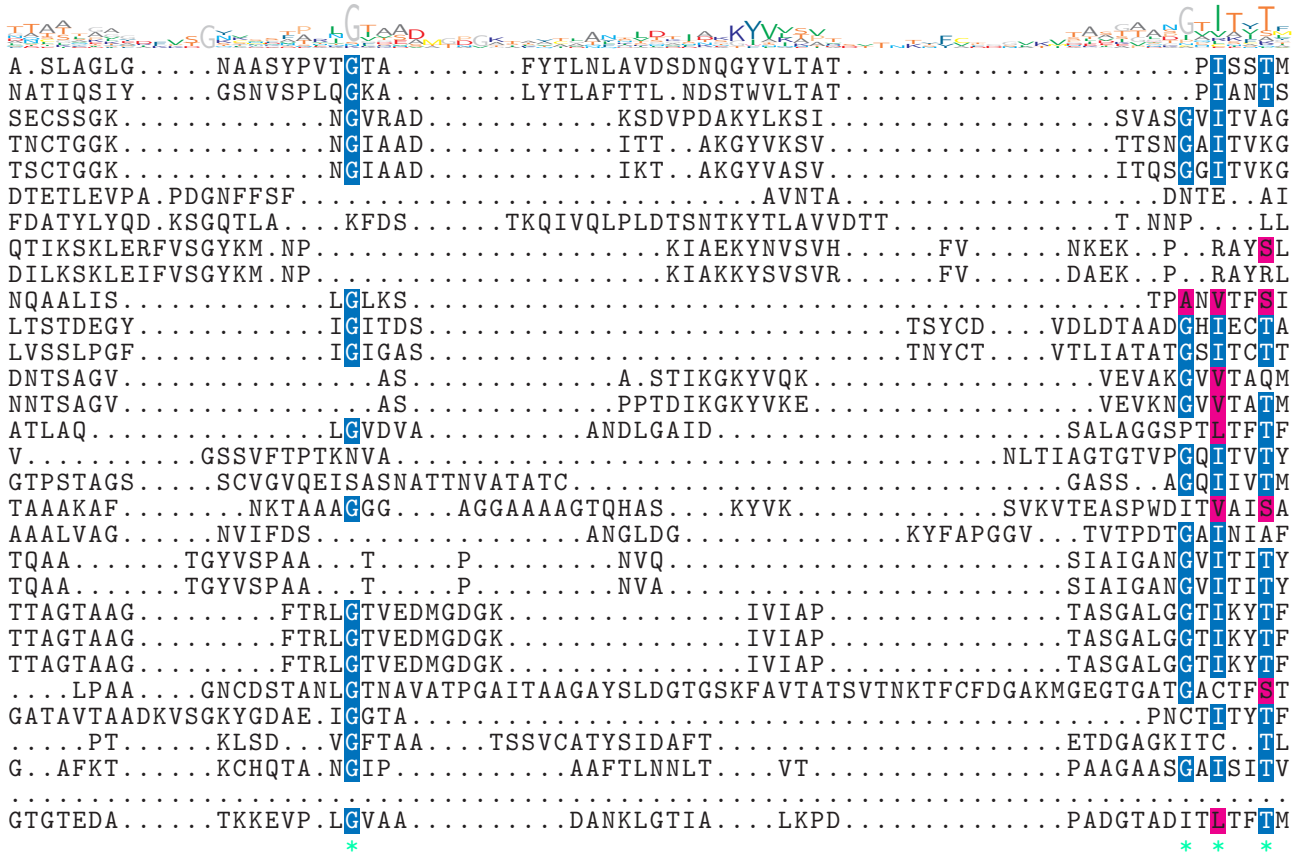


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NTDB id 1383 A4U84 RS08530 WP 005711781.1	MKKIRLIRPLAKAFTLIELMIVIAIIGILATIAIPSYTSYTQ	KAALSELLSASASYKSDVEICIYNVGDKKI	72
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NTDB id 1140 NGFG RS02440 WP 025455870.1	MEQKGFTLIELMIVVAIIGILSVIAIPSYQSYIEKGYQS	SQLYTEMVGINNISKQFILKNPLDDN	64
NTDB id 1033 TT RS11945 WP 011173287.1	MRNAKGFTLIELLVITAIIGILAAVLIPNLLAARKRAND	TVVTAYLNDVAVKFQEMYQIDNNSYTS	65
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQMKGFTLIELMIVVAIIGILAAIAPAYQDYTVRSNAAA	ALAEITPGKIGFEQAIN	66
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSSCRGFSLIELMIVVAIIGILAAIAPAYQNYTVR	STAASALAEITPAKAAFEQAI	68
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTLQKGFTLIELMIVIAIIGILAAVALPAYQDYTARAQV	SEAILLAEGQKSAVTEYYLNHGIWPK	66
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTLQKGFTLIELMIVIAIIGILAAVALPAYQDYTARAQV	SEAILLAEGQKSAVTEYYLNHGWPE	66
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NTDB id 1188 NELON RS01755 WP 003772458.1	MKAMQKGFTLIELMIVIAIIGILAAIAPAYQDYTVRSKV	SEALIAASSPKSALSEAFQTDGITGM	66
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NTDB id 1258 GC085 RS09745 WP 014844212.1	MRQKGFTLIELMIVVAIIGILAAIAPAYQDYTIRARVTEGL	TLADSAKLAVSETAITNNALPA	64
NTDB id 1274 LPP RS09530 WP 011214168.1	MRQKGFTLIELMIVVAIIGILAAIAPAYQDYTIRARVTEGL	TLADSAKLAVSETAITNNALPA	64
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NTDB id 1167 A1552VC RS11065 WP 000649326.1	MKAYKNKQKGFTLIELMIVVAIIGVLAALAAIPQYQNYVVK	SAIGVGLANITALKTNIEDYIATEGSFPA	70
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NTDB id 1079 FDQ49 RS11140 WP 022575996.1	MNAQKGFTLIELMIVVAIIGILAAIAPAYQNYIAKSQASE	AFITLADGLKTTINTNLQAGTCFAG	65
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNAQKGFTLIELMIVVAIIGILAAIAPAYQNYIAKSQVNS	GFSEISSIKTGYENAVNEGAV...	62
NTDB id 1400 DSB67 RS12660 WP 010643261.1	MKTNKQKKQKGFTLIELMIVVAIIGILAAIAPAYQDYTKKAT	MAEFPKIASSTKLAVELCAH.EHAADA	69
NTDB id 1065787 ACH105 RS15855 WP 272890074.1	MKSIAKQKGFTLIELMIVVAIIGILAAIAPQFNEYRAKAND	STAQADAKNILTVMAAAQR.....	61
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKAQKGFTLIELMIVVAIIGILAAIAPQYQNYVARSEGAS	ALASVNPLKTTVEEALSRGWSVKS	65
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 QAGDGI.....IC.....LNDQGG...KFWAKGATDCALSA.....SS 137
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved