

T-LSI and C-AIR Joint Symposium on Life Science Innovation and Artificial Intelligence

Date: Monday, May 8, 2017

Venue: Room 308, Innovative Medical Research Institute, University of Tsukuba

Access: <http://www.tsukuba.ac.jp/access/gmap/gmap.php?i=281010>

Program:

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| 13:40 - 13:45 | Opening |
| 13:45 - 14:00 | <i>Bioinformatics field in T-LSI</i> Hiroko Isoda Program Leader, Master's/Doctoral Program in Life Science Innovation, University of Tsukuba |
| 14:00 - 14:30 | <i>Phenotype switching in cancer progression: the potential of pharmacophenomics in cancer therapy</i> Colin Goding Group Leader, Ludwig Institute for Cancer Research, University of Oxford |
| 14:30 - 15:00 | <i>Protein informatics for antibody drug discovery</i> Hiroki Shirai Executive Fellow, Modality Research Laboratories, Astellas Pharma, Co Ltd. |
| 15:00 - 15:30 | <i>An efficient regulatory network inference algorithm from single-cell RNA-Seq during differentiation</i> Hiroataka Matsumoto Research Fellow of JSPS, Advanced Center for Computing and Communication, RIKEN |
| 15:30 - 15:45 | Coffee break |
| 15:45 - 16:15 | <i>Computational and systems biology approaches to personalized medicine</i> Francesca Buffa Group Leader, Department of Oncology, University of Oxford |
| 16:15 - 16:45 | <i>The leading-edge applications of mass spectrometry in the drug discovery and diagnosis</i> Taka-Aki Sato Director, R&D Center for Precision Medicine, University of Tsukuba |
| 16:45 - 17:15 | <i>A scalable eigenanalysis engine for large-scale simulations and data analysis</i> Tetsuya Sakurai Director, Center for Artificial Intelligence Research (C-AIR), University of Tsukuba |
| 17:15 - 17:20 | Closing |