

logo

NTDB id 1323 RAOC RS07335 WP 004916218.1	LREPLESGEIIISRAAKYFPAFQLYAAMNPSPGHHYGGONRTPQQLRYLRLSGPFLDRFDMSIEYPALPQG	397
NTDB id 1271 LPP RS03220 WP 011213198.1	MRQPLEDRVITISRAKFTVNYPASFMLVASMNPSPSGFFPDDPNNT.SSSFEMQRYLNKLSGPLLDRIDITHIEVQKVVEFD	394
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESGHICISRAAAQIEFPAKFQLIAAMNPPCGQWGNSSQANCMCTPDRIISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1073356 ACI51Y RS20215 WP 405078589.1	LREPLESGHICISRAAAQIEFPAKFQLIAAMNPPCGQWGNSSQANCMCTPDRIISRYLAKLSAPLLDRIDMQVTIHALSQE	393
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESGEIIISRTAKVCYPARVQLVAAMNPSPSGHHYQGVHNR..LPAQQILRYLSKLSGPFLLDRFDLSIEVPLLPQG	392
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESGEIVISRAQKTRFPAFQLVGALNPSPGTGYEGSSQAR..ANPQSILRYLSRLSGPLLDRFDMSIEIPALPKG	392
NTDB id 1349 HI 1117 AAC22771.1	LREPLESGEIIISRAQKTRFPAFQIVGALNPSPGTGYEGNQAR..TNPQAAILRYLGRLSGPLLDRFDMSLEIPALPKG	396
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESGEIIISRANAKIQFPAFQLVAAAMNPSPGTGHYTGTNHR..TSPQQIMRYLNRLSGPFLLDRFDLSIEVPLLPQG	395
consensus	*!*****!!!!* **** *!!!!*****!!!!!!* * ** *	

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NTDB id 1323 RAOC RS07335 WP 004916218.1	QLSD..KRKGESSEIRQVRLKAREIQQERYQDLAISYNAQMGPKEIEHFCELDVSLLLTKNAMEKLNLSARAYDRILK	475
NTDB id 1271 LPP RS03220 WP 011213198.1	ELIKPNTHLEKQSLAIQREKVTMKHEIQMARQD...SLNANLNSKTCCEMVCELGSEEQFLREVMSKIKLSARGYHRLIK	470
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPNTHLEKQSLAIQREKVTMKHEIQMARQD...SLNANLNSKTCCEMVCELGSEEQFLREVMSKIKLSARGYHRLIK	470
NTDB id 1073356 ACI51Y RS20215 WP 405078589.1	VLSQQ.HYQGESSATIRERVLIARQIQLKRAN...KINALLTSREIEKHCGLEMA DATYLEEVMNKLGLSVRAWHRLIK	468
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLANG.GDRGESTAAVRQRVVVARERMLARSG...KVNALLQSREIEQYCPLLKADAEFLESALHRLGLSIRAYHRIK	467
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLSEG.GDRGEPTAIIVKERVNQARDCMLARSG...KVNALLQSREIEAFCLPQKSDAEFLENALHRLGLSIRAYHRIK	467
NTDB id 1349 HI 1117 AAC22771.1	SLQNT.GDRGETSAQVREKVLKVREIQMERAG...KINAYLNSKEIERDCKLNDKDAFFLEKALNKLGLSVRAYHRILK	471
NTDB id 1387 A4U84 RS06900 WP 020457569.1	ALQNNTDQRGETTEQARKRVLEARHLQFNRRG...KINAQLTTKEIEQDCKLTDRDALFLENALT KLGLSVRAYHRILK	471
consensus	! ***** *!!!! ***** ! **!! * ***** ! ! ! ** *** ** *!!!!!!*!!!!!!*	

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NTDB id 1323 RAOC RS07335 WP 004916218.1	VSRTIADLERETNIQSHHIAEAIQYRSLDRDFWKV.....	510
NTDB id 1271 LPP RS03220 WP 011213198.1	VSRTIADMNSSKKVLLNHLQQALSYPKQNLHLPK.....	503
NTDB id 1248 GC085 RS03150 WP 011213198.1	VSRTIADMNSSKKVLLNHLQQALSYPKQNLHLPK.....	503
NTDB id 1073356 ACI51Y RS20215 WP 405078589.1	VARTIADLGERDNIERKHLAEALSYRCMDRLLIQTHKSLE	508
NTDB id 1153 A1552VC RS00115 WP 000521788.1	VARTIADLQGEAQIARPHLAEALGYRAMDRLLKQLSAQNV	507
NTDB id 1397 DSB67 RS15690 WP 005535577.1	VARTIADLEGSPQIERTHLAEALGYRAMDRLLKQLTAQAV	507
NTDB id 1349 HI 1117 AAC22771.1	VSRTIADLQGEQQIFQPHLAEALGYRAMVRLKQLSNM..	509
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQPHLAEALGYRAMDRLLQRLQGES.	510
consensus	!***** ** !****! * !* ***** *	

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