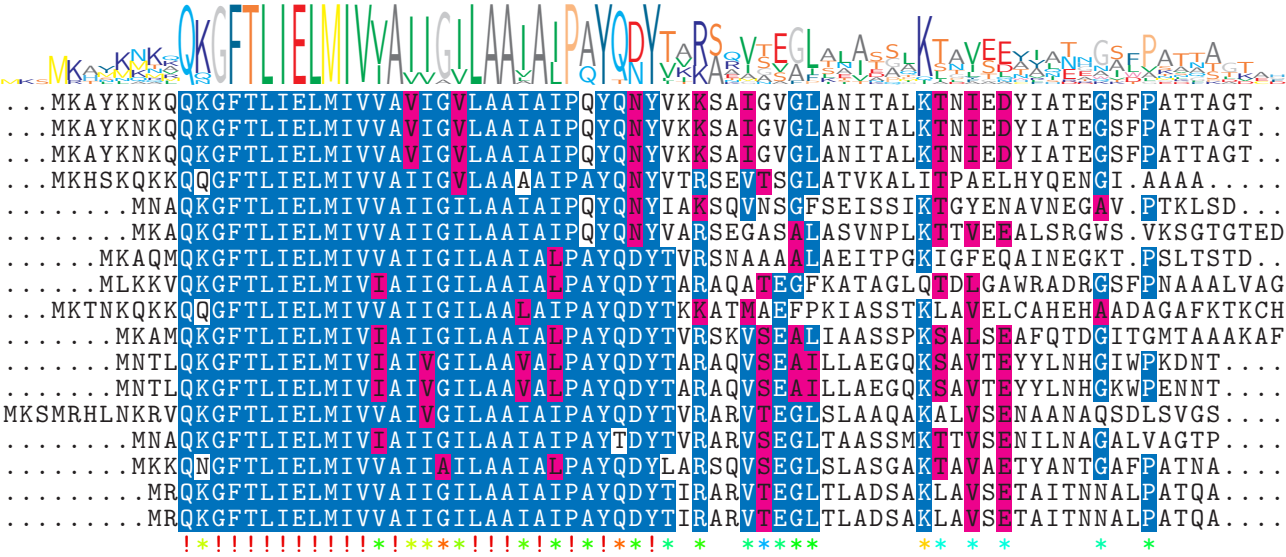


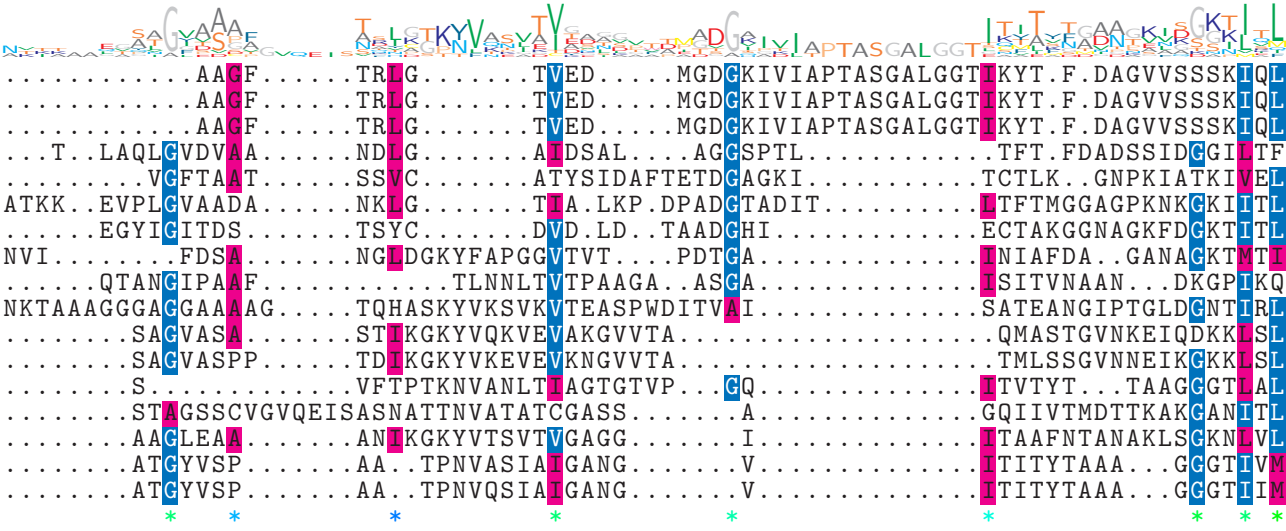
logo

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	1292	VP	RS12235	WP	005479693.1
NTDB	id	1038	HON27	RS01510	WP	000993725.1
NTDB	id	1192	PAKAF	RS23870	WP	024947914.1
NTDB	id	1197	PSJM300	03940	AFN	76866.1
NTDB	id	1191	AAC28468.1	484..945()		
NTDB	id	1400	DSB67	RS12660	WP	010643261.1
NTDB	id	1188	NELON	RS01755	WP	003772458.1
NTDB	id	1109	OK783	RS10670	WP	157147416.1
NTDB	id	1119	NGFG	RS11465	WP	017147169.1
NTDB	id	1424	RS	RS02795	WP	011000517.1
NTDB	id	1002	ACIAD	RS15100	WP	004923779.1
NTDB	id	999957	ABFU70	RS16205	WP	011038214.1
NTDB	id	1274	LPP	RS09530	WP	011214168.1
NTDB	id	1258	GCO85	RS09745	WP	014844212.1
consensus						



logo

NTDB	id	1184	GTF74	RS02335	WP	000649326.1
NTDB	id	1167	A1552VC	RS11065	WP	000649326.1
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NTDB	id	1274	LPP	RS09530	WP	011214168.1
NTDB	id	1258	GCO85	RS09745	WP	014844212.1
consensus						



logo

NTDB id 1184	GTF74	RS02335	WP 000649326.1	AR.DANGLWTCST.....TVTSEIAPKGC	TAGATIN.....	153	
NTDB id 1167	A1552VC	RS11065	WP 000649326.1	AR.DANGLWTCST.....TVTSEIAPKGC	TAGATIN.....	153	
NTDB id 1160	ABDM36	RS02330	WP 000649326.1	AR.DANGLWTCST.....TVTSEIAPKGC	TAGATIN.....	153	
NTDB id 1292	VP	RS12235	WP 005479693.1	TR.DAATGWSCLP.....TAPAIA.VDGC	S.....	139	
