

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGRLVEATIEAEDLRTAARLRDRGLFVAETKEPGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPKNRYRWKGINSNGKKVSGQMLAISEIEVRDKLKDQHIIQIKKLKXGSVSLLAR....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAQPLKNRYRWKGINSNGKKTSGNVLAMTIEIEVERLERDAQHIIKIKKLKXGSISFITK....ISH	62
NTDB id 1112	NGAF RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIIEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTS...SSKR	65
NTDB id 1113	AAAB5695.1 219..1451()	MAKNGGFSLF AKKEKR FIIEGRHSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTS...SSKR	65
NTDB id 1038914	ACCP93 RS14605 WP 033836749.1	MSVARNVIKKQPADRSNSQM QP FVWEGTD XRG I K MKGEQAARNVNMLRAELRRQGITPTV VXP KP K PLFGA....AGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLLT FH YQGINKAGQKMEGD IQARSLAI AKA D LRKQGIVTNKVVKRK K PLFDR....KNK	62
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKA I K NSV FTWEGLDRQGA K IKGELSGVSPALVKAQLRKQGINPQKV R KK SISLFS....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVM PV FAYEGVDRKG I K LK GEL P ARNMALAKVT LRKQGITIRTIREKRKN IFEG....LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKK AQMMPTFA YEGVDRKG I K LK GEL P AKNMALAKVT LRKQGVTVRNIREKRKN ILEG....LFKK	64
consensus		* * * * * * * * * * ! * * * * * * * * *	

[illegible][illegible]

