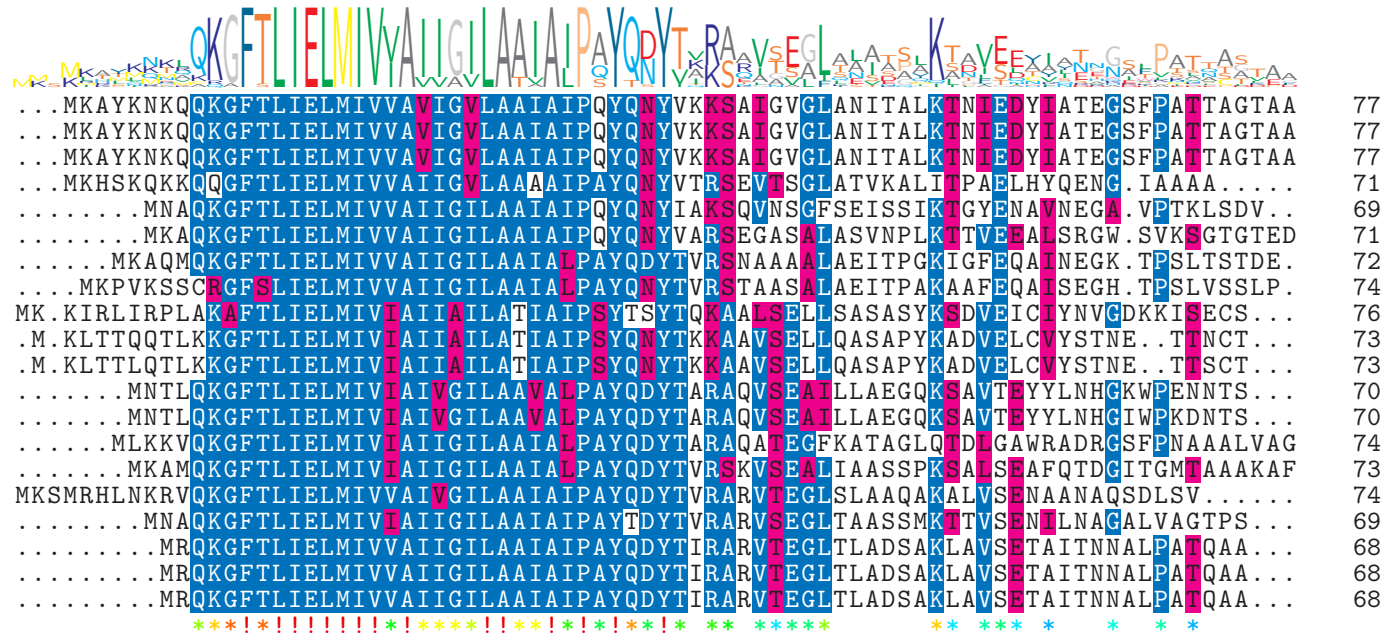


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consensus

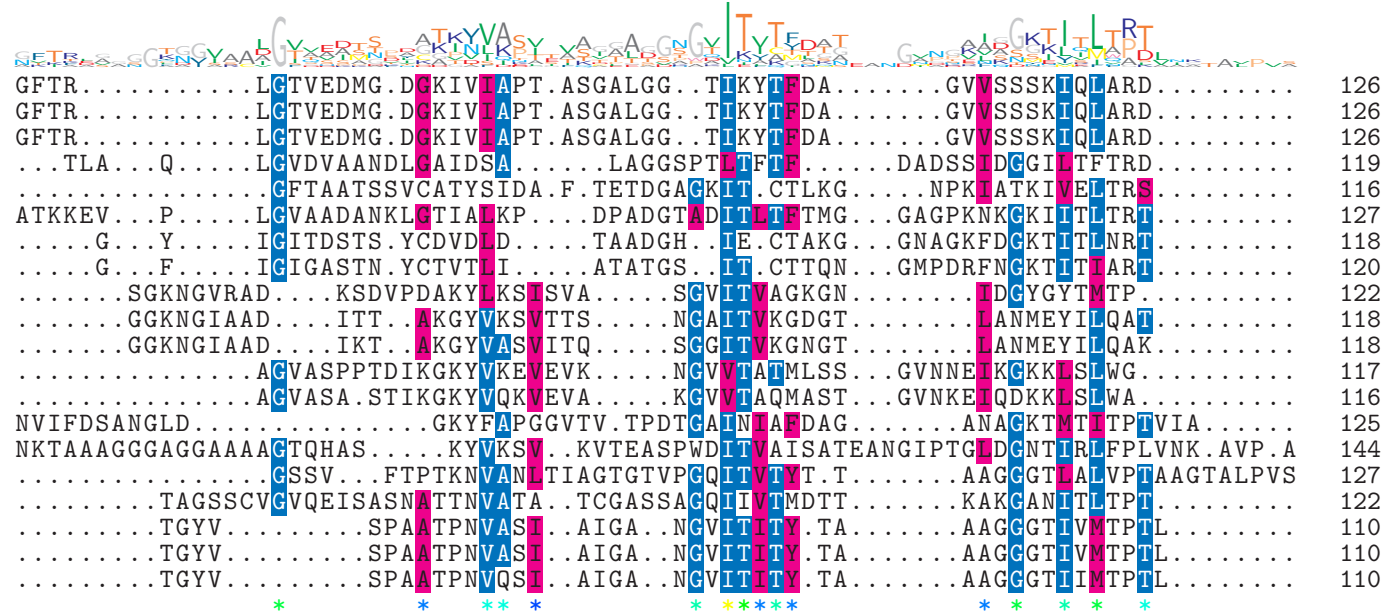
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consensus

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logo

NTDB id 1184	GTF74	RS02335	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGC	TAGATIN.....	153
NTDB id 1167	A1552VC	RS11065	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGC	TAGATIN.....	153
NTDB id 1160	ABDM36	RS02330	WP 000649326.1	ANGLWTCSTT.....VTSEIAPKGC	TAGATIN.....	153
NTDB id 1292	VP	RS12235	WP 005479693.1	AATGWSCLPT.....APAIA.VDGCS		139
NTDB id 1038	HON27	RS01510	WP 000993725.1	TDGAWTCVSD.....VPTETLPKGC	TAGTPTEGAI	149
NTDB id 1192	PAKAF	RS23870	WP 024947914.1	AADGLWKCTSD.....QDEQFIPKGC	SK.....	150
NTDB id 1197	PSJM300	03940	AFN76866.1	ADGEWSCAST.....LDAKYKPGKCS		139
