

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.LVEATIEAEDLRTAARLRDRGLFVAEIKPEPKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKNYRWKGINSNGKK.VSGQMLAISEIEVRDKLKDQHIQIKKLLKKGVSLLAR.....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIIAPQLKNYRWKGINSNGKK.TSGNVLAMATEIEVRERLDAQHIIKKLLKKGVSISFITK.....ISH	62
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSFLAKKEKRFYFEGRHSASDKLVNGEVSATTEEEARKKLAKRGIPLQITRV.....KTS...SSKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSFLAKKEKRFYFEGRHSASDKLVNGEVSATTEEEARKKLAKRGIPLQITRV.....KTS...SSKR	65
NTDB id 159126	APY30 RS17040 WP 033836749.1	MSVARNVIKKQPADRSNSQMMPFVWEGTDKRGIK.MKGEQAARNVNNMLRAELRRQGITPTVVKPKPKPLFGA.....AGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLLT FHYQGINKAGQK.MEGDIQARSLAIKADLRKQGIVTNKVVKKRKPLFDR.....KNK	62
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKAIKNSVFTWEGLDKRGQAK.IKGELSGVSPALVKAQLRKQGINPQKVRKKKSISLFS.....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVYFAYEGVDRKGIK.LKGELPARNMALAKVTLRKQGITRTIREKRKNIFEG....LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKAMMPYFAYEGVDRKGIVK.IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG....LFKK	64
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NTDB id 1029	TT RS02230 WP 011228203.1	GPGLKDLAIFSRQLATMLGAGLTLLQALAILERQTENRKFREILKQVRTDVEGGMAFSEALSKHK.IFSRLVYNLVRAGE	139
NTDB id 1169	A1552VC RS11075 WP 000648511.1	RVKSKDITILTRQLATMLTTGVPIVQAALKLVGDNHRKAEMKSILAQITKSVEAGTPLSKAMRTASAHFDTLVYDLVETGE	143
NTDB id 1402	DSB67 RS12670 WP 010643257.1	RVGKGKDITIFTRQTATMLMTGVPIVQAALKLVSENHKKAEEMKSILMSVTRAVEAGTPMSKAMRTASTHFDPPLYTDLTIATGE	142
NTDB id 1112	NGFG RS09215 WP 003689811.1	KITQEDITVFTRQLSTMINKAGLPMLQAFEIVARGHGNSPSTEMLEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGE	145
NTDB id 1113	AAA85695.1 219..1451()	KITQEDITVFTRQLSTMINAGLPLMQAFEIVARHGHNPSMTEMLEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGE	145
NTDB id 159126	APY30 RS17040 WP 033836749.1	KVGPKDIAFFSRQMATTMMKSGVPVIGSLEIIGEGHKNPRMRKMVGQIRTDIEGGSSLHEATISKHPVQFDELYRNLVRAGE	154
NTDB id 1252	GC085 RS07730 WP 011213805.1	KITQADITVFSRQLATMIESGIPLVQAFDIVAKGQSNKRCLKDLIETIKHDIETGLTLAESLIKHPLYFNELFCNLVDAGE	142
NTDB id 1198	PSJM300 03950 AFN76868.1	KIKPMDIALFTRQMATTMMKAGVPLQSFDIIGEGFDNPNNMRKLVDLKKQEVAAGNSFASSLRKKPPQYFDDLYCNLVDSDGE	141
NTDB id 1016	ACIAD RS01680 WP 004920476.1	KVKPLDIAIFTRQLATMMKAGVPLVQAFEIVAEGLENPAMRDVVVLGLKNEVEGGNTFAGALRKYPQYFDNFLFCSLTESGE	144
NTDB id 1059	ABD1 RS01610 WP 000279215.1	KVTTLDTITIFTRQLATMMKAGVPLVQFEIVAEGLENPAMREVVLGIKGEVEGGSTFASALRKYPQHFDNFLFCSLVESGE	144
consensus		** !*****!****!*!!!!!!*****!*****!*****!*****!*****!*****!*****!*****!	

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consensus

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