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NTDB id 1115 OK783 RS00485 WP 003698035.1	MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIV.NPTGFVTVSSPARIALDFEQTGISMD..Q	77
NTDB id 1092 CAD91899.1 1..2286( )	MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIV.NPTGFVTVSSPARIALDFEQTGISMD..Q	77
NTDB id 1100 NMB RS09455 WP 002225652.1	MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIV.NPTGFVTVSSPARIALDFEQTGISMD..Q	77
NTDB id 1186 GTF74 RS01320 WP 000788426.1	.....	0
NTDB id 1175 A1552VC RS12105 WP 000788426.1	.....	0
NTDB id 1408 DSB67 RS13980 WP 010643500.1	.....	0
NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKIIAILIL..ISASLSMAYSKDNSLISVKVLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDIFISTDMQIPSDQ	78
NTDB id 1257 GC085 RS04855 WP 163621000.1	MRKIIAILIL..ISASLSLAYS KDNSLISVKVLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDIFISTDMQIPSDQ	78
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARILDFDKAGQKLA..K	77
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLK..Q	77
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLK..Q	77
NTDB id 1093781 ACMGGG RS22505 WP 414155836.1	.....	0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNSGLSRLGIALLA.MFAPALLAADLEKLDVAAIPGDRVELKLFQFDEPVA.APRGYTIEQPARIALDLPGVQNKLG..T	76
consensus	* * * * * * * * * * * * * * *	

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NTDB id 1115 OK783 RS00485 WP 003698035.1	QVLEYADPLLSKISAAQNSSRARLVNLNKPQQYNTVEVRGNKVVIFINESDDTVSAPARPAVKAAAPAAPAKQQA.....	151
NTDB id 1092 CAD91899.1 1..2286( )	QVLEYADPLLSKISAAQNSSRARLVNLNKPQQYNTVEVRGNKVVIFINESDDTVSAPARPAVKAAAPAAPAKQQAAPSTK	157
NTDB id 1100 NMB RS09455 WP 002225652.1	QVLEYADPLLSKISAAQNSSRARLVNLNKPQQYNTVEVRGNKVVIFINESDDTVSAPARPAVKAAAPAAPAKQQAAPSTK	157
NTDB id 1186 GTF74 RS01320 WP 000788426.1	.....MKNGLKT..YVAQWTWLTWVGLALC.....A.....	24
NTDB id 1175 A1552VC RS12105 WP 000788426.1	.....MKNGLKT..YVAQWTWLTWVGLALC.....A.....	24
NTDB id 1408 DSB67 RS13980 WP 010643500.1	.....MKMRHGLTTPRKLTLGFLSLWFSLLLA.....P.....	28
NTDB id 1284 ACE17W RS12370 WP 010946666.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSLVINGKSNDFENS.....	136
NTDB id 1257 GC085 RS04855 WP 163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSLVINGKSNDFENS.....	136
NTDB id 1009 ACIAD RS15175 WP 004923730.1	SLIPVTTNEASSIDVSSDDKRARVVVNLKDAGAFTRVEGNVFIKINNIQSTAT.....PVAT.....	136
NTDB id 1052 ABD1 RS15850 WP 001017033.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDAGAFTRVEGNVFIKINSAQTSNK.....PLPV.....	136
NTDB id 1039 HON27 RS01460 WP 031993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDAGAFTRVEGNVFIKINSAQTSNK.....PLPV.....	136
NTDB id 1093781 ACMGGG RS22505 WP 414155836.1	.....	0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	KNRELSVGNTRSVTVVEAKDRTRLINLITLSSYTTRVEGNLNFVVVGNPAGASVASTAPVKASPAPAS.....	146
consensus	* * * * * * * * * * * * * *	

