

NTDB id 1029 TT RS02230 WP 011228203.1MPVYQYKARDQGR.LVEATTEAEDLRATAARLRDRGLFVAETIKPEPGKGLQAEVRIPALERGPGGLK 65
NTDB id 1402 DSB67 RS12670 WP 010643257.1 .MKT.....IAPQLKNYRMKGINSCKK.TSGNYLAMTEIEVRERLDAQHTIKIKLKKGSISFITK....ISHRVKKGK 67
NTDB id 1169 A1552VC RS11075 WP 000648511.1 MKAT.....QTLPLKNYRMKGINSNGKK.VSGQMLAISEIEVRDKLKDQHIQIKLKKGSVSLAR....LTHRVKSK 68
NTDB id 1113 AAA85695.1 219..1451(.) .MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRVK.....T....SSKRKITQE 70
NTDB id 1112 NFGF RS09215 WP 003689811.1 .MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRVK.....T....SSKRKITQE 70
NTDB id 1252 GC085 RS07730 WP 011213805.1 .MDK.....NSPPLTLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKQGVITNKVVKRRKPLFDR....KNKKITQA 67
NTDB id 1016 ACIAD RS01680 WP 004920476.1 MAAK.....KTQVMVPVFAYEGVDRKGKIK.LKGELPARNMALAKVTLRKQGITTRTIREKRNIFEG....LLKKKVKPL 69
NTDB id 1059 ABD1 RS01610 WP 000279215.1 MAVK.....KAQMMPTFAYEGVDRKGVK.LKGELPAKNMALAKVTLRKQGVTVRNIREKRNILEG....LFFKKVTTL 69
NTDB id 1001668 ABEP14 RS27185 WP 003107299.1 .MAD.....KALKTSVFVWEGTDRKGT.KVKGELSSQNPTLVKAQLRKQGITPVKVRKKGISLLG.....AGKKIKPM 66
NTDB id 1198 PSJM300 03950 AFN76868.1 .MAQ.....KAIKNSVFTWEGTDKQGA.K.LKGELSGVSPALVKAQLRKQGINPQKVRKKGISLFS.....AGKKIKPM 66
consensus **** * ** ** ** ** * * * * * ! * * * * * * * * * *

[illegible][illegible]

