



logo

NTDB id 1184	GTF74 RS02335 WP 000649326.1	...R.DANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1167	A1552VC RS11065 WP 000649326.1	...R.DANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1160	ABDM36 RS02330 WP 000649326.1	...R.DANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1038	HON27 RS01510 WP 000993725.1	...R.STDGAWTCVSD.....VPTEFLPKGCTAGTPTEGAIITSL	149
NTDB id 1197	PSJM300 03940 AFN76866.1	...R.TADGEWSCAST.....LDAKYKPGKCS.....	139
NTDB id 1192	PAKAF RS23870 WP 024947914.1	...RTAADGLWKCTSD.....QDEQFIPKGCSSK.....	150
NTDB id 1038996	ACCP94 RS05195 WP 218533820.1	...DNNGSISWGCNS.....GTTIDQKYLPTS CRTAATP.....	146
NTDB id 1119	NGFG RS11465 WP 017147169.1	...RENCVSKWFCGQPVTRADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK.....	166
NTDB id 1109	OK783 RS10670 WP 157147416.1	...RQDGSVKWFCGQPVTRTGDNDDTVA..DANNAIDTKHLPSTCRDESSAT.....	164
NTDB id 1191	AAC28468.1 484..945( )	...ASGQISTWTCAPGAANG.....IETBRLPTSCQ*.....	153
NTDB id 1400	DSB67 RS12660 WP 010643261.1	TATYASNGVTSSECKDATN.....AVQTSYCP.....	154
NTDB id 1188	NELON RS01755 WP 003772458.1	..AATSTGAIDWACASA AKETADKRFGTGSGVAVGTLPAKYAPSECK.....	188
NTDB id 1424	RS RS02795 WP 011000517.1	VSSAPSCPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAECR.....	168
NTDB id 1002	ACIAD RS15100 WP 004923779.1	...YASGAVTWKCTTT.....S.DKKYVPSECRG.....	147
NTDB id 1274	LPP RS09530 WP 011214168.1	...QANGDVTWICTGG.....TLLAKYRPASCRP.....	136
NTDB id 1258	GC085 RS09745 WP 014844212.1	...QANGDVTWICTGG.....TLLAKYRPASCRP.....	136
consensus		* * !** * * * ! *	

- non conserved
- similar
- ≥ 50% conserved