

logo

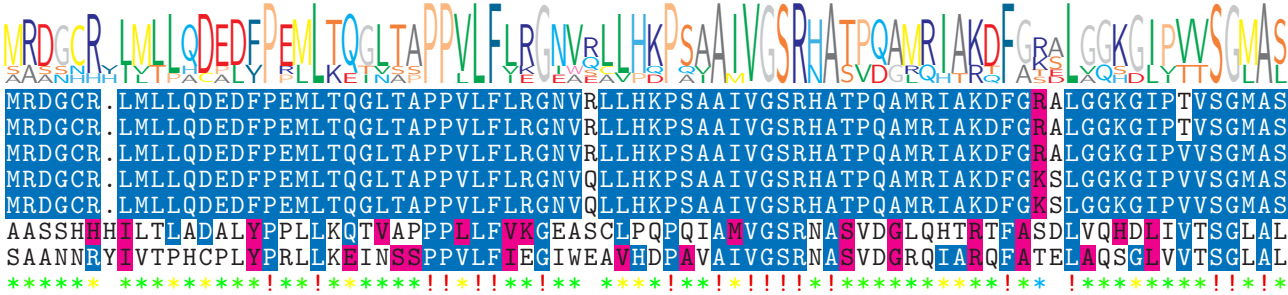
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NTDB id 1120 NGFG RS10355 WP 020997408.1
NTDB id 190923 A6L26 RS04295 WP 002216225.1
NTDB id 1094 KZH42 RS01320 WP 002224767.1
NTDB id 1089 NMB RS00600 WP 002224767.1
NTDB id 1399 DSB67 RS15600 WP 010645755.1
NTDB id 1152 A1552VC RS00200 WP 000654772.1
consensus



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NTDB id 1122 OK783 RS09595 WP 003705341.1
NTDB id 1120 NGFG RS10355 WP 020997408.1
NTDB id 190923 A6L26 RS04295 WP 002216225.1
NTDB id 1094 KZH42 RS01320 WP 002224767.1
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NTDB id 1399 DSB67 RS15600 WP 010645755.1
NTDB id 1152 A1552VC RS00200 WP 000654772.1
consensus



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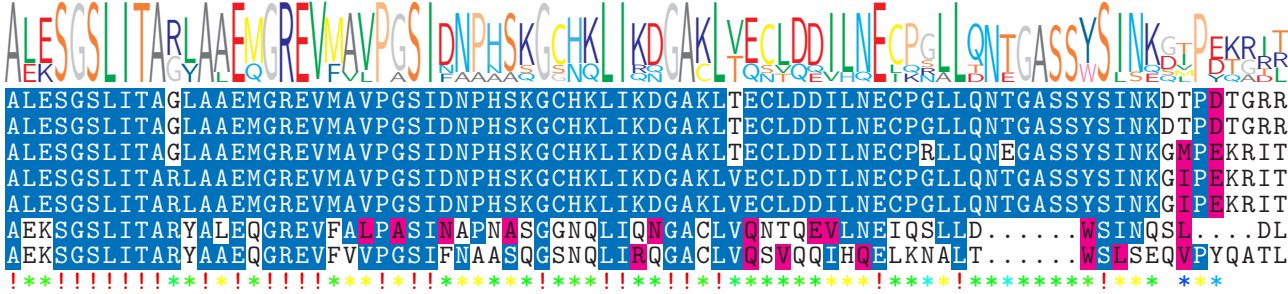
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NTDB id 1152 A1552VC RS00200 WP 000654772.1
consensus



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NTDB id 1122 OK783 RS09595 WP 003705341.1
NTDB id 1120 NGFG RS10355 WP 020997408.1
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NTDB id 1152 A1552VC RS00200 WP 000654772.1
consensus



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NTDB id 1122 OK783 RS09595 WP 003705341.1	AVQTASDQLSLPEGKMPSEKTENRPVGGSLDRMGFDPVHPDVL	AGQLAMPAADLYAALLELELDGSVAAMPGGRYQRIR	394
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NTDB id 190923 A6L26 RS04295 WP 002216225.1	AVQTASDQLSLPEGKMPSEKTENRPVGGSLDRMGFDPVHPDVL	AGQLAMPAADLYAALLELELDGSVAAMPGGRYQRIR	396
NTDB id 1094 KZH42 RS01320 WP 002224767.1	AVQTASDQLSLPEGKMPSEKTENRPVGGSLDRMGFDPVHPDVL	AGQLAMPAADLYAALLELELDGSVAAMPGGRYQRIR	396
NTDB id 1089 NMB RS00600 WP 002224767.1	AVQTASDQLSLPEGKMPSEKTENRPVGGSLDRMGFDPVHPDVL	AGQLAMPAADLYAALLELELDGSVAAMPGGRYQRIR	396
NTDB id 1399 DSB67 RS15600 WP 010645755.1	FSA.....P.....IDEEELPFPQLLANVGN	EATPVDILASRTNIPVQEVMMQLLELELSGHVVAVSGGYIRKGR	368
NTDB id 1152 A1552VC RS00200 WP 000654772.1	FSA.....V.....QSDDEELPFPELLANVGI	EATPIDILASRTQIPVQDILMMQLLELELLGHVVAVPGGYIRKGR	370
consensus	*****	* **** * *	

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NTDB id 1122 OK783 RS09595 WP 003705341.1	T	395
NTDB id 1120 NGFG RS10355 WP 020997408.1	T	395
NTDB id 190923 A6L26 RS04295 WP 002216225.1	T	397
NTDB id 1094 KZH42 RS01320 WP 002224767.1	T	397
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NTDB id 1399 DSB67 RS15600 WP 010645755.1	G	369
NTDB id 1152 A1552VC RS00200 WP 000654772.1	G	371
consensus	*	

- non conserved
- similar
- ≥ 50% conserved