

Join in the Race to Unravel the SARS-CoV-2 Genome

As COVID-19 virus continues to spread, there is a need to decipher the SARS-CoV-2 viral genome which is constantly evolving. Next-generation sequencing (NGS) provides an effective high-throughput solution for not just screening multiple samples but also for genome discovery projects.

Applications for SARS-CoV-2 Genome Sequencing



SARS-CoV-2

Genome Discovery

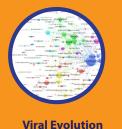


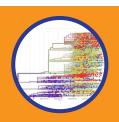




Viral Surveillance and Epidemiology Research







1st BASE

Phylogenetic Studies

High-Performance & Reproducible SARS-CoV-2 Genome Sequencing Services from 1st BASE

Adopts a targeted NGS approach with the ARTIC network v3 panel and is a valuable tool to accurately sequence the SARS-CoV-2 genome for the characterization of viruses. The service features bioinformatics analysis that **maps the genome** to the reference **to identify genome variants** as well as a **de novo assembly pipeline for discovery works.** The relationship of each samples will also be presented with a phylogenetic tree.



All-in-One Package From sample preparation to bioinformatics analysis.



High Quality Genome Assembly Improved assembly with genome polishing. >200 sequencing coverage per sample.



Fast Turnaround Time 4-6 weeks from sample to analysis.



Sanger Sequencing Services



Viral Transport Medium (VTM)

Viral DNA/RNA Extraction Kit

2019-nCoV Plasmid Positive Controls

Please contact our distributor below for orders and enquiries.



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