Supplementary Information

**Discovery and genetic characterization of diverse smacoviruses in Zambian non-human primates**

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Table S1. List of primers used to examine the PCR prevalence of detected CRESS DNA viruses

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| **No.** | **Primer name** | **Sequences (5’ to 3’)** |
| 1. | ssDNAV-1F | ATGGTGCCTCTAGGGTCTATTATCC |
| 2. | ssDNAV-1R | AATCTCCGACAATCGARGGCTGAC |
| 3. | ssDNAV-2F | GGTCAGCGTACTGTTGTTCAGGAGA |
| 4. | ssDNAV-2R | TGATAAAAGACTCATCACGCCATTC |
| 5. | ssDNAV-3F | CCGATCGTTTATTCTGTTGTCTCTG |
| 6. | ssDNAV-3R | GAAGAGCATAATACATGCTGAACTG |
| 7. | ssDNAV-4F | GTCCGTATCCTGCTTTGCCTACC |
| 8. | ssDNAV-4R | AGCGCCAGGACGATTAATAGCAAC |
| 9. | ssDNAV-5F | CGATTATGCTTTGTCGACCCAATCC |
| 10. | ssDNAV-5R | AAATAGGACATCCCCGTTTCAGG |
| 11. | ssDNAV-6F | GTAATTCTTATCATGCGACAATGCC |
| 12. | ssDNAV-6R | GGAGAAGCATTACCACGCATACCC |