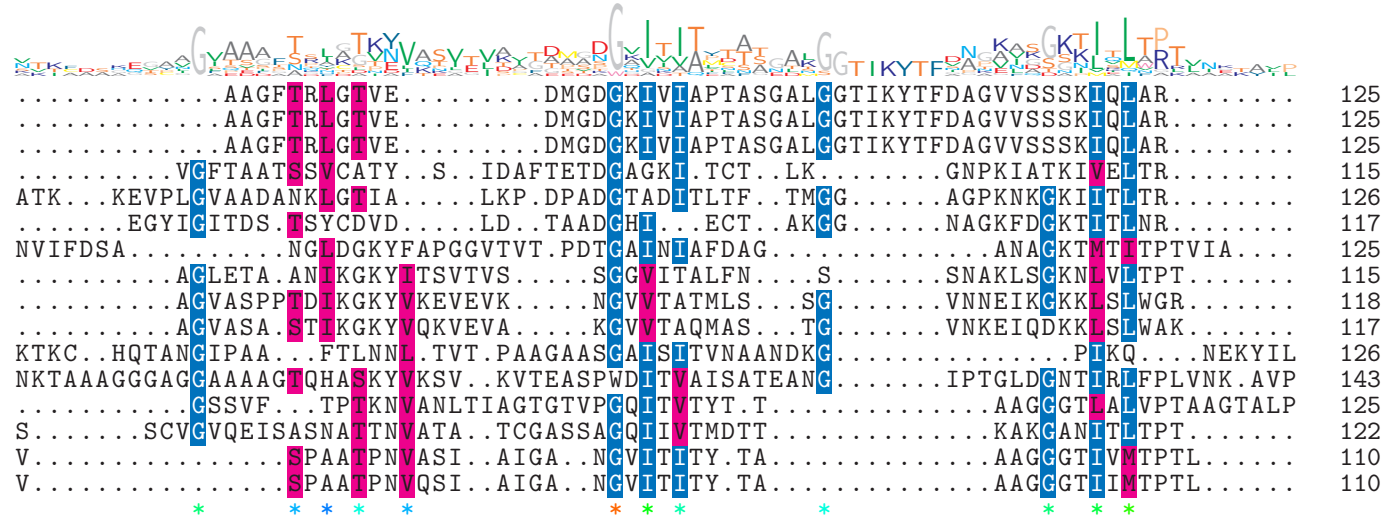
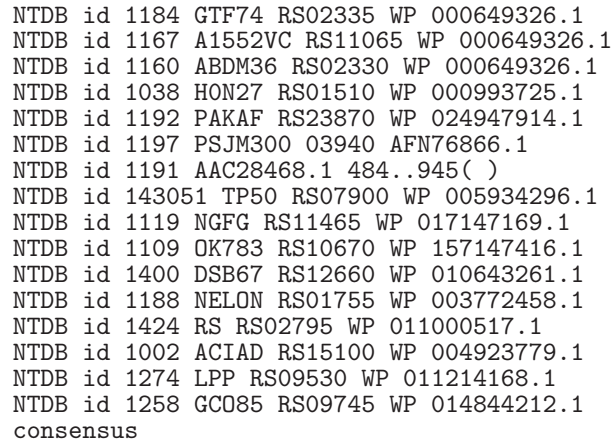


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NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1
NTDB id 1160 ABDM36 RS02330 WP 000649326.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
NTDB id 1192 PAKAF RS23870 WP 024947914.1
NTDB id 1197 PSJM300 03940 AFN76866.1
NTDB id 1191 AAC28468.1 484..945( )
NTDB id 143051 TP50 RS07900 WP 005934296.1
NTDB id 1119 NGFG RS11465 WP 017147169.1
NTDB id 1109 OK783 RS10670 WP 157147416.1
NTDB id 1400 DSB67 RS12660 WP 010643261.1
NTDB id 1188 NELON RS01755 WP 003772458.1
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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<0

logo

NTDB id 1184	GTF74	RS02335	WP 000649326.1	DANGLWTCST.....	TVTSEIAPKGCTAGATIN.....	153	
NTDB id 1167	A1552VC	RS11065	WP 000649326.1	DANGLWTCST.....	TVTSEIAPKGCTAGATIN.....	153	
NTDB id 1160	ABDM36	RS02330	WP 000649326.1	DANGLWTCST.....	TVTSEIAPKGCTAGATIN.....	153	
NTDB id 1038	HON27	RS01510	WP 000993725.1	STDGAWTCVS.....	DVPTFLPKGCTAGTPTEGAITSL	149	
NTDB id 1192	PAKAF	RS23870	WP 024947914.1	TAADGLWKCTS.....	DQDEQFIPKGCSK.....	150	
NTDB id 1197	PSJM300	03940	AFN76866.1	TADGEWSCAS.....	TLDAKYKPGKCS.....	139	
NTDB id 1191	AAC28468.1	484..945()		...ASGQISTWTCAPGA.....	ANGIETRRLPTSCQ*.....	153	
NTDB id 143051	TP50	RS07900	WP 005934296.1	...DNNGSTISWGCST.....	GTIDQKYLPTSCRTAATP.....	146	
NTDB id 1119	NGFG	RS11465	WP 017147169.1	...RENGSVKWF CGQPVT	RADDDTVADA..KDGKEIDT	KHLPSTCRDKASDAK.....	166
NTDB id 1109	OK783	RS10670	WP 157147416.1	...RQDGSVKWF CGQPVT	RTGDNDDTVA..DANNAIDT	KHLPSTCRDESSAT.....	164
NTDB id 1400	DSB67	RS12660	WP 010643261.1	TATYASNGVTWSSECKDATN.....	AVQTSYCP.....	154	
NTDB id 1188	NELON	RS01755	WP 003772458.1	.AATSTGAIDWACASAAKETADKRF	GTGSGVAVGTLPAKYAPSECK.....	188	
NTDB id 1424	RS	RS02795	WP 011000517.1	VSSAPSGPIMWTCYAQNKAQAASSVAPS...	GTMSLAAKYAPAECP.....	168	
NTDB id 1002	ACIAD	RS15100	WP 004923779.1	...YASGAVTWKCTIT.....	S.DKKYVPSECRG.....	147	
NTDB id 1274	LPP	RS09530	WP 011214168.1	...QANGDVTWTCCTGG.....	TLLAKYRPASCRP.....	136	
NTDB id 1258	GC085	RS09745	WP 014844212.1	...QANGDVTWTCCTGG.....	TLLAKYRPASCRP.....	136	
consensus				* * ! * *	** ** ! *		

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved