NGS Analytics and Genome Informatics Program (Hall 3)

| 02 nd Feb. | TOPIC | SPEAKER |
|-----------------------|---|---|
| 10:30 – 11:15 | Next-Generation Sequencing (NGS) Basics | |
| | Sequencing Technologies, NGS File Format, Command Line Computing for Genomics, NGS Data Analytics, Genomics Data Storage, Fast Parallel Computing using GPU Machines | Dr Raja Mugasimangalam Founder & CEO – Genotypic & QTLomics Chairman – iBioM, India |
| 18:30 - 19:15 | Insights into The Microbiome Analysis Using NGS Data | |
| 19:15 – 20:00 | 16S rRNA shotgun sequencing data analysis and Visualization Whole Genome Metagenome - Assembly, Gene prediction and Visualization, Commander Demonstration | Dr Prasanna Koti Associate Principal Scientist Genotypic Technology, India |

| 03 rd Feb. | TOPIC | SPEAKER |
|-----------------------|---|--|
| 08:30 - 09:30 | Insights into Transcriptome and Small RNA data analysis | Ms Garima Sanoria Scientist, Genome Informatics Genotypic Technology, India |
| 11:15 - 12:15 | Insights into Whole Genome Sequencing: Alignment to Variant Calling and Reporting | Mr Nihar Bachan Das Scientist, Genome Informatics Genotypic Technology, India |
| 12:15 - 12:45 | Making sense of variants and mutations; FastQ to BAM to VCF to SNP table> annotations | Dr Chellappa Gopalakrishnan Founder and CEO, Genomatics Pvt. Ltd., India |
| 12:45 - 13:15 | Advanced computational approaches for chromosome-scale haplotype-resolved genomics | Dr Shilpa Garg Senior Researcher, Novo Nordisk Genomics Sustainable Solutions, Denmark |
| 14:30 - 15:00 | Long Read Sequencing for whole genome assembly and scaffolding | Dr Zuwei Qian Director, Marketing, Asia Pacific Pacific Biosciences |
| 15:00 – 15:30 | Long Read Sequencing for whole genome assembly and scaffolding | Dr Khi Pin, Chua Senior Bioinformatics Scientist, Pacific Biosciences |
| 15:30 – 15:45 | NGSA Test (Closure of NGSA) | |