

**「分子認識と分子計算」分野**  
**Selected Oral Presentations**  
**(Molecular recognition and molecular modeling)**

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**口頭発表演題** (1演題 10分:発表 7分、質疑応答 3分)

**1. [P1-01] Development status of ABINIT-MP program in 2019**

Yuji Mochizuki

Rikkyo University

**2. [P1-03] Binding Energy Calculation of Protein-Peptide Complex Using Unbiased MD Simulations and MSM Analysis**

Hiroaki Hata

Tokyo Institute of Technology

**3. [P1-04] Interaction Analyses between Calcite/Apatite and Peptides by Fragment Molecular Orbital Method**

Ryo Hatada

Rikkyo University

**4. [P1-05] Fragment Molecular Orbital Method Applied to Factor Xa Inhibitors**

Kazufumi Ohkawa

Asahi Kasei Pharma Corporation

5. [P1-08] Regulation mechanism of agonistic / antagonistic activities of vitamin D receptor analyzed by generalized ensemble method  
Takafumi Kudo  
Yokohama City University
  
6. [P1-12] Development of the CHARMM force field for Cyclosporine A and application to molecular dynamics simulations using a membrane-water system  
Tsutomu Yamane  
Yokohama City University
  
7. [P1-20] Selectivity of phosphodiesterase-10A inhibitor for phosphodiesterase family elucidated by free energy perturbation approach  
Toru Ekimoto  
Yokohama City University
  
8. [P1-21] *De Novo* Binding Prediction using gREST  
Suyong Re  
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