**Supplementary Table S1.** Distance matrix of multiple alignments of amino acid sequences from ORF1, calculated in relative numbers using the “Simple Similarity” option using UGENE.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | **NODE99\_E3** | **NODE35\_E3** | **NODE28\_E3** | **NODE58\_E3** | **NODEA508\_E2** | **NODEA98\_E3** | **NODEA297\_E3** |
| NODE99\_E3 | 100% |  |  |  |  |  |  |
| NODE35\_E3 | 40% | 100% |  |  |  |  |  |
| NODE28\_E3 | 34% | 36% | 100% |  |  |  |  |
| NODE58\_E3 | 44% | 36% | 32% | 100% |  |  |  |
| NODEA508\_E2 | 99% | 40% | 33% | 41% | 100% |  |  |
| NODEA98\_E3 | 27% | 27% | 25% | 26% | 25% | 100% |  |
| NODEA297\_E3 | 28% | 28% | 25% | 27% | 25% | 100% | 100% |