

PacBio Office Hours

Hosted by the UC Riverside Genomics Core

Wednesday, February 14th, 2024

10am – noon

Location: Keen Hall, Room 2018

Stop by and chat with the PacBio team! Get questions answered about current projects, help scoping future ones, or learn about the latest and greatest in whole genome sequencing, RNA sequencing, and more!

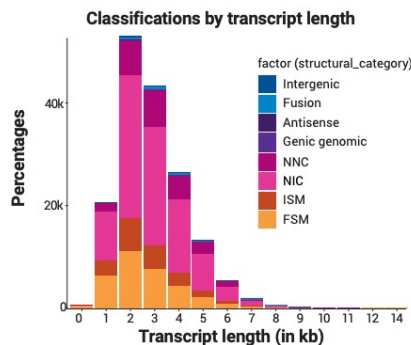
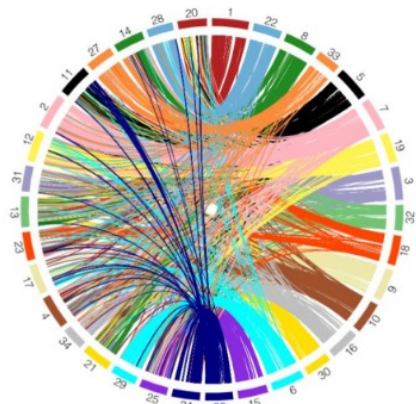
Coffee and snacks will be provided.



Register here!

What can you do with HiFi Sequencing?

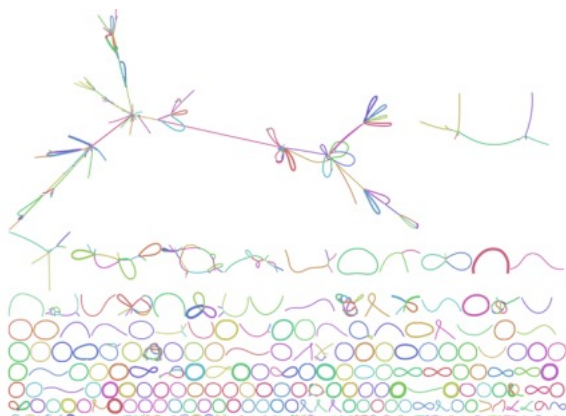
Reference-quality genome assemblies No-assembly required transcriptomes



- 162,290 transcripts
- Min: 80 bp
- Max: 14,288 bp
- Mean: 3,347 bp

Detect full-length transcripts up to 14 kb

Shotgun Metagenome Profiling



Native Methylation Detection

