

# 東アジア人類遺伝学会連合 第14回大会シンポジウム / The 14th Annual Meeting of the East Asian Union of Human Genetics Societies

## 東アジア人類遺伝学会連合 第 14 回大会 / The 14th Annual Meeting of the East Asian Union of Human Genetics Societies

日 時 : 11 月 20 日 (木) 9:30 ~ 10:50、11:00 ~ 12:20、14:45 ~ 16:25 第 2 会場 (小ホール)  
Date : Nov. 20 (Thu.) 9:30-10:50, 11:00-12:20, 14:45-16:25 Room 2 (Small Hall 5F)

Chairs : Xue Zhang (Chinese Society of Human Genetics)  
Masao Nagasaki (Tohoku University)

### EAUHGS-1 Korean Reference Genome Variation Project (KRGVP): Contents and Implication

9:30

○ Seong Beom Cho

Division of Biomedical Informatics, Center for Genome Science, National Institute of Health, KCDC

### EAUHGS-2 One thousand Japanese whole genome project in ToMMo

9:50

○ Masao Nagasaki

a Department of Integrative Genomics, Tohoku Medical Megabank Organization, Sendai, 980-8575, Japan

### EAUHGS-3 Molecular diagnosis of hereditary hearing loss and its related translational medicine

10:10

○ Guangxin Xiang

Translational Medicine Institute, CapitalBio Corporation & National Engineering Research Center for Beijing  
Biochip Technology, Beijing, China

### EAUHGS-4 The role of *de novo* variants in rare disease pathogenesis

10:30

○ Murim Choi

Department of Biomedical Sciences, Seoul National University College of Medicine, Seoul, Korea.

10:50-11:00 break

Chairs : Jin-Sung Lee (Korean Society of Medical Genetics)  
Yoko Aoki (Tohoku University)

### EAUHGS-5 MeCP2 in autism — displays allelic expression imbalance and affects both nerve system and heart development by regulating Notch pathway

11:00

Yufang Zheng, Yumeng Wang, Yahui Liu, Zhangmin Yang, Yanqin He, Keping Hu,

○ Hongyan Wang

State Key Laboratory of Genetic Engineering, School of Life Sciences Fudan University

### EAUHGS-6 Brain Somatic Mutations in Neurodevelopmental Disorders: Lessons from Epilepsy Patients

11:20

○ Jeong Ho Lee

Graduate School of Medical Science and Engineering, KAIST (Korea Advanced Institute of Science and  
Technology), Daejeon 305-701, Korea

### EAUHGS-7 Next-generation sequencing applications in pre- and postnatal genetic diagnostics

11:40

○ Desheng Liang<sup>1,2)</sup>、Weigang Lv<sup>1)</sup>、David S Cram<sup>3)</sup>、Lingqian Wu<sup>1,2)</sup>

1) State Key Laboratory of Medical Genetics, Central South University, Changsha, China

2) Hunan Jiahui Genetics Hospital, Changsha, Hunan, China

3) Berry Genomics C., Limited

- EAUHGS-8 12:00** 次世代シーケンサーを用いた RASopathies の遺伝子解析 /  
Molecular analysis of RASopathies using next generation sequencer
- 青木 洋子<sup>1)</sup> (Yoko Aoki)、新堀 哲也<sup>1)</sup> (Tetsuya Niihori)、井上 晋一<sup>1)</sup> (Shin-ichi Inoue)、  
松原 洋一<sup>1,2)</sup> (Yoichi Matsubara)
- 1) 東北大学大学院 医学系研究科 遺伝病学分野  
(Department of Medical Genetics, Tohoku University School of Medicine, Sendai, Japan)
  - 2) 国立成育医療研究センター研究所  
(National Research Institute for Child Health and Development, Tokyo, Japan)

12:20-14:45 Lunch break

Chairs : Kyuyoung Song (University of Ulsan College of Medicine)  
Taisei Mushiroda (RIKEN)

- EAUHGS-9 14:45** Genome-wide association analyses in Han Chinese identify two new susceptibility loci for  
amyotrophic lateral sclerosis
- Min Deng<sup>1)</sup>、Ling Wei<sup>2)</sup>、Xianbo Zuo<sup>3)</sup>、Xiaodong Ju<sup>4)</sup>、Kai Wang<sup>2)</sup>
- 1) Medical Research Center, Peking University Third Hospital, Beijing, 100191, China.
  - 2) Department of Neurology, No.1 Hospital, Anhui Medical University, Hefei, Anhui, 230022, China.
  - 3) State Key Laboratory Incubation Base of Dermatology, Ministry of National Science and Technology, Hefei, Anhui, 230032, China.
  - 4) Institute of Sports Medicine, Peking University Third Hospital, Beijing, 100191, China.

- EAUHGS-10 15:05** Esophageal squamous cell carcinoma in China: interaction of environmental and genetic  
factors on esophageal carcinogenesis
- Li Dong Wang
- Henan Key Laboratory for Esophageal Cancer Research The First Affiliated Hospital of Zhengzhou University,  
Zhengzhou, Henan Province 450052

- EAUHGS-11 15:25** 大規模ヒト疾患ゲノム解析を通じた疾患病態解明・新規創薬への展望 /  
Comprehensive strategies to utilize large scale human genetic studies for elucidation of novel  
disease biology and drug discovery
- 岡田 随象<sup>1,2)</sup> (Yukinori Okada)
- 1) 東京医科歯科大学 大学院医歯学総合研究科 疾患多様性遺伝学分野  
(Department of Human Genetics and Disease Diversity, Tokyo Medical and Dental University Graduate School  
of Medical and Dental Sciences, Tokyo, Japan.)
  - 2) 理化学研究所 統合生命医科学研究センター 統計解析研究チーム  
(Laboratory for Statistical Analysis, Center for Integrative Medical Sciences, RIKEN, Yokohama, Japan.)

- EAUHGS-12 15:45** *NUDT15* R139C is a risk factor for thiopurine-induced leukopenia
- Kyuyoung Song
- Department of Biochemistry and Molecular Biology, University of Ulsan College of Medicine, Seoul, Korea

- EAUHGS-13 16:05** 薬疹関連ゲノムバイオマーカーの同定と医学的有用性の検証 /  
Identification of genomic biomarkers associated with cutaneous adverse drug reactions and  
validation of medical utility of genetic testing
- 蒔田 泰誠 (Taisei Mushiroda)
- 理化学研究所 統合生命医科学研究センター  
(RIKEN Center for Integrative Medical Sciences)