

logo

	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLFQFDEPVAAPRGYTI EQPARIALDLPGVQNKLGTKNRE	
NTDB id 93009 SEEP7378 RS01675 WP 000832822.1	0
NTDB id 1433 BW25113 RS17570 WP 000815987.1	0
NTDB id 1347 HI 0435 AAC22094.1	0
NTDB id 1373 NTHI RS02645 WP 011272046.1	0
NTDB id 1193 PA5040 NP 253727.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLFQFDEPVAAPRGYTI EQPARIALDLPGVQNKLGTKNRE	80
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
consensus	0

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	LSVGNTRSVTWVEXDKRIRITNLRASLSLTTTRVEGRNFFSVCNAAAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	
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NTDB id 1433 BW25113 RS17570 WP 000815987.1	0
NTDB id 1347 HI 0435 AAC22094.1	0
NTDB id 1373 NTHI RS02645 WP 011272046.1	0
NTDB id 1193 PA5040 NP 253727.1	LSVGNTRSVTVVEAKDRTRLIINLTALSSYTTRVEGNNLFVVVGNPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	160
NTDB id 1186 GTF74 RS01320 WP 000788426.1 MKNGLKT..YVAQW.....LTLWVGLALCAS.....S.....MVFSAESATA	36
NTDB id 1175 A1552VC RS12105 WP 000788426.1 MKNGLKT..YVAQW.....LTLWVGLALCAS.....S.....MVFSAESATA	36
NTDB id 1408 DSB67 RS13980 WP 010643500.1 MKMRHGLTTLTPRKLTLGL.....FSLWFSLLAP.....FVAEEVQS	37
consensus	0

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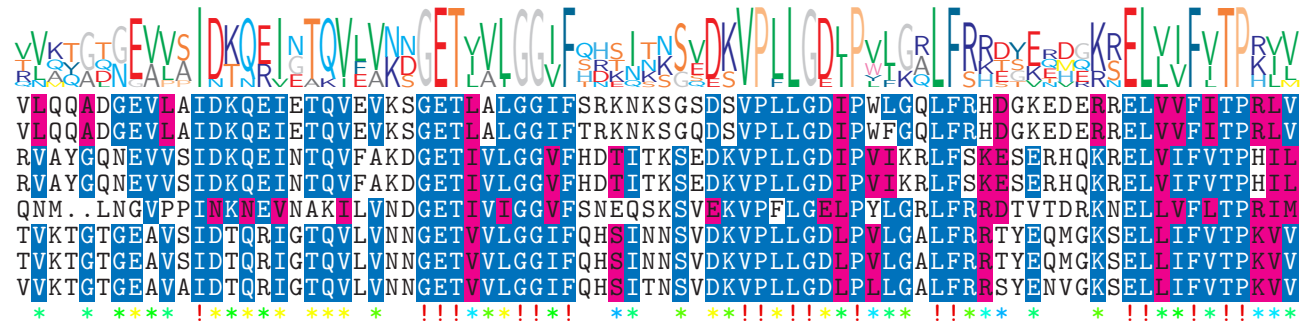
	NQLENIDFRVNKEKAAVLWELASPSAVVDVQKVEGLSIELLKTVDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	
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NTDB id 1347 HI 0435 AAC22094.1	0
NTDB id 1373 NTHI RS02645 WP 011272046.1	0
NTDB id 1193 PA5040 NP 253727.1	RAIRNIDFQRGEKGEVVIDLSDPTLSPDIQEGGKIRLDFAKTQLPDLRVRDLVKDFATPVQFVNASASQSDRTSITI	240
NTDB id 1186 GTF74 RS01320 WP 000788426.1	NQLENIDFRVNKEKAAVLWELASPSAVVDVQKVEGLSIELLKTVDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKAAVLWELASPSAVVDVQKVEGLSIELLKTVDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1408 DSB67 RS13980 WP 010643500.1	NSLKSIDFRNTKDKDAVIVVELVSPAIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTPVESVEVFRDSTTRLVA	117
consensus	0

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	TVDGEYQHDYTLKGRYLEVLEKLPKADKPKSVLAKKCKLISINFDIPVQVQLLADQNDNINWSDVSGNLT	
NTDB id 93009 SEEP7378 RS01675 WP 000832822.1 MKRWIAIILIVLMP.A.....AQAGKAAKVTLVVDVVPVQVLQTLAEQERQNLVVSQDVSQGLS	59
NTDB id 1433 BW25113 RS17570 WP 000815987.1 MKQWIAALLMLIP.G.....VQAAPQKVTLMVDVVPVQVLQALAEQEKLNLVSPDVSQGLS	59
NTDB id 1347 HI 0435 AAC22094.1 MKKYFLKCGYFLVCFCLPLIVFA.N.....PKTDNERFFIRLSQAPLAQTLQALAFQDQVNLVIGDILENKIS	67
NTDB id 1373 NTHI RS02645 WP 011272046.1 MKKYFLKCGYFLVCFCLPLIVFA.N.....PKTDNECFIRLSQAPLAQTLQALAFQDQVNLVIGERLEGNIS	67
NTDB id 1193 PA5040 NP 253727.1	EPSPGLYDYLVTQDNRITVS.IKPMTTEDAERRKKDNFAYTGEKLSINFDIPVQVQLLADQNDNINWSDVSGNLT	319
NTDB id 1186 GTF74 RS01320 WP 000788426.1	TVDGEFQHDYTLKGRYLEVLEKLPKADKPKSVLAKKCKLISINFDIPVQVQLLADQNDNINWSDVSGNLT	194
NTDB id 1175 A1552VC RS12105 WP 000788426.1	TVDGEFQHDYTLKGRYLEVLEKLPKADKPKSVLAKKCKLISINFDIPVQVQLLADQNDNINWSDVSGNLT	194
NTDB id 1408 DSB67 RS13980 WP 010643500.1	YIDEDYSHDYRLTGKYLEVLEKLPKADKPKSVLAKKCKLISINFDIPVQVQLLADQNDNINWSDVSGNLT	195
consensus	0

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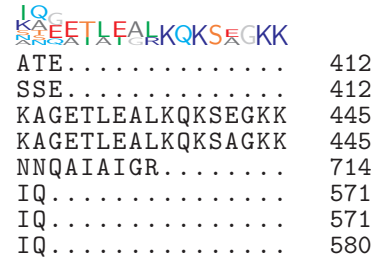
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 consensus



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 NTDB id 1175 A1552VC RS12105 WP 000788426.1
 NTDB id 1408 DSB67 RS13980 WP 010643500.1
 consensus



non conserved
 similar
 ≥ 50% conserved