NTDB id 1038	HON27	RS01510	WP 000993725.1	TR.STDGAWTCVS.....DVPTEFLPKGC	TAGTPTEGAITS	149	
NTDB id 1192	PAKAF	RS23870	WP 024947914.1	TRTAADGLWKCTS.....DQDEQFIPKGC	SK.....	150	
NTDB id 1197	PSJM300	03940	AFN76866.1	NR.TADGEWSCAS.....TIDAKYKPGKCS		139	
NTDB id 1191	AAC28468.1	484..945()		TPTVI.....AASGQISTWTCAPG.....AANGIETRRLPTSCQ*		153	
NTDB id 1400	DSB67	RS12660	WP 010643261.1	NEKYILTATYASNGVTWSSECKD.....ATNAVQTSYCP		154	
NTDB id 1188	NELON	RS01755	WP 003772458.1	FPLVNKA.VP.AATSTCAIDWACASAAKETADKRFGTGS	GVAVGTLPAKYAPSECK.....	188	
NTDB id 1109	OK783	RS10670	WP 157147416.1	WA.....KRQDCSVKWFCCGQPVRTGTGDNDDTVAD..ANNAIDTKHLPSTCR	DESSAT.....	164	
NTDB id 1119	NGFG	RS11465	WP 017147169.1	WG.....RRENGSVKWFCCGQPVT	RADDDTVADAK..DGKEIDTKHLPSTCRDKASDAK.....	166	
NTDB id 1424	RS	RS02795	WP 011000517.1	VPTAAGTALPVSSAPSGPIMWTCYAQNKAQAAS.SVAPS..GTMSLA	AAKYAPAECCR.....	168	
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NTDB id 999957	ABFU70	RS16205	WP 011038214.1	TP.....TDNNGSISWGC	TNG.....TTIDQKYLPTS	CRTAAAP.....	146
NTDB id 1274	LPP	RS09530	WP 011214168.1	TPT.....LQANGDVTWTC	TGG.....TLL.AKYRPAS	CRP.....	136
NTDB id 1258	GCO85	RS09745	WP 014844212.1	TPT.....LQANGDVTWTC	TGG.....TLL.AKYRPAS	CRP.....	136
consensus							

- non conserved
- similar
- ≥ 50% conserved