagaki, Jiyoung Kang, Masaru Tateno
- 3P06** Hybrid QM/MM simulation of bovine cytochrome *c* oxidase (II): Dynamical mechanisms of ligand recognition  
Toru Matsuoka, Jiyoung Kang, Masaru Tateno
- 3P07** Standard free energy of binding for theophylline and caffeine to RNA aptamer using alchemical transformation  
Yoshiaki Tanida, Azuma Matsuura
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