日時: 2019年10月22日16:00-17:30

場所: 2 階平安

「創薬データサイエンス」分野

Selected Oral Presentations (Drug Discovery Data Science)

モデレーター: 池田 和由 Kazuyoshi Ikeda

慶應義塾大学 Keio University

口頭発表演題 (1演題 10分:発表 8分、質疑 2分)

 [P7-08] Prediction of Compound-Protein Interactions and Visualization Based on Graph Convolutional Networks

池口 茉莉恵 Marie Ikeguchi

Kyoto University

2. [P7-03] Predicting drug indications and therapeutic target modules based on disease similarity by interpretable machine learning models

澤田 隆介 Ryusuke Sawada

Kyushu Institute of Technology

3. [P7-09] Development of data curation and integration protocol for the chemical library in early drug discovery

清水 祐吾 Yugo Shimizu

Keio University

4. [P7-10] Clustering therapeutic drugs based on similarities of indications and side effects reported in public database

新川 龍太郎 Ryutaro Shinkawa

Mie University School of Medicine

5. [P7-11] Drug Discovery Raid Battle 2018: an open challenge to discover PD-1/PD-L1 small-molecule inhibitors

山本 一樹 Kazuki Yamamoto

Isotope Science Center, University of Tokyo

6. [P7-05] The Multiple Representation of Protein Sequence Motifs Using Sequence Binary Decision Diagrams

加藤 博明 Hiroaki Kato

National Institute of Technology, Hiroshima College

7. [P7-12] Automatic Reading of Tables and Figures in Scientific Papers

進藤 裕之 Hiroyuki Shindo

Nara Institute of Science and Technology