NTDB id 1204	PSJM300	03935	AFN76865.1	TDGLWDCTSD.....LDSRYKPGKCL		141
NTDB id 1383	A4U84	RS08530	WP 005711781.1	KFANNITWATSCQGA...DISL.....FPANFCGQKATP		154
NTDB id 1365	NTHI	RS01955	WP 011271973.1	GNAATGVTWTTTCKGT...DASL.....FPANFCGSVTQ		149
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NTDB id 1188	NELON	RS01755	WP 003772458.1	ATSTCAIDWACASAAKETADKRFGTGSGVAVGTLPAKYAPSECK		188
NTDB id 1424	RS	RS02795	WP 011000517.1	SAPSGPIMWTCYAQNKAQAASSVAPS...GTMSLAAYAPAECR		168
NTDB id 1002	ACIAD	RS15100	WP 004923779.1	.YASGAVTWKCTTT.....S.DKKYVPSECRG		147
NTDB id 1122509	EL201	RS09825	WP 011214168.1	.QANGDVTWTCCTGG.....TLLAKYRPASCRP		136
NTDB id 1274	LPP	RS09530	WP 011214168.1	.QANGDVTWTCCTGG.....TLLAKYRPASCRP		136
NTDB id 1258	GCO85	RS09745	WP 014844212.1	.QANGDVTWTCCTGG.....TLLAKYRPASCRP		136
consensus				* * ! * * *	* * * * * *	

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