

logo

NTDB id 1323 RAOC RS07335 WP 004916218.1
NTDB id 1106873 ACLKM6 RS00755 WP 075167834.1
NTDB id 1271 LPP RS03220 WP 011213198.1
NTDB id 1248 GC085 RS03150 WP 011213198.1
NTDB id 1397 DSB67 RS15690 WP 005535577.1
NTDB id 1153 A1552VC RS00115 WP 000521788.1
NTDB id 1387 A4U84 RS06900 WP 020457569.1
NTDB id 1349 HI 1117 AAC22771.1
consensus



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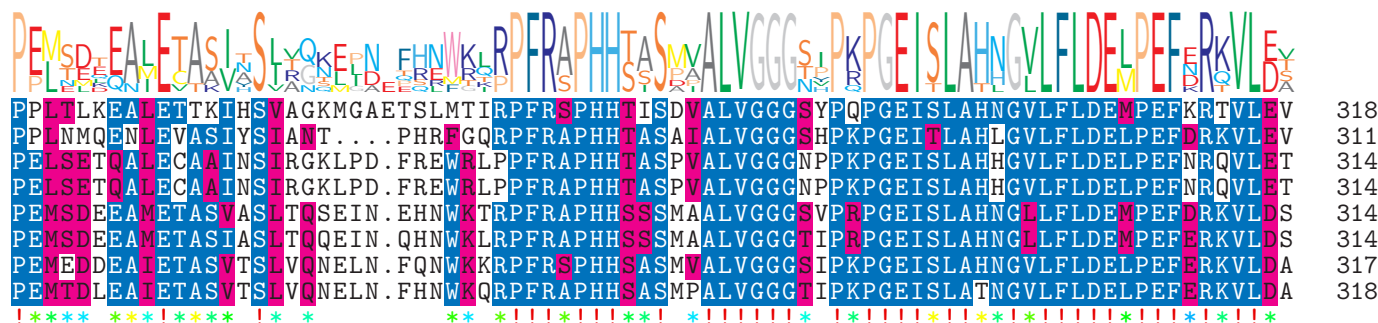
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NTDB id 1323 RAOC RS07335 WP 004916218.1	MRQPLEDREVTISRAKFTVNYPASFMLVASMNPSPSGFFPDDPNNT.SSSFEMQRYLNKLSGPLLDRIDITHIEVQKVVEFD	397
NTDB id 1106873 ACLKM6 RS00755 WP 075167834.1	LRQPLESKEIVISRASRQITFPASFQLIAAMNPPCGYAFNQDIRCQCSAESIKRYQNRISGPLLDRIDLHIDVPPLQAH	391
NTDB id 1271 LPP RS03220 WP 011213198.1	LREPLESGHICISRAAAQIEFPAKFQLIAAMNPPCGQWGNSSQANCMCTPDRIISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESGHICISRAAAQIEFPAKFQLIAAMNPPCGQWGNSSQANCMCTPDRIISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESGEIIISRAQGKTRFPAQFQIVGALNPSPITGYIEGNQAR..TNPQAILRYLGRLSGPLLDRFDMSEIIPALPKG	392
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESGEIVISRAQGKTRFPAQFQIVGALNPSPITGYIEGNQAR..ANPQSILRYLSRLSGPLLDRFDMSEIIPALPKG	392
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESGEIISIRANAKVQFPACFQLIAAMNPSPITGHYQGTNR..TSPQQLMRYLNRLSGPFLDRFDLSIEVPLLPQG	395
NTDB id 1349 HI 1117 AAC22771.1	LRQPLESGEIISIRANAKIQQFPAQFQLVAAAMNPSPITGHYGTNR..TSPQQIMRYLNRLSGPFLDRFDLSIEVPLLPQG	396
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NTDB id 1323 RAOC RS07335 WP 004916218.1	QLSDKR..KGESSEITRQVRLKAREIQQERYQDLAISYNAQMGPKEIEHFCELDEVSLLLIKNAMEKLNLSARAYDRILK	475
NTDB id 1106873 ACLKM6 RS00755 WP 075167834.1	ELHDKT..AVENS DTVRQRVIQAYERQMQRQGM...LNMNLSPKQLEQFAALATPAQNMMDLAQQRLNLSARAYHRVLR	465
NTDB id 1271 LPP RS03220 WP 011213198.1	ELIKPNTHLEKQSLAIREKVTMKMHEIQMARQDS...LNANLNSKTCCEMVCELGSEEQLFLREVMSKLLKSARGYHRLK	470
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPNTHLEKQSLAIREKVTMKMHEIQMARQDS...LNANLNSKTCCEMVCELGSEEQLFLREVMSKLLKSARGYHRLK	470
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLSEG.GDRGEPTAIVKERVNQARDCLARSGK...VNALLGSREIEAFQPLQKSDAEFLENALHRLGLSIRAYHRIK	467
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLANG.GDRGESTAAVRQRVVVARERMLARSGK...VNALLQSREIEQYCPLLKADAEFLESALHRLGLSIRAYHRIK	467
NTDB id 1387 A4U84 RS06900 WP 020457569.1	ALQNNTDQRGETTEQARKRVLEARHLQFNRRGK...INAQLTTKEIEQDCKLTDRDALFLENALTKLGLSVRAYHRILK	471
NTDB id 1349 HI 1117 AAC22771.1	SLQNT.GDRGETSAQVREKVLKVREIQMERAGK...INAYLNSKEIERDCKLNDKDAFFLEKALNKLGLSVRAYHRILK	471
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NTDB id 1323 RAOC RS07335 WP 004916218.1	VSRTIADLERETNIQSHHIAEAIQYRSIDRDFWKV.....	510
NTDB id 1106873 ACLKM6 RS00755 WP 075167834.1	VARSIAADLAESSEIQTALHSEALS YRAQSSP.....	496
NTDB id 1271 LPP RS03220 WP 011213198.1	VSRTIADYNSSKKVLLNHLQQALS YKQNLHLPK.....	503
NTDB id 1248 GC085 RS03150 WP 011213198.1	VSRTIADYNSSKKVLLNHLQQALS YKQNLHLPK.....	503
NTDB id 1397 DSB67 RS15690 WP 005535577.1	VARTIADLEGSPQIERTHLAEALGYRAMDRLLKQLTAQAV	507
NTDB id 1153 A1552VC RS00115 WP 000521788.1	VARTIADLQGEAQIARPHLAEALGYRAMDRLLKQLSAQNV	507
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQPHLAEALGYRAMDRLLQRLQGGS.	510
NTDB id 1349 HI 1117 AAC22771.1	VSRTIADLQGEQQIFQPHLAEALGYRAMVRLQLKLSNM..	509
consensus	!***** ** !***** !*** ** *	

- X non conserved
- X similar
- X ≥ 50% conserved