

Search Details

|                                       |                                                                                                    |
|---------------------------------------|----------------------------------------------------------------------------------------------------|
| Job Title/ Assay Name : Test sequence | Amplicon Sequence:<br>GACCCCAAAATCAGCGAAATGCACCCCGCATTACGTTTGGTGGACC<br>CTCAGATTCAACTGGCAGTAACCAGA |
| Effective Date : 2025-11-07           | Forward: GACCCCAAAATCAGCGAAAT                                                                      |
| Database : core_nt                    | Reverse: TCTGGTTACTGCCAGTTGAATCTG                                                                  |
| Alignment algorithm : blastN          | Probe: ACCCCGCATTACGTTTGGTGGACC                                                                    |
| Exclusive Organism:                   | Inclusive Organism:                                                                                |
| RID: GVFWM794014                      | Total Result: 5000                                                                                 |
| App Version: 1.0                      |                                                                                                    |

| No of Accessions                                             | Accession Details    | Oligo | Mutation     | BDS Analysis Performance Prediction |
|--------------------------------------------------------------|----------------------|-------|--------------|-------------------------------------|
| Betacoronavirus pandemicum/3418604 Analyzed Accessions: 5000 |                      |       |              |                                     |
| <a href="#">5000</a>                                         | <a href="#">5000</a> | NA    | No Mutations | <div>✓✓✓</div> Detected             |