Figure S7

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SP|O09800|GN_HHV11      -MGPPRRVCRAGLLFVLLVALAAGD----AGPRGEP-PGEEG-G--------------------RDGIGGARCETQNTGQMSAPGALVPFYVGMAS---------------------- 69
SP|Q86539|GN_HHV2H      MTGKPARLGRWV--VLLFVA---LV----AGVPGEP-PNAAG-A--------------------RGVIGDAQCRGDSAGVVSVPGVLVPFYLGMTS---------------------- 65
SP|Q65ZG0|GN_VZVD       ---------------MG---------------------------SITASFILIT-----MQILF---FC---EDSSGEPNFAERNFWHASCSARGVYIDGSMITTLFFY--ASLL 60
TR|B9VXN1|B9VXN1_HCMVT  --------MEWNTLVLGLLVLSVAASSNHTSTASTPSPSSSTHTSTTVKATTTATTSTTTVTSTTSSTTSTKPGSTHDPNVMRPHAHNDFYKAHCTSHMYELSLSSFAAWWTMLNALIL 112
SP|P0C6Z3|GN_EBVA8      ---------------MGKVLR-------------K--PFA-------KAVPLLFLAATWL---LTGVLPAGASSPTNAAAASLTEAQDQFYSYTCNADTFSPSLTSFASIWALLTLVLV 79
                                                    2              1                                    2                               1
                                                                                           .  *:    .


SP|O09800|GN_HHV11      MG-VCIIAHVCQICQRLLAAGHA---- 91
SP|Q86539|GN_HHV2H      MG-VCMIAHVYQICQRALAAGSA---- 87
SP|Q65ZG0|GN_VZVD       GVCVALISLAYHACFRLFTRSVLRSTW 87
TR|B9VXN1|B9VXN1_HCMVT  MGAFCIVLR--HCCFQNFTATTTKGY- 136
SP|P0C6Z3|GN_EBVA8      IIASAIYLM--YVCFNKFVNTLLTD-- 102
                          .:          *  . :.
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