

CT_{RNA} (gene, kit)	CT_{PK+HID} (gene, kit)
30.7 (N, DisCoVery) 30.9 (ORF1ab, DisCoVery) 27.4 (RP, DisCoVery)	nd (N, DisCoVery) nd (ORF1ab, DisCoVery) 23.2 (RP, DisCoVery)
23.5 (N, DisCoVery) 21.1 (ORF1ab, DisCoVery) 26.2 (RP, DisCoVery)	nd (N, DisCoVery) nd (ORF1ab, DisCoVery) 26.0 (RP, DisCoVery)
31.3 (N, GeneFinder) 31.3 (RdRp, GeneFinder) 32.1 (E, GeneFinder) 27.3 (RP, Genefinder)	40.1 (N, GeneFinder) nd (RdRp, GeneFinder) nd (E, GeneFinder) 30.3 (RP, Genefinder)
34.6 (N, GeneFinder) 34.5 (RdRp, GeneFinder) 38.8 (E, GeneFinder) 27.2 (RP, Genefinder)	41.3 (N, GeneFinder) nd (RdRp, GeneFinder) nd (E, GeneFinder) 28.2 (RP, Genefinder)
35.9 (N, GeneFinder) 37.0 (RdRp, GeneFinder) nd (E, GeneFinder) 28.4 (RP, Genefinder)	nd (N, GeneFinder) nd (RdRp, GeneFinder) nd (E, GeneFinder) 30.6 (RP, Genefinder)

S3 Table. CT values of false-negative samples detected with the PK+HID method, using the results obtained with purified RNA samples as reference.