

logo

NTDB id 1193 PA5040 NP 253727.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTIEQPARIALDLPGVQNKLGTKNRE	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
NTDB id 48024 H7R55 RS02185 WP 227673968.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1465 ASU2 RS06170 WP 014991934.1		0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
consensus		

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NTDB id 1193 PA5040 NP 253727.1	LSVGNTRSVTVVEAKDRTRLIINLTALSSYTTRVEGNNLFVVVGNSPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	160
NTDB id 1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQTW.....LTLWVGLALCAS.....S....MVFSAESATA	36
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTW.....LTLWVGLALCAS.....S....MVFSAESATA	36
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTGL.....FSLWFSLLLAP.....FVAEEVQS	37
NTDB id 48024 H7R55 RS02185 WP 227673968.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1465 ASU2 RS06170 WP 014991934.1		0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
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NTDB id 1193 PA5040 NP 253727.1	RAIRNIDFQRGEKGEGNVVIDLSDPTLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASDAQSDRTSITI	240
NTDB id 1186 GTF74 RS01320 WP 000788426.1	NQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1408 DSB67 RS13980 WP 010643500.1	NSLKSIDFRTNKDKDAVIVVELVSPA AIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDSTTTRLVA	117
NTDB id 48024 H7R55 RS02185 WP 227673968.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1465 ASU2 RS06170 WP 014991934.1		0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
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NTDB id 1193 PA5040 NP 253727.1	EPGSLYDYL VYQTDNRLT VSIKPTTTEDAERRKKDNFAYTGEKLSLNFDIDIVRSVLQLIADFTDLNLVASDVTQGNITL	320
NTDB id 1186 GTF74 RS01320 WP 000788426.1	TVDGEFQHDYTLKGGKYLEVVISKLK.ADEKPKPKSVLEKEGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	195
NTDB id 1175 A1552VC RS12105 WP 000788426.1	TVDGEFQHDYTLKGGKYLEVVISKLK.ADEKPKPKSVLEKEGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	195
NTDB id 1408 DSB67 RS13980 WP 010643500.1	YIDEDYSHDYRLTGRYLEVKVGKLG.ANEKAPDKSILDKGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	196
NTDB id 48024 H7R55 RS02185 WP 227673968.1	MSLLTLLILA.....PLSPA KNTVLVVDV PVAQVLQSLADLEQKNLVIAPDVQGSLSL	54
NTDB id 1433 BW25113 RS17570 WP 000815987.1	MKQ.....WIAALLMLIP.....GVQA AKPQKVTLMVDV PVAQVLQALAEQEKLNLVSPDVS GTVSL	60
NTDB id 1465 ASU2 RS06170 WP 014991934.1	MKRM.....ILLIFF.....ILPPVIAHAFSISLKNAPTAKILSYLAEEHGKNIVLSDNIETNTTL	56
NTDB id 1347 HI 0435 AAC22094.1	MKKYFLKCGYFLVCFCPLPLIVF.....ANPKTDNERFFIRLSQAPLAQTLEQLAFQ QDVNLVIGDILENKISL	68
NTDB id 1373 NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCPLPLIVF.....ANPKTDNECFFIRLSQAPLAQTLEQLAFQ QDVNLV MGERLEGNISL	68
consensus	*** * * * **** * ** * * * * * ! * * ! * * ! * ! * * * * * !	

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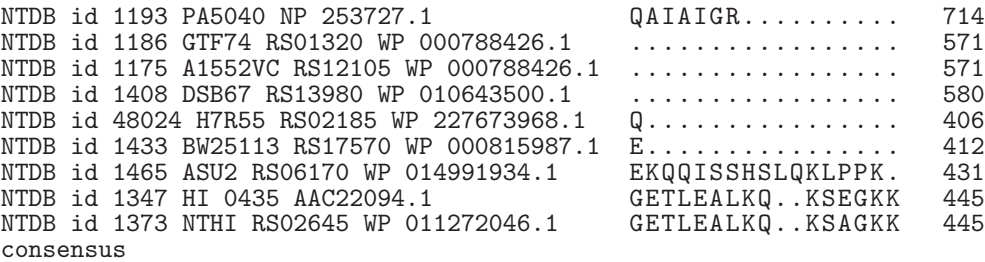
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logo



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non conserved

similar

≥ 50% conserved