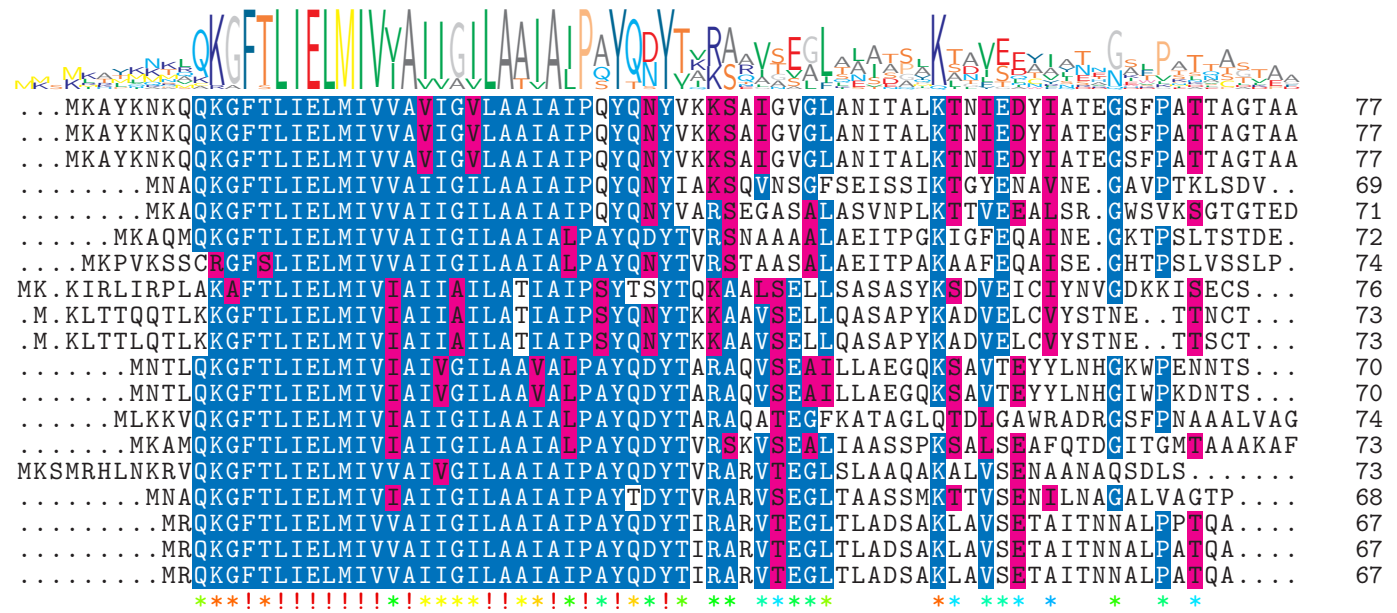


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NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1
NTDB id 1160 ABDM36 RS02330 WP 000649326.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
NTDB id 1192 PAKAF RS23870 WP 024947914.1
NTDB id 1197 PSJM300 03940 AFN76866.1
NTDB id 1204 PSJM300 03935 AFN76865.1
NTDB id 1383 A4U84 RS08530 WP 005711781.1
NTDB id 1365 NTHI RS01955 WP 011271973.1
NTDB id 1354 HI 0299 AAC21963.1
NTDB id 1119 NGFG RS11465 WP 017147169.1
NTDB id 1109 OK783 RS10670 WP 157147416.1
NTDB id 1191 AAC28468.1 484..945( )
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 183637 AQE97 RS09330 WP 016356941.1
NTDB id 1258 GC085 RS09745 WP 014844212.1
NTDB id 1274 LPP RS09530 WP 011214168.1
consensus

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NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1
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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 183637 AQE97 RS09330 WP 016356941.1
NTDB id 1258 GC085 RS09745 WP 014844212.1
NTDB id 1274 LPP RS09530 WP 011214168.1
consensus

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NTDB id 1184	GTF74	RS02335	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
NTDB id 1167	A1552VC	RS11065	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGCTAGATIN.....	153
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NTDB id 1192	PAKAF	RS23870	WP 024947914.1	AADGLWKCTSD.....QDEQFIPKGC	150
NTDB id 1197	PSJM300	03940	AFN76866.1	ADGEWSCAST.....LDAKYKPGKCS	139
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NTDB id 1383	A4U84	RS08530	WP 005711781.1	KFANNNTTWTATSCQGA...DISL.....FPANFCGQKATP	154
NTDB id 1365	NTHI	RS01955	WP 011271973.1	GNAATGVTWTTTCKGT...DASL.....FPANFCGSVTQ	149
NTDB id 1354	HI	0299	AAC21963.1	GNAAGVTWTTTCKGT...DASL.....FPANFCGSVTK	149
NTDB id 1119	NGFG	RS11465	WP 017147169.1	RRENGSVKWFVCGQPVTRADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK	166
NTDB id 1109	OK783	RS10670	WP 157147416.1	KRQDGSVKWFCGQPVTRTGDNDDTVA..DANNAIDTKHLPSTCRDESSAT	164
NTDB id 1191	AAC28468.1	484..945()		A.....ASGQISTWTCAPGAANG.....IETRRRLPTSCQ*	153
NTDB id 1188	NELON	RS01755	WP 003772458.1	K.AVP.AATSTGATDWACASAAKETADKRFGTGSGVAVGTLPAKYAPSECK	188
NTDB id 1424	RS	RS02795	WP 011000517.1	GTALPVSSAPSGPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAE	168
NTDB id 1002	ACIAD	RS15100	WP 004923779.1	YASGAVTWKCTTT.....S.DKKYVPSECRG	147
NTDB id 183637	A9E97	RS09330	WP 016356941.1	QANGDVTWCTTGG.....TLLAKYRPASCRP	136
NTDB id 1258	GCO85	RS09745	WP 014844212.1	QANGDVTWCTTGG.....TLLAKYRPASCRP	136
NTDB id 1274	LPP	RS09530	WP 011214168.1	QANGDVTWCTTGG.....TLLAKYRPASCRP	136
consensus				* * ! * * *.....* * * * *	

- ⓧ non conserved
- ✗ similar
- ✕ ≥ 50% conserved