

<0

logo

NTDB id 1184	GTF74	RS02335	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1167	A1552VC	RS11065	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1160	ABDM36	RS02330	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1038	HON27	RS01510	WP 000993725.1	TDGAWTCVSD.....VPTEFLPKGCTAGTPTEGAITSL	149
NTDB id 1192	PAKAF	RS23870	WP 024947914.1	AADGLWKCTSD.....QDEQFIPKGC	150
NTDB id 1197	PSJM300	03940	AFN76866.1	ADGEWSCAST.....LDAKYKPGKCS	139
NTDB id 1204	PSJM300	03935	AFN76865.1	TDGLWDCTSD.....LDSRYKPGKCL	141
NTDB id 1383	A4U84	RS08530	WP 005711781.1	KFANNNTTWTATSCQGA...DISL.....FPANFCGQKATP	154
NTDB id 1365	NTHI	RS01955	WP 011271973.1	GNAATGVTWTTTCKGT...DASL.....FPANFCGSVTQ	149
NTDB id 1354	HI 0299	AAC21963.1		GNAAAGVTWTTTCKGT...DASL.....FPANFCGSVTK	149
NTDB id 1119	NGFG	RS11465	WP 017147169.1	RRENGSVKWFVCGQPVTRADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK	166
NTDB id 1109	OK783	RS10670	WP 157147416.1	KRQDGSVKWVCGQPVTRTGDNDDTVA..DANNAIDTKHLPSTCRDESSAT	164
NTDB id 1191	AAC28468.1	484..945()		A.....ASGQISTWTCAPGAANG.....IETRRRLPTSCQ*	153
NTDB id 1188	NELON	RS01755	WP 003772458.1	K.AVP.AATSTGATDWACASAAKETADKRFGTGSGVAVGTLPAKYAPSECK	188
NTDB id 1424	RS	RS02795	WP 011000517.1	GTALPVSSAPSGPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAE	168
NTDB id 1002	ACIAD	RS15100	WP 004923779.1	YASGAVTWKCTTT.....S.DKKYVPSECRG	147
NTDB id 184604	A9E93	RS09335	WP 016356941.1	QANGDVTWCTTGG.....TLLAKYRPASCRP	136
NTDB id 1258	GCO85	RS09745	WP 014844212.1	QANGDVTWCTTGG.....TLLAKYRPASCRP	136
NTDB id 1274	LPP	RS09530	WP 011214168.1	QANGDVTWCTTGG.....TLLAKYRPASCRP	136
consensus				* * ! * * *.....* * * *	

- ⓧ non conserved
- ✖ similar
- ⓧ ≥ 50% conserved