Genomics, Epigenomics and Sequencing Core

GES Core service and collaboration

Illumina next generation sequencing

- RNA-seq; miRNA-seq
- DNA-seq; ChIP-seq; exome and targeted sequencing; amplicon-seq
- WGS and WGBS
- Whole mitochondrial genome sequencing (new)
- Methyl-seq
- Ultra-low input RNA-seq, miRNA-seq and ChIP-seq
- Single cell RNA-seq (manual, new)
- 16S metagenomic sequencing (new)
- RRBS and ultra-low input RRBS

Illumina Infinium HD beadchip assay

- DNA methylation association study: MethylationEPIC
- Beadchip-based 5-hmC assay (new)
- Genotyping and CNV assay

DNA/RNA extraction

- Cell; Tissue; Blood; FFPE sample, etc
- DNA/RNA QC
 - Agilent Bioanalyzer assay
 - DNA quantification by fluorescent-based Qubit assay
 - Agarose gel electrophoresis

NGS-based multiplexed bisulfite PCR sequencing (new)

- Methylation analysis of multiple targeted regions in a single assay at single CpG resolution

Bisulfite PCR sequencing

- Promoter CpG island analysis and primer design for bisulfite PCR
- Cloning and sequencing
- Data analysis

Real-time qPCR

- Real-time RT-PCR for RNA-seq data validation (new)
- Taq-Man CNV/SNP assay

Other service/research: please inquire

Proposal and manuscript preparation

- Collaboration
- Consultation
- Materials and Methods
- Supporting letter

Contact

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