|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **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** |
| OQ818316 | *Bat picornavirus 1* *strain LMH22A* | polyprotein [*African bat icavirus*] | 93.18/0 | 46.51/81 | HQ595341.1 | YP\_009121743.1 |
| OQ818317 | *Kunsagivirus B clone Bat/CAM/KuV-P2/2013* | polyprotein [*Kunsagivirus B*] | 81.93/95 | 91.61/90 | NC\_033818.1 | YP\_009345896.1 |
| OQ818318 | *Bat picornavirus 1 strain LMH22A* | polyprotein [*Teschovirus A*] | 81.95/1 | 44.42/90 | HQ595341.1 | AYD75972.1 |
| OQ818319 | NA | polyprotein [*Bat sapovirus*] | NA/NA | 67.15/99 | NA | AQQ78883.1 |
| OQ818320 | *Bat sapelovirus clone Bat/CAM/Sap-P24/2013* | polyprotein [*Bat sapelovirus*] | 94.37/99 | 98.95/96 | KX644938.1 | YP\_009345901.1 |
| OQ818321 | *Bat sapelovirus clone Bat/CAM/Sap-P24/2013* | polyprotein [*Bat sapelovirus*] | 73.31/96 | 75.75/95 | KX644938.1 | YP\_009345901.1 |
| OQ818322 | *Eidolon dupreanum kobuvirus* | polyprotein [*Eidolon dupreanum kobuvirus*] | 99.42/98 | 99.88/87 | OP287812.1 | WBP49885.1 |
| OQ818323 | *Sapelovirus-like porcine picornavirus* | polyprotein [*teschovirus A6*] | 77.36/89.90 | 44.26/88 | LC386162.1 | QWS67609.1 |
| OQ818324 | *Feline picornavirus* | polyprotein [*Porcine teschovirus 16*] | 78.74/1 | 43/89 | NC\_016156.1 | BBE37019.1 |
| OQ818325 | *Marmot sapelovirus 2 strain HT6* | polyprotein [*Bat picornavirus 3*] | 78.86/3 | 59.57/91 | KY855433.1 | YP\_004782204.1 |
| OQ818326 | NA | polyprotein 1 [*Mulberry mosaic leaf roll associated virus*] | NA/NA | 39.51/77 | NA | QIK02125.1 |
| OQ818327 | NA | polyprotein P2-V [*Arabis mosaic virus*] | NA/NA | 28.50/62 | NA | ADJ39333.1 |
| OQ818328 | *Marmot sapelovirus 2 strain HT6* | polyprotein [*Bat picornavirus 3*] | 78.86/3 | 59.90/91 | KY855433.1 | YP\_004782204.1 |
| OQ818329 | *Bat sapelovirus clone Bat/CAM/Sap-P24/2013* | polyprotein [*Bat sapelovirus*] | 84.34/4 | 64.02/96 | KX644938.1 | YP\_009345901.1 |