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NTDB id 1323 RAOC RS07335 WP 004916218.1	LREPLESGEIIISRAAKYFPAFQLYAAMNPSPTCGYGSGQNRTPQQLRYLRRLSGPFLDRFDMSIEIPALPQG	397
NTDB id 1271 LPP RS03220 WP 011213198.1	MRQPLEDRVITISRAKFTVNPASFMLVASMNPSPSGFFPDDP.NNTSSSFEMQRYLNLKLSGPLLDRIDITHIEVQKVVEFD	394
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESGHICISRAAAQIEFPAKFQLIAAMNPPCGQWGNSSQANCMCTPDRIISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1079304 ACKKH6 RS19930 WP 004915428.1	LREPLESRQVIISRAKAKVCFPANFQLIAALNPSPTGQYQGDY..CRSSSTKILRYLSRVSGPFLDRFDLSIEIPLPLPG	393
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESGEIVISRAQKTRFPAKFQLVGALNPSPTGYYEGSQ..ARANPQSILRYLSRLSGPLLDRFDMSIEIPALPKG	392
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESGEIIISRAQKTRFPAKFQIVGALNPSPTGYYEGNQ..ARTNPQAAILRYLGRLSGPLLDRFDMSLEIPALPKG	392
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESGEIIISRANAKVQFPACFQLIAAMNPSPTGHYQGTH..NRTSPQQLMRYLNLRLSGPFLDRFDLSIEVPLLPQG	395
NTDB id 1349 HI 1117 AAC22771.1	LRQPLESGEIIISRANAKIQFPAKFQLYAAMNPSPTGHYTGTH..NRTSPQQIMRYLNLRLSGPFLDRFDLSIEVPLLPQG	396
consensus	*!!!!!!!!!!!!!!!!**!*!!!!*****!!!!!!* ** **** *!!!! *!!!!*!!!!!!*!!!!***** ** *	

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NTDB id 1323 RAOC RS07335 WP 004916218.1	TSPGDRGEISSEIQRVLRKAREIQMARCKNALNSKEIECDALFLEALKGLSARAYHRLK	475
NTDB id 1271 LPP RS03220 WP 011213198.1	QLSD..KRKGESSEIRQRVLKAREIQQERYQDLAISYNAQMGPKEIEHFCELDEVSLLLIKNAMEKLNLSARAYDRILK	470
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPNTHLEKQSLAIREKVTMKHEIQMARQDS...LNANLNSKTCCEMVCELGSEEQLFLREVMSKCLKLSARGYHRLK	470
NTDB id 1079304 ACKKH6 RS19930 WP 004915428.1	ELIKPNTHLEKQSLAIREKVTMKHEIQMARQDS...LNANLNSKTCCEMVCELGSEEQLFLREVMSKCLKLSARGYHRLK	468
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLSHQ.QYQSENSEIQRARVILAREIQMKRMGK...LNSQLTARETTNLCQLTSKDSLFLEHALNKLGLSTRAYHRLK	467
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLANG.GDRGESTAAVRQRVWVARERMLARSGK...VNALLQSREIEQYCPLLKADAEFLESALHRLGLSIRAYHRIK	467
NTDB id 1387 A4U84 RS06900 WP 020457569.1	TLSEG.GDRGEPTAIVKERVNQARDCMLARSGK...VNAQLTKEIEQDCCKLTDRDALFLENALTKLGLSVRAYHRIK	471
NTDB id 1349 HI 1117 AAC22771.1	ALQNNTDQRGETTEQARKRVLEARHLQFNRRGK...INAQLTKEIEQDCCKLTDRDALFLENALTKLGLSVRAYHRIK	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 1079304 ACKKH6 RS19930 WP 004915428.1	VARTIADLQNTNVEIKPHLEALGYRAMDKLLHLQKQVS	508
NTDB id 1153 A1552VC RS00115 WP 000521788.1	VARTIADLQGEAQIARPHLAEALGYRAMDRLLKQLSAQNV	507
NTDB id 1397 DSB67 RS15690 WP 005535577.1	VARTIADLEGSPQIERTHLAEALGYRAMDRLLKQLTAQAV	507
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQPHLAEALGYRAMDRLLQRLQGGS.	510
NTDB id 1349 HI 1117 AAC22771.1	VSRTIADLQGEQIFQPHLAEALGYRAMVRLKQLSNM..	509
consensus	!!!!!!!** ** !****!*!!!!***** *	

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