205F

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 0-PB1.1 MRVMGMSRNY QQWWIWGILG FWMLMICSVM GNLWVTVYYG VPVWREAKTT LFCASDAKAY EREVHNVWAT HACVPTDPNP QEMFLKNVTE NFNMWKNDMV  100
 0-PL4.1 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 8-PB2.3 ---------- ---------- ---------- ---------- --------P- ---------- ---------- ---------- ---------- ----------
14-PB5.4 ---------- -------T-- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
20-PB2.2 ---------- -------V-- --------GV ---------- ---------- ---------- ---------- ---------- -----E---- D---------
20-PB5.2 ---------- ---------- --------G- ---------- ---------- ---------- ---------- ---------- -----E---- ----------
26-PB2.2 ---------- ---------- --------G- ---------- ---------- ---------- ---------- ---------- ---------- D---------

                                                                  V1                                                  V2
 0-PB1.1 DQMHEDIISL WDQSLKPCVK LTPLCVTLSC SNYSNCNDTY S......... .TANCTSGGE IKNCSFNATT EIRDKNRKEY ALFYRPDIVP LKPNDSNSRE  200
 0-PL4.1 --------N- ---------- ---------- ---------- -......... .--------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- ---N------ -......... .--------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------- ---------P -......... .--------- ---------- ---------- ---------- ----------
 8-PB2.3 ---------- ---------- ---------- ---------- -N.....DTN SI-------- ---------- ---------- ---------- --------S-
14-PB5.4 ---------- ---------- --------N- -D------NS T......... ..D------- ---------- ---------- ---------- --------S-
20-PB2.2 ---------- ---------- ---------- ---------D -N...YNKTY NK-------- ---------- ---------- ---------- -----N--S-
20-PB5.2 ---------- ---------- ---------- ---------- -NGTYSNDTN S-V----R-- ---------- ---------- ---------- ----EG--S-
26-PB2.2 ---------- ---------- --------N- -D---G-G-N N......... .---Y----- ---------- ---------- ---------- -----NK-S-

 0-PB1.1 YILINCNTST IAQACPKVSF DPIPIHYCAP AGYAILKCND NKTFNGTGPC HNVSTVQCTH GIKPVISTQL LLNGSLAEED IIIRSENLAN NVKTIIVHLN  300
 0-PL4.1 ---------- ---------- ---------- ---------N ---------- ---------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 8-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
14-PB5.4 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
20-PB2.2 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
20-PB5.2 ---------- ---------- ---------- ---------- ---------- R--------- ---------- ---------- ---------- ----------
26-PB2.2 -L-------- ---------- ---------- ---------- ---------- R--------- ---------- ---------- --------T- ----------

                                V3                                                  α2
 0-PB1.1 ESVEINCTRP NNNTSRGIRI GPGQTFFATG RIIGNIRQAY CSINKSKWND TLQKIKGKLQ EHFPNKTIHF EPHSGGDLEI TTHSFNCRGE FFYCNTSELF  400
 0-PL4.1 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 8-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
14-PB5.4 ---------- ---------- ---------- ---------- ----R----- -----R---- ---------- K--------- ---------- ----------
20-PB2.2 ---------- ---------- ---------- ----D----- ---------- -----E---- --------Q- ---------- ---------- ----------
20-PB5.2 ---------- ------SV-- ---------- ----D----- ----R----- -----E---- --------D- ---------- ---------- ----------
26-PB2.2 ---------- ---------- ---------- ----D----H -N-T------ ---R------ --------Q- K--------- ---------- ----------

                    V4                                                              V5
 0-PB1.1 NISRLNSTSS IITLPCRIKQ FINMWQKVGR AMYAPPIEGK ITCNSSITGL LLTRDGGNNT NNTETFRPGG GDMRDNWRSE LYKYKVVEIK PLGIAPTGSK  500
 0-PL4.1 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- --A------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 8-PB2.3 ---G------ ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
14-PB5.4 ---------- N---T----- ---------- ---------- ---------- ---------- -D-------- ---------- ---------- ----------
20-PB2.2 ------K--P ---------- ---------- ---------- ---------- --------T- ---------- ---------- ---------- ----------
20-PB5.2 ---------P ---------- ---------- ---------- ---------- ---------- ST-------- ---------- ---------- ----------
26-PB2.2 ..T------P ---------- ---------- ---------- ---------- ---------S S.N------- ---------- ---------- ----------

