

Chúc Phụ Nữ Việt Nam luôn vui tươi xinh!



MGMA 2024

Long Read Microbiome

20 Oct 2024

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Content

- Projects find home
- PacBio and ONT present their microbiome platforms
- ONT: De novo assembly + mapping + variant calling
- PacBio: 16S Full-length

Projects find home

- Each group select a project either ONT WGS, ONT full length 16S, ONT shotgun, PacBio WGS, PacBio full length 16S and PacBio shotgun
- One or 2 or 3 students per group
- Presentation's day: 23/11/2024 (30 min = present 20 min + Q&A 10 min)
- List of Projects and register at

https://docs.google.com/spreadsheets/d/1M_KTe991iK4zjKD3HL1fml-FMEHXirKvB65XYLUnVbl/edit?gid=0#gid=0

| | A | B | C | D | E | F | G | H |
|---|-----|---|--|--|------------|--------------|----------------|---------------------|
| 1 | No. | Title of Papers | Doi/link | NCBI/ENA Accession Number | Type | Tutor's name | Student's name | Presentation's Date |
| 2 | 1 | Multiplex PCR Approach for Rapid African Swine Fever Virus Genotyping | https://doi.org/10.3390/v16091460 https://www.mdpi.com/1999-4915/16/9/1460 | https://www.ncbi.nlm.nih.gov/bioproject/PRJEB75237 https://www.ncbi.nlm.nih.gov/sra?term=ERP159826 | ONT WGS | Phát và Huy | | 23 Nov 2024 |
| 3 | 2 | Evaluation of Nanopore sequencing for Mycobacterium tuberculosis drug susceptibility testing and outbreak investigation: a genomic analysis | https://doi.org/10.1016/S2666-5247(22)00301-9 https://www.thelancet.com/journals/lanmic/article/PIIS2666-5247(22)00301-9/fulltext | https://www.ebi.ac.uk/ena/browser/view/PRJEB49093 | ONT WGS | Phát và Huy | | 23 Nov 2024 |
| 4 | 3 | A comparison between full-length 16S rRNA Oxford nanopore sequencing and Illumina V3-V4 16S rRNA sequencing in head and neck cancer tissues | doi: 10.1007/s00203-024-03985-7 | https://www.ncbi.nlm.nih.gov/bioproject/?term=PRJNA1087430 | ONT 16S FL | Khải và Kim | | 23 Nov 2024 |

PacBio and ONT present their microbiome platforms

MGMA2024

Online & Free Course

LONG READ MICROBIOME

27-10-2024 | 07:00 - 08:30 PM | SUNDAY

PACBIO HIFI SEQUENCING FOR HIGH-RESOLUTION MICROBIOME RESEARCH

Pacific Biosciences (PacBio) has emerged as a leader in the field of microbiome research, leveraging its advanced long-read sequencing technologies (HiFi sequencing) to provide deeper insights into the microbial world. HiFi sequencing enables the generation of reference-grade, highly accurate, closed chromosomes, and plasmids, even from the most repeat-dense and GC-rich genomes. Additionally, PacBio's sequencing solutions allow for species- and strain-level identification through reliable and cost-effective full-length 16S rRNA sequencing. For shotgun metagenomic application, HiFi sequencing provides microbiome researchers the power to precisely profile the function and taxonomy of a microbial community, while also improving the recovery of high-quality metagenome-assembled genomes (MAGs), including circular, single-contig MAGs, even at lower sequencing coverage.

Contact us:
✉ bioinformatics.mgma@gmail.com
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MICROBIAL GENOME & MICROBIOME ANALYSIS

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LONG READ MICROBIOME

27-10-2024 | 08:30 - 10:00 PM | SUNDAY

WHAT YOU'RE MISSING MATTERS - DELIVERING THE FUTURE OF MICROBIAL GENOMICS WITH OXFORD NANOPORE

Genomics and Epigenetics

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