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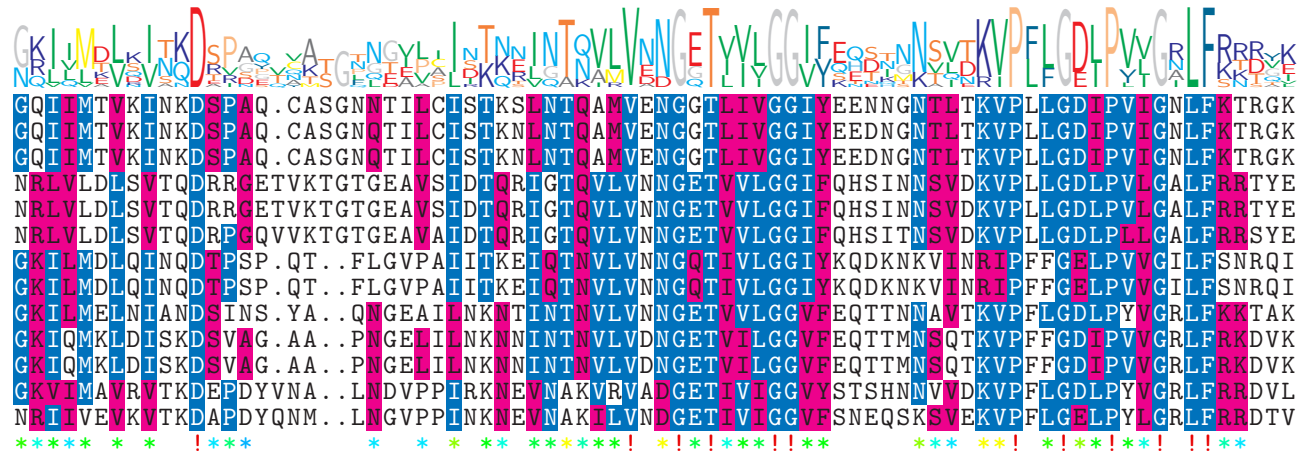
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NTDB id 1092 CAD91899.1 1..2286( )	SAVSVSEPFPAKQQAAPFTESVSVSAPFSPAKQQAASAQKQTA.....APAKQQAAPAKQTNIDFRKDGKNA	229
NTDB id 1100 NMB RS09455 WP 002225652.1	SAVSVSEPFPAKQQAAPFTESVSVSAPFSPAKQQAASAQKQQAAPAKQQAAPAKQTNIDFRKDGKNA	237
NTDB id 1186 GTF74 RS01320 WP 000788426.1	.....SSMV.....FSAESATANQLENIDFRVNKEKA	51
NTDB id 1175 A1552VC RS12105 WP 000788426.1	.....SSMV.....FSAESATANQLENIDFRVNKEKA	51
NTDB id 1408 DSB67 RS13980 WP 010643500.1	.....F.....VAEEVQSNLSKSIDFRNTKDKD	52
NTDB id 1284 ACE17W RS12370 WP 010946666.1	.....KEVFI.....TNQVVNAKHEIKRIDFRGIEKQG	164
NTDB id 1257 GC085 RS04855 WP 163621000.1	.....KEVFI.....TNQVVNAKHEIKRIDFRGIEKQG	164
NTDB id 1009 ACIAD RS15175 WP 004923730.1	.....SA.....PAPQQIGNIGFQRGAQGE	157
NTDB id 1052 ABD1 RS15850 WP 001017033.1	.....VS.....AQPQVSNIGFQRGSQGE	156
NTDB id 1039 HON27 RS01460 WP 031993779.1	.....VS.....AQPQVSNIGFQRGSQGE	156
NTDB id 1093781 ACMGGG RS22505 WP 414155836.1	.....	0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	.....Y.....AQPIK.....PKPYVPAGRAIRNIDFRGEKGE	175
consensus	* * * * * * * * * * * * * *	





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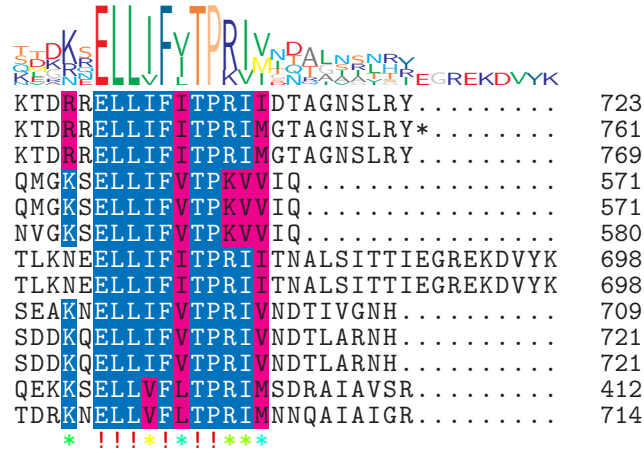
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 NTDB id 1193 PAKAF RS26665 WP 003161249.1  
 consensus



non conserved  
 similar  
 ≥ 50% conserved