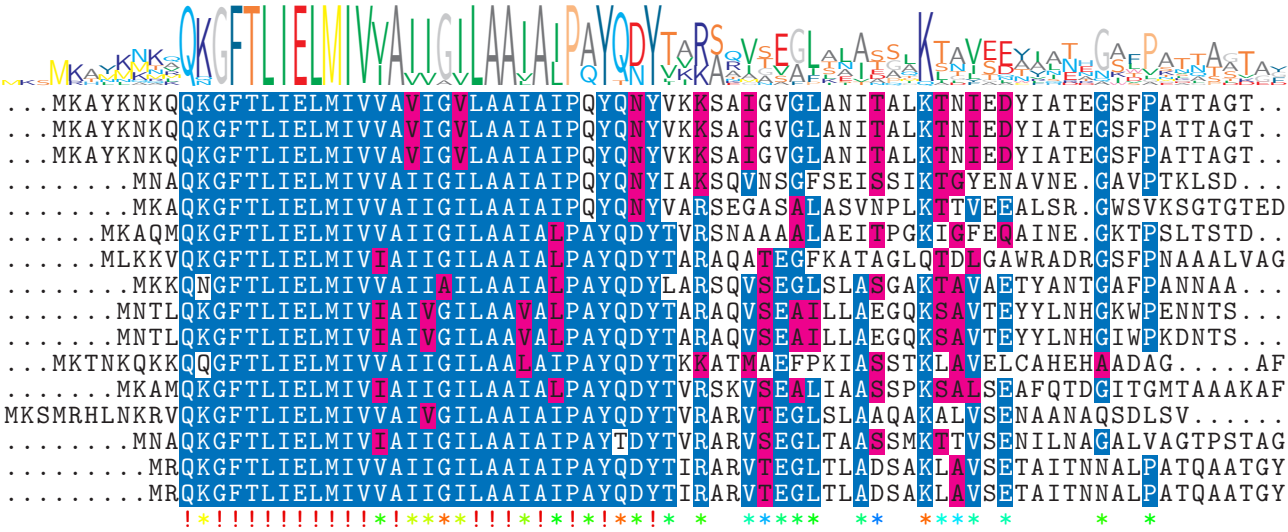


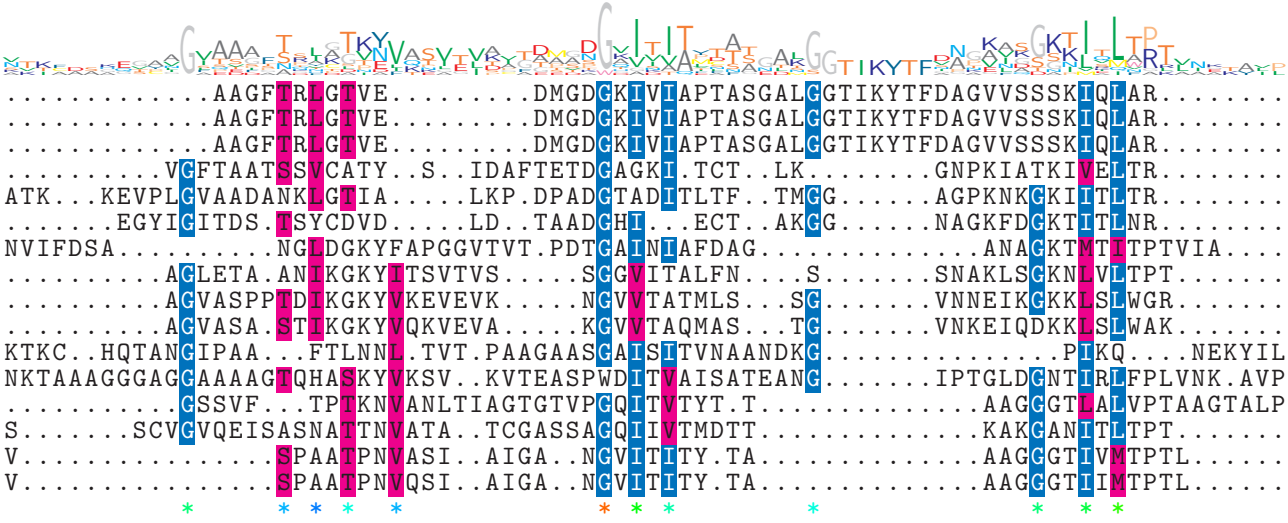
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NTDB id	1184	GTF74	RS02335	WP	000649326.1
NTDB id	1167	A1552VC	RS11065	WP	000649326.1
NTDB id	1160	ABDM36	RS02330	WP	000649326.1
NTDB id	1038	HON27	RS01510	WP	000993725.1
NTDB id	1192	PAKAF	RS23870	WP	024947914.1
NTDB id	1197	PSJM300	03940	AFN76866.1	
NTDB id	1191	AAC28468.1	484..945()		
NTDB id	150873	AC614	RS07105	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	GCO85	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
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NTDB id	1258	GCO85	RS09745	WP	014844212.1
consensus					



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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
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NTDB id 1192	PAKAF	RS23870	WP 024947914.1	TAADGLWKCTS.....	DQDEQFIPKGCSK.....	150
NTDB id 1197	PSJM300	03940	AFN76866.1	TADGEWSCAS.....	TLDKYPKGCS.....	139
NTDB id 1191	AAC28468.1	484..945()		...ASGQISTWTCAPGA.....	ANGIETRRLPTSCQ*.....	153
NTDB id 150873	AC614	RS07105	WP 005934296.1	...DNNGSTISWGCST.....	GTIDQKYLPTSCRTAATP.....	146
NTDB id 1119	NGFG	RS11465	WP 017147169.1	...RENGSVKWF CGQPVT	RADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK.....	166
NTDB id 1109	OK783	RS10670	WP 157147416.1	...RQDGSVKWFCGQPVT	RTGDNDDTVA..DANNAIDTKHLPSTCRDESSAT.....	164
NTDB id 1400	DSB67	RS12660	WP 010643261.1	TATYASNGVTWSSECKDATN.....	AVQTSYCP.....	154
NTDB id 1188	NELON	RS01755	WP 003772458.1	.AATSTGAIDWACASAAKETADKRFGT	SGVAVGTLPAKYAPSECK.....	188
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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				* * ! * *	** ** ! *	

- non conserved
- similar
- ≥ 50% conserved