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Human Herpes Simplex Virus Type 1 in Confiscated Gorilla

Technical Appendix

Technical Appendix Figure. Sequence alignment of the regions of *UL30* from the gorilla amplicon, HSV-1, HSV-2, MaHV-1, PaHV-2, and VZV with a majority sequence presented representing the amino acid found in most of the individual sequences. Differences from the majority sequence are underlined. The 3 aa that are boxed (aa 92–94) represent a cluster of amino acids (VAL) unique to the gorilla amplicon and not found in other alphaherpesviruses, particularly the V at position 92. However, the following 3 aa are found at the corresponding region in other HHVs: HHV-4 (LAI), HHV-5 (MAL), HHV-6 (LAL), and HHV-8 (LAI) (not shown). EHV, equid herpesvirus; HSV, herpes simplex virus; HHV, human herpesvirus; MaHV, Macacine herpesvirus; PaHV, Papiine herpesvirus.