**Table S7. Homology among L open reading frame sequences within the BWAV/PGAV clade**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | **Nucleotide Identity (%)** | | | | |
| **Amino acid identity (%)** |  | **BWAV**  **(M459)** | **BWAV**  **(UgAr 1888)** | **PGAV**  **(SA AR 1)** | **PGAV**  **(191B-07)** |
| **BWAV**  **(M459)** |  | **94.6** | **70.8** | **70.8** |
| **BWAV**  **(UgAr 1888)** | **97.6** |  | **70.9** | **71.1** |
| **PGAV**  **(SA AR 1)** | **75.1** | **75.3** |  | **98.2** |
| **PGAV**  **(191B-07)** | **75.1** | **75.2** | **99.5** |  |