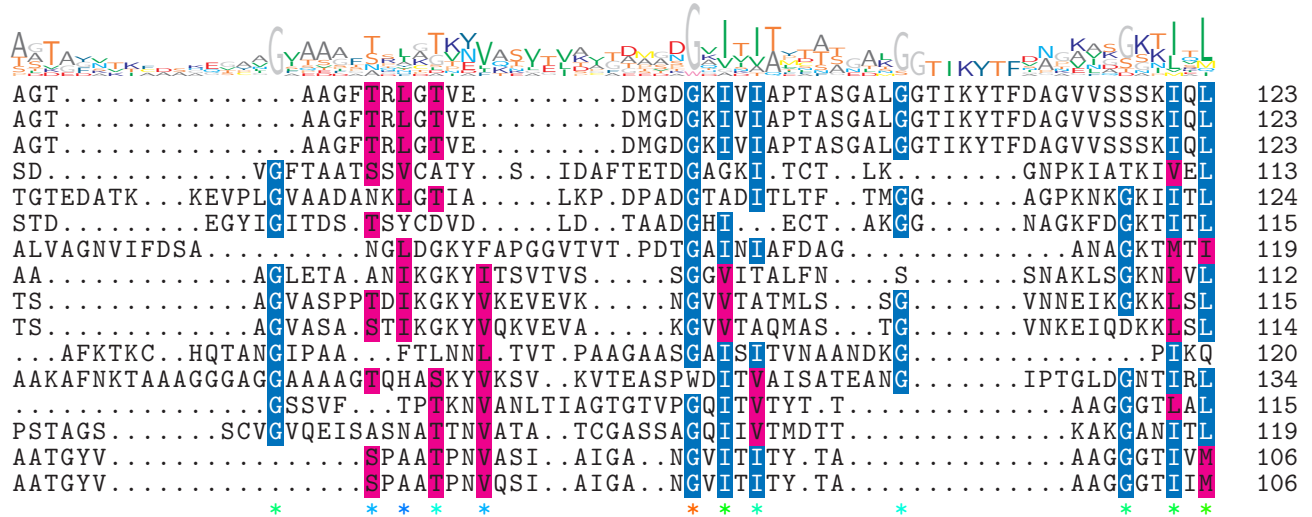
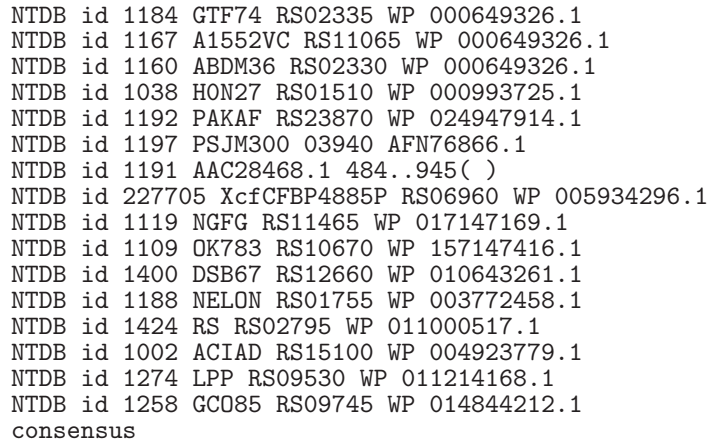


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NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1
NTDB id 1160 ABDM36 RS02330 WP 000649326.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
NTDB id 1192 PAKAF RS23870 WP 024947914.1
NTDB id 1197 PSJM300 03940 AFN76866.1
NTDB id 1191 AAC28468.1 484..945( )
NTDB id 227705 XcfCFBP4885P RS06960 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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logo

Protein	Sequence	Score
AR	DANGLWTCST	153
AR	DANGLWTCST	153
AR	DANGLWTCST	153
TR	STDGAWTCVS	149
TR	TAAIDGLWKCTS	150
NR	TADGEWSCAS	139
TPTVIA	ASGQISTWTCAPGA	153
TPT	DNNGSISWGCST	146
WGR	RENGSVKWF	166
WAK	RQDGSVKWFCGQPVTRTGDNDTVA	164
	NEKYILTATYASNGVTWSSECKDATN	154
FPLVNK.AVP	AATSTCAIDWACASAAKETADKRFGTSGVAVGTLPAKYAPSECK	188
VPTAAGTALPVSSAPSGPIMWTCYAQNKAQAASSVAPS	GTMSLAAKYAPAECR	168
TPT	YASCAVTWKCTIT	147
TPTL	QANGDVTWICTGG	136
TPTL	QANGDVTWICTGG	136

- X non conserved
- X similar
- X $\geq 50\%$ conserved