 0-PB1.1 RAVVEREKRA VGLGAVFLGF LGAAGSTMGA ASITLTVQAR QVLAGIVQQQ SNLLRAIEAQ QHMLQLTVWG IKQLQARVLA IERYLKDQQI LGIWGCSGKL  600
 0-PL4.1 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
 8-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
14-PB5.4 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
20-PB2.2 ----K----- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
20-PB5.2 ----Q----- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
26-PB2.2 ----K----- ---------- ---------- ---------- ---------- -S-------- ---------- ---------- ---------- ----------

 0-PB1.1 ICTTAVPWNA SWSNKSQEDI WNNTTWRQWD REISNYTSII YNLLEESQNQ QERNEKDLLA LDSWKNLWNW FDISRWLWYI KIFIIIVGGL IGLRIIFAVL  700
 0-PL4.1 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----M----- ----------
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----M----- ----------
 2-PB2.3 ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----M----- ----------
 8-PB2.3 ---------- ---------- ---M------ ---------- ---------- ---------- ---------- ---------- ----M----- ----------
14-PB5.4 ---------- ---------- ---M--M--- ---------- ---------- ---------- ---------- ---------- ----M----- ----------
20-PB2.2 ---------- ---------- ---M--M--- ---------- ---------- ---------- ---------- ---------- ----M----- ----------
20-PB5.2 ---------- ---------- ---M--M--- ---------- ---------- ---------- ---------- ---------- ----M----- ----------
26-PB2.2 ---------- -------D-- ---M--M--- ---------- ---------- ---------- ---------- -N-------- ----M----- ----------

 0-PB1.1 SIVNRVRQGY SPLSFQTLTP NPRELDRLGR VEEEGGEQDR DKSIRLVNGF LALFWDDLRS LCLFSYHHLR DFILIAARVV ELLGRRGWEA LKYLGSLVQY  800
 0-PL4.1 ---------- ---------- ---------- I--------- ---------- --------- ---------- ---------- ---------- ----------
 0-PL6.3 ---------- ---------- ---------- I--------- ---------- --------- ---------- ---------- ---------- ----------
 2-PB2.3 ---------- ---------- ---------K I--------- ---------- --------- ---------- ---------- ---------- ----------
 8-PB2.3 ---------- ---------- ---------- I--------- ---------- --------- ---------- --V------- ---------- ----------
14-PB5.4 ---------- ---------- ---------K I--------- ---------- --------- ---------- ---------- ---------- ----------
20-PB2.2 ---------- ---------- ---------- I--------- ---------- --------- ---------- ---------- ---------- ----------
20-PB5.2 ---------- ---------- ---------- I--------- ---------- --------- ---------- ---------- ---------- ----------
26-PB2.2 ---------- ---------- ---------- I--------- ---------- --------- ---------- ---------- ---------- ----------

 0-PB1.1 WGLELKKSAI SLFDTIAIAV AEGTDRIIEL IQRFGRGILN IPTRIRQGLE AALQ 854
 0-PL4.1 ---------- ---------- ---------- ---------- ---------- ----
 0-PL6.3 ---------- ---------- ---------- ---------- ---------- ----
 2-PB2.3 ---------- ---------- --------D- ---------- ---------- ----
 8-PB2.3 ---------- ---------- ---------- ---------- ---------- ----
14-PB5.4 ---------- --L------- ---------- ---------- ---------- ----
20-PB2.2 ---------- --L------- ---------- ---------- ---------- ----
20-PB5.2 ---------- --L------- ---------- ---------- -L-------- ----
26-PB2.2 ---------- --L------- ---------- ---------- ---------- ----
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