|  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Novel**  **sequence** | **Virus** | **Country** | **Host** | **Corresponding African bat picornavirus** |  | **Country** | **Host** | **aa similarity (%)** | **nt similarity (%)** | |
| **OQ818316** | *Mischivirus* | Madagascar | *Pteropus rufus* | NC\_026470 |  | DRC | *Macronycteris gigas* | 40% | | 49% |
| **OQ818317** | *Kunsagivirus* | Madagascar | *Eidolon dupreanum* | NC\_033818 |  | Cameroon | *Eidolon helvum* | 89% | | 81% |
| **OQ818319** | *Sapovirus* | Madagascar | *Eidolon dupreanum* | KX759623 |  | Cameroon | *Eidolon helvum* | 70% | | 62% |
| **OQ818320** | *Sapelovirus* | Madagascar | *Eidolon dupreanum* | NC\_033820 |  | Cameroon | *Eidolon helvum* | 99% | | 94% |
| **OQ818321** | *Sapelovirus* | Madagascar | *Eidolon dupreanum* | NC\_033820 |  | Cameroon | *Eidolon helvum* | 72% | | 71% |
| **OQ818322** | *Kobuvirus* | Madagascar | *Eidolon dupreanum* | OP287812 |  | Madagascar | *Eidolon dupreanum* | 99% | | 98% |
| **OQ818329** | *Sapelovirus* | Madagascar | *Rousettus madagascariensis* | NC\_033820 |  | Cameroon | *Eidolon helvum* | 60% | | 64% |
| **OQ818337** | *Hepatovirus* | Madagascar | *Eidolon dupreanum* | NC\_028366 |  | Ghana | *Eidolon helvum* | 86% | | 82% |
| **OQ818340** | *Sapovirus* | Madagascar | *Eidolon dupreanum* | KX759623 |  | Cameroon | *Eidolon helvum* | 68% | | 31% |
| **OQ818342** | *Sapelovirus* | Madagascar | *Eidolon dupreanum* | NC\_033820 |  | Cameroon | *Eidolon helvum* | 99% | | 95% |
| **OQ818343** | *Sapelovirus* | Madagascar | *Eidolon dupreanum* | NC\_033820 |  | Cameroon | *Eidolon helvum* | 99% | 92% | |
| **OQ818344** | *Sapelovirus* | Madagascar | *Eidolon dupreanum* | NC\_033820 |  | Cameroon | *Eidolon helvum* | 91% | | 78% |
| **OQ818345** | *Sapovirus* | Madagascar | *Rousettus madagascariensis* | KX759623 |  | Cameroon | *Eidolon helvum* | 9% | | 10% |
| **OQ818347** | *Sapovirus* | Madagascar | *Rousettus madagascariensis* | KX759623 |  | Cameroon | *Eidolon helvum* | 66% | | 28% |
| **OQ818348** | *Sapovirus* | Madagascar | *Rousettus madagascariensis* | KX759623 |  | Cameroon | *Eidolon helvum* | 43% | | 31% |