

```

NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1
NTDB id 1160 ABDM36 RS02330 WP 000649326.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
NTDB id 1192 PAKAF RS23870 WP 024947914.1
NTDB id 1197 PSJM300 03940 AFN76866.1
NTDB id 1204 PSJM300 03935 AFN76865.1
NTDB id 1191 AAC28468.1 484..945( )
NTDB id 1109 OK783 RS10670 WP 157147416.1
NTDB id 1119 NGFG RS11465 WP 017147169.1
NTDB id 1060269 ACF3M1 RS03335 WP 394003921.1
NTDB id 1424 RS RS02795 WP 011000517.1
NTDB id 1002 ACIAD RS15100 WP 004923779.1
NTDB id 1274 LPP RS09530 WP 011214168.1
NTDB id 1258 GC085 RS09745 WP 014844212.1
consensus

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NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1
NTDB id 1160 ABDM36 RS02330 WP 000649326.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
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NTDB id 1258 GC085 RS09745 WP 014844212.1
consensus

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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.....QDEQFIPKGCCK.....	150
NTDB id 1197	PSJM300	03940	AFN76866.1	...ADGEWSCAST.....LDAKYKPGKCS.....	139
NTDB id 1204	PSJM300	03935	AFN76865.1	...TDGLWDCSTSD.....LDSRYKPGKCL.....	141
NTDB id 1191	AAC28468.1	484..945()		ASGQISTWTCAPG.....AANGIETRRLLPTSCQ*.....	153
NTDB id 1109	OK783	RS10670	WP 157147416.1	RQDCSVKWFQCGQPVTRTGDNDDTVADANNAIDTKHLPSTCRDESSAT.....	164
NTDB id 1119	NGFG	RS11465	WP 017147169.1	RENC SVKWFQCGQPVTRADDDTVADAKDGKEIDTKHLPSTCRDKASDAK.....	166
NTDB id 1060269	ACF3M1	RS03335	WP 394003921.1	TSDQTVSWDCEGT.....VDPYRYPSSCR.....	140
NTDB id 1424	RS	RS02795	WP 011000517.1	APSGPIMWTCYAQNKAQAASS.VAPSGTMSLAAPKYAPAECP.....	168
NTDB id 1002	ACIAD	RS15100	WP 004923779.1	YASCAVTWKCTTT.....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				* !*! * ** ! !	

- X

 non conserved
- X

 similar
- X

 ≥50% conserved