

ASI 杉田理論生物化学研究室

誌 上 発 表 Publications

[雑誌]

(原著論文) *印は査読制度がある論文誌

Re S., Miyashita N., Yamaguchi Y., and Sugita Y.: "Structural Diversity and Changes in Conformational Equilibria of Biantennary Complex-Type N-Glycans in Water Revealed by Replica-Exchange Molecular Dynamics Simulation", *Biophys. J.* **101**, No. 10, pp. L44-L46 (2011). *

Imai T., Miyashita N., Sugita Y., Andriy K., Hirata F., and Kidera A.: "Functionality Mapping on Internal Surfaces of Multidrug Transporter AcrB Based on Molecular Theory of Solvation: Implications for Drug Efflux Pathway", *J. Phys. Chem. B* **115**, No. 25, pp. 8288-8295 (2011). *

Shibata N., Sato H., Sakaki S., and Sugita Y.: "Theoretical Study of Magnesium Fluoride in Aqueous Solution", *J. Phys. Chem. B* **115**, No. 35, pp. 10553-10559 (2011). *

Yasuda S., Yoshidome T., Harano Y., Roth R., Oshima H., Oda K., Sugita Y., Ikeguchi M., and Kinoshita M.: "Free-energy function for discriminating the native fold of a protein from misfolded decoys", *Proteins: Struct. Funct. Bioinf.* **79**, No. 7, pp. 2161-2171 (2011). *

Ogushi F., Ishitsuka R., Kobayashi T., and Sugita Y.: "Rapid flip-flop motions of diacylglycerol and ceramide in phospholipid bilayers", *Chem. Phys. Lett.* **522**, 96-102 (2012). *

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李秀栄, 杉田有治: "QM/MM 法を用いたリン酸化反応の遷移状態モデリング", *薬学雑誌* **131**, No. 8, pp. 1171-1182 (2011).

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Miyashita N., Maenaka K., and Sugita Y.: "Simulation studies on the differences in the binding mechanism of LILRB1/HLA-G and LILRB2/HLA-G", Algorithms in MacroMolecular Modeling Conference, (University of Texas), Austin, USA, Nov. (2009).

Ogushi F., Ishitsuka R., Kobayashi T., and Sugita Y.: "Flip-flop motions of lipid molecules in mixed lipid bilayer systems", Biophysical Society 54th Annual Meeting, (Biophysical society), San francisco, USA, Feb.-

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Pisliakov A. and Sugita Y.: "Computer simulations of proton transfer in cytochrome c oxidase and nitric oxide reductase", 16th European Bioenergetics Conference, Warsaw, Poland, July (2010).

Ogushi F., Ishitsuka R., Kobayashi T., and Sugita Y.: "Flip-flop motions of ceramides, diacylglycerols, and cholesterol in phospholipid bilayer systems", ISSP/SOFT2010, (Institute of solid state physics, the Univ. of Tokyo), Kashiwa, Aug.-Aug. (2010).

Pisliakov A. and Sugita Y.: "Computer Simulations Reveal Proton Transfer Pathways in Nitric Oxide Reductase", Frontiers in the Simulation of Macromolecules, Los Angeles, USA, Nov. (2010).

Pisliakov A. and Sugita Y.: "Water dynamics and proton transfer in nitric oxide reductase: Insights from computer simulations", 2nd Cold Spring Harbor Asia conference on Membrane Proteins: Structure and Function, (Cold Spring Harbor), Suzhou, China, May (2011).

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- (CBSB11), (Julich), Julich, Germany, July (2011).
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- Sugita Y., Ikeguchi M., and Toyoshima C.: "Relationship between Ca^{2+} -Affinity and Shielding of Bulk Water in the Ca^{2+} -ATPase Revealed Molecular Dynamics Simulations", From Computational Biophysics to Systems Biology 2011 (CBSB11), (NIC Coordination Bureau Julich Supercomputing Centre (JSC) Institute for Advanced Simulation (IAS) Forschungszentrum Julich D-52425 Julich, Germany), Julich, Germany, July (2011).
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- Nishima W., Miyashita N., Yamaguchi Y., Sugita Y., and Re S.: "Modulation mechanism on the conformational diversities of biantennary complex-type N-glycans in water", 71st Okazaki Conference on "New perspectives on molecular science of glycoconjugates", (Institute for Molecular Science), Okazaki, Oct. (2011).
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- 李帆=, 宮下尚之, 石戸聰, 杉田有治: “A multi-dimensional umbrella sampling approach for structure prediction of transmembrane helical dimer”, 第 49 回日本生物物理学年会, (日本生物物理学学会), 姫路, 9 月 (2011).
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