

[illegible]

NTDB id 1292	VP RS12235 WP 005479693.1	TLAQLGVDVAAN.....DL....GAIDSALAGGSPTLTFTFDADSSIDGGILT.....	115
NTDB id 1184	GTF74 RS02335 WP 000649326.1	GFTRLGTVEDMG.....DGKIVTAPTASGALG..GTIKYTFDAGV.VSSSKIQ.....	122
NTDB id 1167	A1552VC RS11065 WP 000649326.1	GFTRLGTVEDMG.....DGKIVTAPTASGALG..GTIKYTFDAGV.VSSSKIQ.....	122
NTDB id 1160	ABDM36 RS02330 WP 000649326.1	GFTRLGTVEDMG.....DGKIVTAPTASGALG..GTIKYTFDAGV.VSSSKIQ.....	122
NTDB id 1383	A4U84 RS08530 WP 005711781.1	GKNGVRADKSDVPDAKYIKSTSVASG....VITVAGKCN.....IDGYGYTM	120
NTDB id 1354	HI 0299 AAC21963.1	GKNGIAADIKT..AKGYVASVITQSG....GITVKNGT.....LANMEYIL	115
NTDB id 1365	NTHI RS01955 WP 011271973.1	GKNGIAADIT..AKGYVKSVTTSNG....AITVKGDCG.....LANMEYIL	115
NTDB id 1191	AAC28468.1 484..945()	NVIFDSAN.....GLDGKYF.....APGGVTVPDPT.....GAINIAFDAGAN.AGKTMITPT....	122
NTDB id 1197	PSJM300 03940 AFN76866.1	E...GYIGI..TDSTSYC.DVDLDTA..ADGHIECT.....AKGGNAGKF.....DGKTIT.....	114
NTDB id 1119	NGFG RS11465 WP 017147169.1	GVASPTTDIKGKYVKEVEVKNG....VVTATMLS.....SGVNNEI.....KGKKLSLW.....	116
NTDB id 1109	OK783 RS10670 WP 157147416.1	GVASA..STIKGKYVQKVEVAKG....VVTAAQMAS.....TGVNKEI.....QDKKLSLW.....	115
NTDB id 1188	NELON RS01755 WP 003772458.1	NKTAAAGGGAGGAAAGTQHASKYVKSVKVTEA..SPWDITVAISA....TEANGITGL.....DGNTIRLFPLVNKA	141
NTDB id 1424	RS RS02795 WP 011000517.1	GSSVFTPTKNVANLTIAGTGTVPGQITVTTYTTAAGGGTLALVPT.....A...AGTAL	124
NTDB id 1002	ACIAD RS15100 WP 004923779.1	GTPSTAGSSCVGVQEIS...ASNATTNVATATCGASSAGQIIVTM.....DTTKAK..GANITL	119
NTDB id 25565	SDE RS04555 WP 011467347.1	GLPTQIASKYVSSVNVG...AGGIITVGYKAAAGVT.....A..ASTIIF	111
NTDB id 1258	GC085 RS09745 WP 014844212.1	GYVSPAATPNVQSIATIG...ANGVITITYTAAAGG.....GTIIM	106
NTDB id 1274	LPP RS09530 WP 011214168.1	GYVSPAATPNVASIATIG...ANGVITITYTAAAGG.....GTIVM	106
consensus		*.....*.....*.....*.....*	

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NTDB id 1292 VP RS12235 WP 005479693.1	...FTRDAATGWSCLPTA.....PAIAVDGCS.....	139
NTDB id 1184 GTF74 RS02335 WP 000649326.1	...LARDANGLWTCSTTV.....TSEIAPKGCTAGATIN.	153
NTDB id 1167 A1552VC RS11065 WP 000649326.1	...LARDANGLWTCSTTV.....TSEIAPKGCTAGATIN.	153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1	...LARDANGLWTCSTTV.....TSEIAPKGCTAGATIN.	153
NTDB id 1383 A4U84 RS08530 WP 005711781.1	TPK.FANNNTITWATSCQ.....GADISLFPANFCGQKATP.	154
NTDB id 1354 HI 0299 AAC21963.1	QAKGNAAAGVTWTTTCK.....GTDASLFPANFCGSVTK..	149
NTDB id 1365 NTHI RS01955 WP 011271973.1	QATGNAATGVTWTTTCK.....GTDASLFPANFCGSVTQ..	149
NTDB id 1191 AAC28468.1 484..945()	.VIAASGQISTWTCAPG.....AANGIETRRLPTSCQ*.....	153
NTDB id 1197 PSJM300 03940 AFN76866.1	...LNRTADGEWSCAST.....LDAKYKPGKCS.....	139
NTDB id 1119 NGFG RS11465 WP 017147169.1	..GRRENGSVKWFQGPVTRADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK	166
NTDB id 1109 OK783 RS10670 WP 157147416.1	..AKRQDGSVKWFCGQPVRTGDNDTTVA..DANNAIDTKHLPSTCRDESSAT.	164
NTDB id 1188 NELON RS01755 WP 003772458.1	VPAATSTGAIDWACASAAKETADKRFGTGSQVAVGTLPKYAPSECK.....	188
NTDB id 1424 RS RS02795 WP 011000517.1	PVSSAPSGPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAECR.....	168
NTDB id 1002 ACIAD RS15100 WP 004923779.1	TPT.YASGAVTWKCTTT.....SDKKYVPSECRG.....	147
NTDB id 25565 SDE RS04555 WP 011467347.1	RPT.FATGTVDWTCCTGG.....TVQDSYRPSNCR.....	139
NTDB id 1258 GC085 RS09745 WP 014844212.1	TPTLQANGDVTWCTTGG.....TLLAKYRPASCRP.....	136
NTDB id 1274 LPP RS09530 WP 011214168.1	TPTLQANGDVTWCTTGG.....TLLAKYRPASCRP.....	136
consensus	* * ! * * *	

 non conserved
 similar
 ≥ 50% conserved