**Table S5. Homology among N and NSs 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)** |  | **97.7 / 99.3** | **80.1 / 87.8** | **80.7 / 88.2** |
| **BWAV**  **(UgAr 1888)** | **99.1 / 98.9** |  | **80.3 / 88.5** | **80.9 / 88.9** |
| **PGAV**  **(SA AR 1)** | **86.0 / 73.1** | **86.4 / 73.1** |  | **99.4 / 99.6** |
| **PGAV**  **(191B-07)** | **86.0 / 74.2** | **86.4 / 74.2** | **100 / 98.9** |  |

\*Values shown for N (ORF) / NSs (ORF)