



#### Taxonomy

- |  |                   |
|--|-------------------|
| Autographiviridae; Unclassified family | Actinomycetota    |
| Caudoviricetes; Unclassified order     | Bacillota         |
| Demerecviridae; Unclassified family    | Bacillota_A       |
| Fuselloviridae; Unclassified kingdom   | Bacteroidota      |
| Herelleviridae; Unclassified family    | Chloroflexota     |
| Iridoviridae                           | Deinococcota      |
| Kyanoviridae; Unclassified family      | Desulfobacterota  |
| Mimiviridae                            | Methanobacteriota |
| Peduviridae; Unclassified family       | Planctomycetota   |
| Phycodnaviridae                        | Pseudomonadota    |
| Rountreeviridae; Unclassified family   | Thermoproteota    |
| Unclassified                           | Verrucomicrobiota |

#### Prediction

- Hi-C
- IPHoP
- both

#### Domain

- Archaea
- Bacteria
- Phage lysogenic
- Phage lytic
- Phage unclear

#### Bat host species

- unknown
- Phyllostomus hastatus*
- Mimon crenulatum*

#### Sequence count

- 20
- 40
- 60