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NTDB id 1323 RAOC RS07335 WP 004916218.1	LREPLESGEIISRAAKIEFPAFQLYAAMNPSPGCHYGONRITPQRLRYLRLSGPFLDRFDMSEIIPALPQG	397
NTDB id 1271 LPP RS03220 WP 011213198.1	MRQPLEDRVITISRAKFTVNPASFMLVASMNPSPSGFFPDDPNNT.SSSFEMQRYLNKLSGPLLDRIDITHIEVQKVVEFD	394
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESGHICISRAAAQIEFPAKFQLIAAMNPPCGQWGNSSQANCMCTPDRIISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1078320 ACJVZ4 RS21195 WP 022649778.1	LREPLESGEIHSRTRAKISYPAQFQLVAAMNPSPGTHYQGNHN..RCTPEQTLRYLSKLSGPFLLDRFDSLIEIPLPPPG	391
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESGEIISRAAQKTRFPAQFQIVGALNPSPGTHYEGNQA..RTNPQATILRYLGRLSGPPLDRFDMSEIIPALPKG	392
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESGEIVISRAAQKTRFPAQFQLVQALNPSPGTHYEGSSQA..RANPQSILRYLSRLSGPPLDRFDMSEIIPALPKG	392
NTDB id 1349 HI 1117 AAC22771.1	LRQPLESGEIISIRANAKIQFPAQFQLVAAMNPSPGTHYTGTN..RTSPQQIMRYLNRLSGPFLDRFDSLIEVPLLPQG	396
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESGEIISIRANAKVFPACFQLIAAMNPSPGTHYQGTN..RTSPQQLMRYLNRLSGPFLDRFDSLIEVPLLPQG	395
consensus	*!*!*!*!*!* *!* * * * *!*!*!*!*!*!*!*!* * * * * * * * * * * * * * * * * *	

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NTDB id 1323 RAOC RS07335 WP 004916218.1	LSD..KRKGESSEIRQVRLKAREIQQERYQDLAISYNAQMGPKEIEHFCELDVSLLLIKNAMEKLNLSARAYDRILK	475
NTDB id 1271 LPP RS03220 WP 011213198.1	ELIKPNTHLEKQSLAIQREKVTMKHEIQMARQDS...LNANLNSKTCCEMVCELGSEEQFLREVMSKIKLSARGYHRLK	470
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPNTHLEKQSLAIQREKVTMKHEIQMARQDS...LNANLNSKTCCEMVCELGSEEQFLREVMSKIKLSARGYHRLK	470
NTDB id 1078320 ACJVZ4 RS21195 WP 022649778.1	LLRQT.GIKGESSAVVRERVIAAQAQYARQNR...LNARLDNSGTRQFCSLNAEDAGWLEETLRFGLSTRAWQRLK	466
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLSEG.GDRGEPTAIVKERVNQARDCLARSGK...VNALLGSREIEAFQPLQKSDAEFLENALHRLGLSIRAYHRIK	467
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLANG.GDRGESTAAVRQVWVARERMLARSGK...VNALLQSREIEQYCPLLKADAEFLESALHRLGLSIRAYHRIK	467
NTDB id 1349 HI 1117 AAC22771.1	SLQNT.GDRGETSAQVREKVLKVREIQMERAGK...INAYLNSKEIERDCKLNDKDAFFLEKALNKLGLSVRAYHRILK	471
NTDB id 1387 A4U84 RS06900 WP 020457569.1	ALQNNTDQRGETTEQARKRVLEARHLQFNRRGK...INAQLTKEIEQDCKLTDRDALFLENALTKLGLSVRAYHRILK	471
consensus	! *	

logo

NTDB id 1323 RAOC RS07335 WP 004916218.1	VSRTIADLERETNIQSHHIAEAIQYRSLDRDFWKV.....	510
NTDB id 1271 LPP RS03220 WP 011213198.1	VSRTIADMNSSKKVLLNHLQQALSQKQNLHLPK.....	503
NTDB id 1248 GC085 RS03150 WP 011213198.1	VSRTIADMNSSKKVLLNHLQQALSQKQNLHLPK.....	503
NTDB id 1078320 ACJVZ4 RS21195 WP 022649778.1	VARTVADVEGCPGTEIRRHLEALSRYAIDRLLHLQKMLA	506
NTDB id 1397 DSB67 RS15690 WP 005535577.1	VARTIADLEGSPQIERTHLAEALGYRAMDRLLKQLTAQAV	507
NTDB id 1153 A1552VC RS00115 WP 000521788.1	VARTIADLQGEAQIARPHLAEALGYRAMDRLLKQLSAQNV	507
NTDB id 1349 HI 1117 AAC22771.1	VSRTIADLQGEQIFQPHLAEALGYRAMVRLKQLSNM..	509
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQPHLAEALGYRAMDRLLQRLQGES.	510
consensus	! *	

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