**Note:**

1. All SARS-CoV-2 whole genome sequences reported here were generated using the ARTIC network tiling amplicon COVID-19 protocol v4.1 (Oxford Nanopore Technology) and analyzed using ncov2019-artic-nf pipeline (artic-ncov2019 bioinformatics SOP implemented in nextflow), with 99% coverage and ~450x depth.
2. WGS results must be interpreted along with available clinical and epidemiological information.
3. This report is electronically signed.

**Reporting Laboratory:** RITM - MOLECULAR BIOLOGY LABORATORY

Date and Time: 06 JUN 2022 1900H

|  |  |  |  |
| --- | --- | --- | --- |
| Performed By: | Analysed By: | Verified By: | Approved By: |
| KAREN IANA TOMAS, RMT  AMALEA DULCENE NICOLASORA, MSc  ARDIANE YSABELLE DOLOR, RMT  CZARINA CHRISTELLE ALYANNAH CELIS, RMT  STEPHEN PAUL ORTIA, RMT | FRANCISCO GERARDO POLOTAN, MSc  GERALD IVAN SOTELO | MA. ANGELICA TUJAN, MSc | LEI LANNA DANCEL  OIC, RITM-MOLECULAR BIOLOGY LABORATORY |
|  |  |  | TIMOTHY JOHN DIZON, MD, MBA  PRINCIPAL INVESTIGATOR, GECO-PH |

|  |
| --- |
|  |