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| --- | --- | --- | --- | --- | --- | --- | --- |
| **Accession** | **Virus** | **BLASTx coverage (%)** | **BLASTx identity (%)** | **BLASTx accession** | **BLASTn coverage (%)** | **BLASTn identity (%)** | **BLASTn accession** |
| OQ818319 | *E. dupreanum* sapovirus 1 | 90 | 67.18 | AQQ78883.1 | 1 | 79.82 | LC504397.1 |
| OQ818340 | *E. dupreanum* sapovirus 2 | 99 | 67.97 | AQQ78883.1 | 63 | 71.4 | KX759619.1 |
| OQ818345 | *R. madagascariensis* sapovirus 1 | 70 | 64.23 | WBG95506.1 | No result | No result | No result |
| OQ818347 | *R. madagascariensis* sapovirus 2 | 99 | 75.8 | WBV74378.1 | 11 | 72.44 | LC380411.1 |
| OQ818348 | *R. madagascariensis* sapovirus 3 | 99 | 81.38 | WGC86364.1 | 89 | 74.92 | OQ709197.1 |
| PP766460 | *E. dupreanum* sapovirus 4 | 70 | 75.26 | AQQ78883.1 | No result | No result | No result |
| PP766461 | *E. dupreanum* sapovirus 3 | 96 | 90.13 | AQQ78883.1 | 92 | 80.7 | KX759623.1 |
| PP766468 | *R. madagascariensis* sapovirus 3 | 100 | 78.75 | WBV74378.1 | No result | No result | No result |
| PP766470 | *R. madagascariensis* sapovirus 2 | 99 | 79.01 | WBV74378.1 | 1 | 86.67 | KU712497.1 |
| PP766473 | *R. madagascariensis* sapovirus 2 | 99 | 80.86 | WBV74378.1 | 1 | 100 | PP554904.1 |
| PP766474 | *R. madagascariensis* sapovirus 2 | 100 | 77.25 | WBV74378.1 | 2 | 94.44 | KX907730.1 |
| PP766476 | *R. madagascariensis* sapovirus 2 | 99 | 70.94 | WBV74378.1 | 19 | 73.71 | KX759623.1 |
| PP766477 | *R. madagascariensis* sapovirus 2 | 34 | 80.51 | WBV74378.1 | No result | No result | No result |