

logo

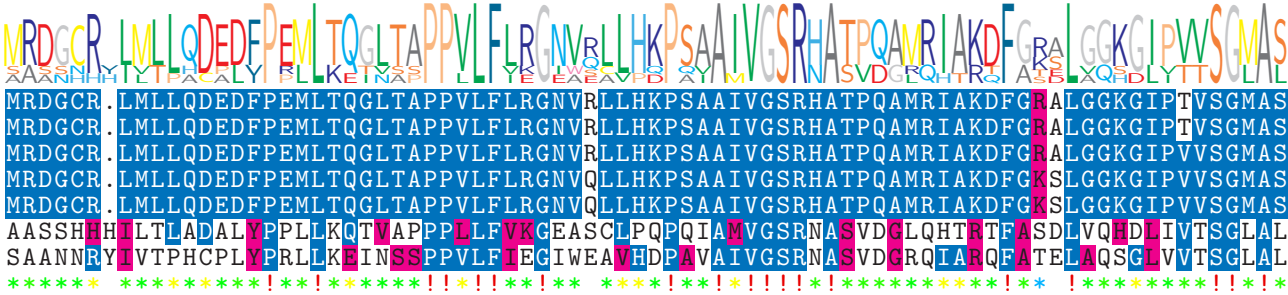
NTDB id 1122 OK783 RS09595 WP 003705341.1
NTDB id 1120 NGFG RS10355 WP 020997408.1
NTDB id 301648 DRA75 RS09855 WP 002216225.1
NTDB id 1094 KZH42 RS01320 WP 002224767.1
NTDB id 1089 NMB RS00600 WP 002224767.1
NTDB id 1399 DSB67 RS15600 WP 010645755.1
NTDB id 1152 A1552VC RS00200 WP 000654772.1
consensus



77
77
77
77
77
73
71

logo

NTDB id 1122 OK783 RS09595 WP 003705341.1
NTDB id 1120 NGFG RS10355 WP 020997408.1
NTDB id 301648 DRA75 RS09855 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



156
156
156
156
156
153
151

logo

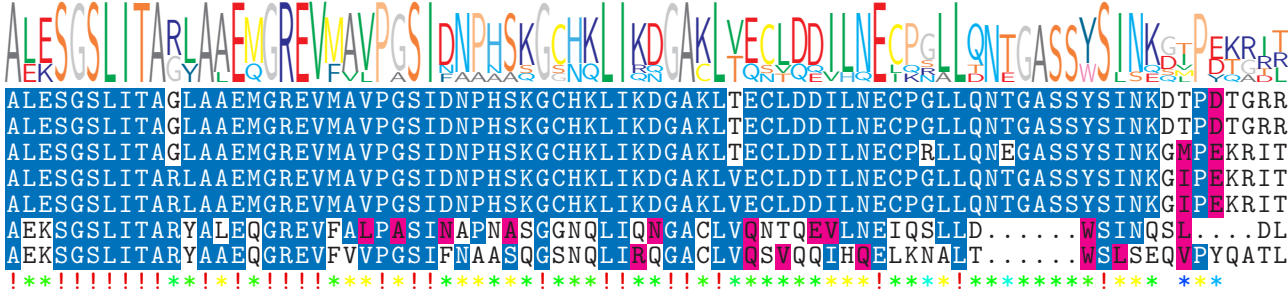
NTDB id 1122 OK783 RS09595 WP 003705341.1
NTDB id 1120 NGFG RS10355 WP 020997408.1
NTDB id 301648 DRA75 RS09855 WP 002216225.1
NTDB id 1094 KZH42 RS01320 WP 002224767.1
NTDB id 1089 NMB RS00600 WP 002224767.1
NTDB id 1399 DSB67 RS15600 WP 010645755.1
NTDB id 1152 A1552VC RS00200 WP 000654772.1
consensus



236
236
236
236
236
233
231

logo

NTDB id 1122 OK783 RS09595 WP 003705341.1
NTDB id 1120 NGFG RS10355 WP 020997408.1
NTDB id 301648 DRA75 RS09855 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



316
316
316
316
316
303
305

logo

NTDB id 1122 OK783 RS09595 WP 003705341.1	AVQTASDQLSLPEGKMPSEKTENRPVGGSLDRMGFDPVHPDVL	AGQLAMPAADLYAALLELELDGSVAAMPGGRYQRIR	394
NTDB id 1120 NGFG RS10355 WP 020997408.1	TVQTAY..APPPAAKMPSEGAAGGTAVGG	ILDKMGFDPVHPDVL	394
NTDB id 301648 DRA75 RS09855 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	*****	* **** * *	

logo

NTDB id 1122 OK783 RS09595 WP 003705341.1	T	395
NTDB id 1120 NGFG RS10355 WP 020997408.1	T	395
NTDB id 301648 DRA75 RS09855 WP 002216225.1	T	397
NTDB id 1094 KZH42 RS01320 WP 002224767.1	T	397
NTDB id 1089 NMB RS00600 WP 002224767.1	T	397
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