

logo

NTDB id 1271 LPP RS03220 WP 011213198.1
NTDB id 1248 GC085 RS03150 WP 011213198.1
NTDB id 1153 A1552VC RS00115 WP 000521788.1
NTDB id 1397 DSB67 RS15690 WP 005535577.1
NTDB id 1349 HI 1117 AAC22771.1
NTDB id 1387 A4U84 RS06900 WP 020457569.1
NTDB id 264115 C1062 RS11905 WP 261981412.1
NTDB id 1323 RAOC RS07335 WP 004916218.1
consensus



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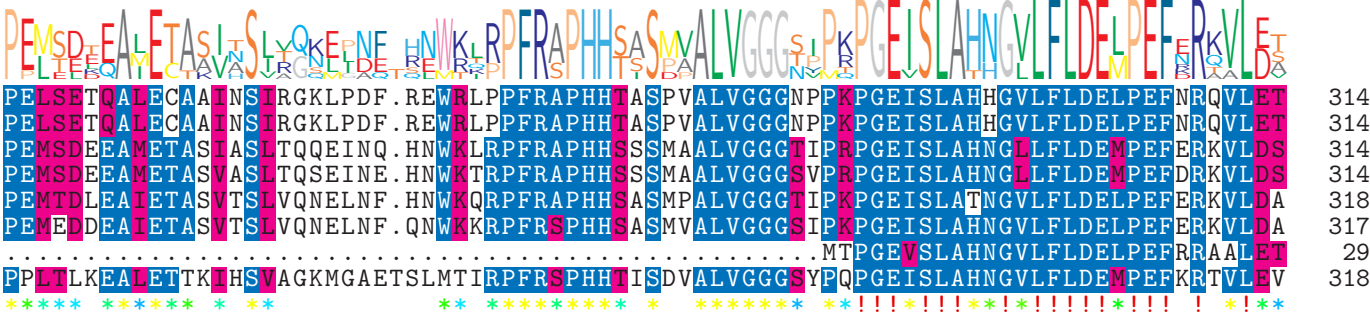
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NTDB id 1271 LPP RS03220 WP 011213198.1	LREPLESGHICISRAAAQIEFPKQFLIAAMNPPCGQWGNISQANCMCTPDRISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESGHICISRAAAQIEFPKQFLIAAMNPPCGQWGNISQANCMCTPDRISRYLAKLSAPLLDRIDMQVTIHALSQE	394
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESGEIVISRAQGGKTRFPARFQLVGALNPSTGYEYEGSQAR..ANPQSILRYLSRLSGPLDRFDMSTIEIPALPKG	392
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESGEIIISRAQGGKTRFPARFQIVGALNPSTGYEYEGNQAR..TNPQAAILRYLGRLSGPLDRFDMSTIEIPALPKG	392
NTDB id 1349 HI 1117 AAC22771.1	LRQPLESGEIIISRAAAQIEFPKQFLIAAMNPPCGQWGNISQANCMCTPDRISRYLAKLSAPLLDRIDMQVTIHALSQE	396
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESGEIIISRAAAQIEFPKQFLIAAMNPPCGQWGNISQANCMCTPDRISRYLAKLSAPLLDRIDMQVTIHALSQE	395
NTDB id 264115 C1062 RS11905 WP 261981412.1	LRQPLETEGQVVISRASGTMTPFCRFMLAAAMNPPCGYLGDRRRRACTCPPAQIARYRRKISGPLLDRFDLLMEVPAVDPS	109
NTDB id 1323 RAOC RS07335 WP 004916218.1	MRQPLEDREVTISRAKFTVNYPASFMLVASMNPSPSGFFPDPPNN.TSSSFEMQRYLNKLSGPLLDRIDIHIEVQKVEFD	397
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NTDB id 1271 LPP RS03220 WP 011213198.1	ELIKPNTHELEKQSLAIREKVTMKHEIQMARQD...SLNANLNSKTCCEMVCELGSEEQLFLREVMSKLLKLSARGYHRLK	470
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPNTHELEKQSLAIREKVTMKHEIQMARQD...SLNANLNSKTCCEMVCELGSEEQLFLREVMSKLLKLSARGYHRLK	470
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLANG.GDRGESTAAVRQVRVVARERMLARSG...KVNALLQSREIEQYCPLLKADAFLSALHRLGLSIRAYHRIK	467
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLSEG.GDRGETAIIVKERNVQARDCLARSG...KVNALLGSREIEAFCLPLQKSDAEFLNALHRLGLSIRAYHRIK	467
NTDB id 1349 HI 1117 AAC22771.1	SLQNT.GDRGETSAQVREKVLKVREIQMERAG...KINAYLNSKEIERDCKLNDKDAFFLEKALNKLGLSVRAYHRILK	471
NTDB id 1387 A4U84 RS06900 WP 020457569.1	ALQNNTDQRGETTEQARKRVLEARHLQFNRRG...KINAQLTTKEIEQDCKLTDRDALFLENALTKLGLSVRAYHRILK	471
NTDB id 264115 C1062 RS11905 WP 261981412.1	ILAS..SPAGECSASTIRERVVAARRLQSSRYAGTPFRNNAALSGKALQKYCRLLPEGRATILLRAVEELALSARAYDRILK	187
NTDB id 1323 RAOC RS07335 WP 004916218.1	QLSD..KRKGESSEIIRQRVLKAREIQQERYQDLAISYNAQMGPKEIEHFCELDVSVLLLTKNAMEKLNLSARAYDRILK	475
consensus	! * * * * * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * *	

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NTDB id 1271 LPP RS03220 WP 011213198.1	VSRTIADLNSSKKVLLNHLQQAISYKQNLHLPK.....	503
NTDB id 1248 GC085 RS03150 WP 011213198.1	VSRTIADLNSSKKVLLNHLQQAISYKQNLHLPK.....	503
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	VSRTIADLQGEQIFQPHLAEALGYRAMVRLKQLSNM..	509
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQPHLAEALGYRAMDRLLQRLQGES.	510
NTDB id 264115 C1062 RS11905 WP 261981412.1	VARTIADLEGTSDIQDVHLYEAVQYRAFEQSLRD.....	221
NTDB id 1323 RAOC RS07335 WP 004916218.1	VSRTIADLERETNIQSHHTAEAIQYRSLDRDFWKV.....	510
consensus	! * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * *	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved