

logo

NTDB id 1323 RAOC RS07335 WP 004916218.1	LREPLESGEIIISRAAAQIEFPKFLIAAMNPPCGQWGN	397
NTDB id 1271 LPP RS03220 WP 011213198.1	LREPLESGHICISRAAAQIEFPKFLIAAMNPPCGQWGN	394
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESGHICISRAAAQIEFPKFLIAAMNPPCGQWGN	394
NTDB id 1056087 ABZP12 RS01645 WP 008574838.1	LREPLESGVVTIARAARSVDFPARFQLVAAMNPPCGWAGDGS	394
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESGEIVISRAAQGKTRFPAFQLVGALNPSTGYEGS	392
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESGEIIISRAAQGKTRFPAFQIVGALNPSTGYEGN	392
NTDB id 1349 HI 1117 AAC22771.1	LRQPLESGEIIISRAANAKIQFPAFQLVAAAMNPSTGHYTGT	396
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESGEIIISRAANAKVQFPACFQLIAAMNPSTGHYQGT	395
consensus	*!!!!!!!!*****!!!!*!*!!!!!!!!!!!!!!!!!!!!*****	

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NTDB id 1323 RAOC RS07335 WP 004916218.1	QLSD..KRKGESSEIRQRVLKAREIQQERYQDLAISYNAQMGPKEIEHF	475
NTDB id 1271 LPP RS03220 WP 011213198.1	ELIKPNTHLEKQSLAIREKVTMKHEIQMARQD...SLNANLNSKTC	470
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPNTHLEKQSLAIREKVTMKHEIQMARQD...SLNANLNSKTC	470
NTDB id 1056087 ABZP12 RS01645 WP 008574838.1	ALRS..GNLGEDSASVRARVVAARQRLARGA...LPNAQLDQADT	468
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLANG.GDRGESTAAVRQRVWVARERMLARSG...KVNALLQSREIEQY	467
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLSEG.GDRGEPTAIVKERVNQARDCMLARSG...KVNALLQSREIEAF	467
NTDB id 1349 HI 1117 AAC22771.1	SLQNT.GDRGETSAQVREKVLKVREIQMERAG...KINAYLNSKEIERD	471
NTDB id 1387 A4U84 RS06900 WP 020457569.1	ALQNNTDQRGETTEQARKRVLEARHLQFNRRG...KINAQLTTKEIEQD	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 1056087 ABZP12 RS01645 WP 008574838.1	VARTIADLDDSPDIATPHLTEATIGYRKLDRAIGTASAA..	506
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	VSRTIADLQGEQKIFQPHLAEALGYRAMVRLLKQLSNM..	509
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQPPHLAEALGYRAMDRLLQRLQGES.	510
consensus	!!!!!!!!*!*!!!!*!!!!*!!!!*!!!!*	

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