|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Novel**  **sequence** | **Top BLASTn hit** | **Top BLASTx hit** | **Identity/coverage top BLAST nt (%)** | **Identity/coverage top BLAST aa (%)** | **Accession number of top BLASTn hit** | **Accession number of top BLASTx hit** |
| OQ818330 | NA | *Arracacha virus B* | NA/NA | 57.24/87 | NA | QCY50806.1 |
| OQ818331 | NA | *Tetnovirus 1* | NA/NA | 62.75/83 | NA | ADK97708.1 |
| OQ818332 | *Victoria bee virus 2* | *Victoria bee virus 2* | 89.41/99 | 92.14/99 | MG995724.1 | AWK77887.1 |
| OQ818333 | NA | *Rondonia picorna-like virus* | NA/NA | 22.87/34 | NA | QHI42122.1 |
| OQ818334 | *Victoria bee virus 2* | *Victoria bee virus 2* | 97.83/100 | 97.64/99 | MG995724.1 | AWK77887.1 |
| OQ818335 | *Bat felisavirus* | *Bat felisavirus* | 80.99/97 | 91.01/56 | KX644943.1 | AQP31140.1 |
| OQ818336 | *Victoria bee virus 2* | *Victoria bee virus 2* | 90.63/100 | 93.03/98 | MG995724.1 | AWK77887.1 |
| OQ818337 | *Hepatovirus H2* | *Hepatovirus H2* | 82.40/97 | 94.75/87 | NC\_028366.1 | YP\_009179216.1 |
| OQ818338 | *Aphid lethal paralysis virus* | *Aphid lethal paralysis virus* | 93.86/100 | 98.34/99 | MF458893.1 | QFI36301.1 |
| OQ818339 | *Eotetranychus kankitus picorna-like virus* | *Picornavirales sp.* | 73.35/14 | 45.56/99 | MN745092.1 | ULF99664.1 |
| OQ818340 | *Bat sapovirus* | *Bat sapovirus* | 71.40/63 | 67.97/99 | KX759619.1 | AQQ78883.1 |
| OQ818341 | *Bat felisavirus* | *Bat felisavirus* | 85.33/99 | 95.74/97 | KX644943.1 | AQP31140.1 |
| OQ818342 | *Bat sapelovirus* | *Bat sapelovirus* | 94.73/99 | 98.47/99 | KX644938.1 | YP\_009345901.1 |
| OQ818343 | *Bat sapelovirus* | *Bat sapelovirus* | 92.28/100 | 98.67/99 | KX644938.1 | YP\_009345901.1 |
| OQ818344 | *Bat sapelovirus* | *Bat sapelovirus* | 94.57/99 | 99.15/99 | KX644938.1 | YP\_009345901.1 |
| OQ818345 | NA | *Bat sapovirus BtSY3* | NA/NA | 54.97/97 | NA | WBV74435.1 |
| OQ818346 | Bat picornavirus | *Wilde virus 3* | 81.94/15 | 57.23/93 | ON168930.1 | QQR34440.1 |
| OQ818347 | *Sapovirus Hu/GI.6/Nichinan/FP05284/2005/JPN* | *Bat sapovirus BtSY2* | 72.44/11 | 75.80/99 | LC380411.1 | WBV74378.1 |
| OQ818348 | NA | *Bat sapovirus BtSY1* | NA/NA | 59.91/99 | NA | WBV74377.1 |