# Supplementary information

**Supplement Table S3.** Percentage of amino acid identity of coding sequence of the E1 and E2 glycoprotein

|  |
| --- |
| **E1 glycoprotein** All four strains: 428/439 identical positions, **97.5% identity** |
| **Strain** | **Thailand 1958** | **Thailand 1975** | **New Caledonia 2011** |
| **Thailand 1975** | 438/439; 99.8% |  |  |
| **New Caledonia 2011** | 435/439; 99.1% | 436/439; 99.3% |  |
| **Cambodia 2011** | 430/439; 97.9% | 429/439; 97.7% | 429/439; 97.7% |
|  |
| **E2 glycoprotein** All four strains: 398/420 identical positions, **94.8% identity** |
| **Strain** | **Thailand 1958** | **Thailand 1975** | **New Caledonia 2011** |
| **Thailand 1975** | 418/420; 99.5% |  |  |
| **New Caledonia 2011** | 414/420; 98.6% | 419/420; 99.8% |  |
| **Cambodia 2011** | 402/420; 95.7% | 402/420; 95.7% | 400/420; 95.2% |