

NTDB id 1029 TT RS02230 WP 011228203.1MPVYQYKARDRQGRLVEATTEAEDLRTAARL LRDRG LFVAETKEPGKGGLQAEVRIPALER GPGLK 65
NTDB id 1402 DSB67 RS12670 WP 010643257.1 .MKT.....IAPQLKNYRWKGINSSGKK.TSGNVLAMTEIEVERERLDAQHTIKIKKLKKSISFITK....ISHRVK GK 67
NTDB id 1169 A1552VC RS11075 WP 000648511.1 MKAT.....QTLP LKNYRWKGIN SNGKK.VSQMLAISIEIVRD KLDQHIIQIKKLKKSVSLLAR....LTHR VKITQE 68
NTDB id 1113 AAA85695.1 219..1451() .MAKNGGFSLF AKKEKR FIFEGRH SASDKLVNGEVS AFTEEEEARKKLAKRGIRPLQITRVK.....T....SSK RKITQE 70
NTDB id 1112 NFGF RS09215 WP 003689811.1 .MAKNGGFSLF AKKEKR FIFEGRH SASDKLVNGEVS AFTEEEEARKKLAKRGIRPLQITRVK.....T....SSK RKITQE 70
NTDB id 1252 GC085 RS07730 WP 011213805.1 .MDK.....NSPPTLT FH YCGINKAGQK.MEGDIQARSLAI AKADLRKQGIVTNKV VVKRRKP LFD R....KNKK ITQA 67
NTDB id 1016 ACIAD RS01680 WP 004920476.1 MAAK.....KTQVM PVFAYEGV DRKG I.KLGELPARNMALAKVT LRKQGITT RTIREKRKN IFEG...LLK KVV KPL 69
NTDB id 1059 ABD1 RS01610 WP 000279215.1 MAVK.....KAQMNP TFAYEGV DRKG VK.IKGELPAKNMALAKVT LRKQGVTVRNIREKRKNILEG...LFK KVV TTL 69
NTDB id 1069272 ACFELM RS04510 WP 023103506.1 .MAD.....KALKTSVF IWE GTD KKGS.VKGE LTGCNPMLVKAQ LRKQGINPLKVRKKIGITLFG.....AGKK IKPM 66
NTDB id 1198 PSJM300 03950 AFN76868.1 .MAQ.....KAIKNSVFT WEG LD RGAK.VKGE SGVSPALVKAQ LRKQGINPQKVRKK SISLFS.....AGKK IKPM 66

consensus *** * * * ** * * * * ! * * * * * * * * *

[illegible][illegible]

