Future of Omics Conference - Nordic Station

4th November, 2024

10.00 Registration in KI BioMedicum (Solnavägen 9, Stockholm)

Microbiology section

10.15-10.30 Welcome speech

<u>10.30-11.00</u> Mapping the human microbiome in individuals, cohorts and populations (Prof. Lars Engstrand, Director of CTMR, Chair of MMHP project)

<u>11.00-11.30</u> Advancing genomics applications for enhanced public health surveillance (Dr. Maike Seifert, Karolinska Institutet)

<u>11.30-12.00</u> The human gut microbiota in health and disease (Prof. Karsten Kristiansen, Copenhagen University, MMHP consortium member)

<u>12.00-12.30</u> HELMI birth cohort study on early life microbiome (Anne Salonen, University of Helsinki, MMHP consortium member)

<u>12.30-13.00</u> From Methods to Microbes: Navigating the Metagenomics investigation of Phage Populations in the infant Gut (Alise Ponsero, Quadram Institute, UK & University of Helsinki)

13.00-14.30 Lunch break

Multi-omics section

<u>14.30-15.00</u> Genetic drivers of heterogeneity in type 2 diabetes pathophysiology (Prof. Torben Hansen, Nova Nordisk Foundation, Center for Basic Metabolic Research, Copenhagen University) <u>15.00-15.30</u> Decipher Clinical Signatures of Gastrointestinal Tumors through Multi-Omics Big Data

<u>15.30-16.00</u> Single-cell transcriptomics of the placenta: bridging the understanding of maternal conditions and offspring health (Prof. Qiaolin Deng, Stockholm University)

16.00-16.20 Coffee break

<u>16.20-16.50</u> Genetic and molecular basis of nervous system disorders (Prof. Jans Mulder, Karolinska Institutet)

<u>16.50-17.20</u> Exposing causal biomarkers for complex diseases using state-of-the-art proteogenomics (Prof. Xia Shen, Karolinska Institutet)

<u>17.20-17.50</u> Past and Future Analyses in Forensic Human Identification (Prof. Marie Allen, Uppsala University)

<u>17.50-18.00</u> Walk to Karolinska Cancer Centre (Change venue location to Cancer Center Karolinska building)

<u>18.15-18:45</u> Unlocking new RNA insights with next-generation single-cell RNA-seq (Dr. Christoph Ziegenhain, CEO of Xpress Genomics)

<u>18:45-19:15</u> Automated NGS total solution for cancer genomics profiling: from sample to report (Ms. Nancy Yang, BD Manager of Gene+)

19:15-21.00 Buffer Dinner with lab tour