日時: 2018年10月11日 14:00-15:30

場所: 研修室

「医薬品研究と ADMET」分野 及び「レギュラトリサイエンス」分野 口頭発表

Selected Oral Presentations

(Information and computing approach for drug design and ADMET study)
(Regulatory Sciences)

座長: 渡邉 怜子 Reiko Watanabe

医薬基盤・健康・栄養研究所

National Institutes of Biomedical Innovation, Health and Nutrition

小村 弘 Hiroshi Komura 日本医療研究開発機構 Japan Agency for Medical Research and Development

「医薬品研究と ADMET」分野

1. [P4-05] Development of a pharmacokinetics prediction system using multiscale integrated modeling: 8. Web application and database consisting of curated public data and newly acquired experimental data

川島 和 Hitoshi Kawashima

医薬基盤・健康・栄養研究所

National Institutes of Biomedical Innovation, Health and Nutrition

2. [P4-11] Development of an informatics system for predicting cardiotoxicity: 5. Quantitative model for hERG blocking small molecules based on the integrated database

佐藤 朋広 Tomohiro Sato

理化学研究所 RIKEN Center for Biosystems Dynamics Research

3. [P4-12] Use of Markov Chain Monte Carlo method to integrate in vitro & in vivo data for prediction of drug interactions caused by inhibition of multiple CYP species

保月静香 Shizuka Hozuki

千葉大学 Chiba University

4. [P4-13] Individual data analysis of patients participated in clinical studies: relationship between longitudinal changes in cardiac functions and mortality risk in CHF

畔 さゆり Sayuri Guro

千葉大学 Chiba University

5. [P4-02] Development of an in vitro evaluation system for drug-induced hepatotoxicity using PXB-cells

稲松 睦 Mutsumi Inamatsu

株式会社フェニックスバイオ PhoenixBio co., Ltd.

「レギュラトリサイエンス」分野

6. [P6-01] Universal Read-Across Approach to predict toxicities 永堀 博久 Hirohisa Nagahori 住友化学株式会社 Sumitomo Chemical

7. [P6-03] Quantitative Structure-Activity Relationship (QSAR) analysis using deep learning based on Deep Snap, a novel molecular image input technique 松坂 恭成 Yasunari Matsuzaka

明治薬科大学 Meiji Pharmaceutical University