

Happy New Year! We are pleased to invite you to join the first of our PacBio APAC Webinar Series of 2021!

PacBio's HiFi sequencing is currently the best-in-class data type for various genomic applications. Specifically in metagenomics applications, the combination of length and accuracy of HiFi reads allow a much higher resolution of the underlying complex populations.

## #1 of PacBio's APAC Webinar Series 2021. Register to Attend Today

In this webinar, we will provide recommendations and best practices for wet-lab workflows in order to generate the highest quality HiFi metagenomic data using PacBio Sequel system.

- Then, we will discuss the bioinformatics workflow starting from CCS to command line analysis of HiFi data.
- Finally, we will also take a deep dive into a shotgun metagenomics analysis to demonstrate how the high accuracy of HiFi sequencing allows the assembly of many complete or near-complete metagenomic-assembled-genomes (MAG).

Register now: https://programs.pacificbiosciences.com/l/1652/2021-01-07/42jdhc