

logo

NTDB id 58368 NGK29 RS20825 WP 022649623.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTI EQPARIALDLPGVQNKLGTKNRE	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	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
consensus		

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NTDB id 58368 NGK29 RS20825 WP 022649623.1	LSVGNTRSVTVWEAKDRTRLIINLTALSSYTTRVEGNNLFVVVGNPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	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	LSVGNTRSVTVVEAKDRTRLIINLTALSSYTTRVEGNNLFVVVGNPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	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
consensus		

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NTDB id 58368 NGK29 RS20825 WP 022649623.1	NQLENIDFRVNKEKAAYLVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLV	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	RAIRNIDFQRGEKGEGNVVIDLSDPTLSPDIQEGGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASAQSDRTSITI	240
NTDB id 1186 GTF74 RS01320 WP 000788426.1	NQLENIDFRVNKEKAAYLVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLV	116
NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKAAYLVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLV	116
NTDB id 1408 DSB67 RS13980 WP 010643500.1	NSLKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSTTRLVA	117
consensus		

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NTDB id 58368 NGK29 RS20825 WP 022649623.1	MK...TWIGLLAALSQP.....LWAATPKPVTLVLDVVPVQVLQAMVAQEDRNLVVSPDVSGSL	59
NTDB id 1433 BW25113 RS17570 WP 000815987.1	MK...QWIAALLIMLIPG.....VQAAPQKVITLMVDVVPVQVLQALAEQEKLNLVSPDVSGT	59
NTDB id 1347 HI 0435 AAC22094.1	MKKYFLKCGYFLVCFCLPLIVFA.N.....PKTDNERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKIS	67
NTDB id 1373 NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVFA.N.....PKTDNECFIRLSQAPLAQTLEQLAFQQDVNLVIGERLEGNIS	67
NTDB id 1193 PA5040 NP 253727.1	EPSGLYDYLVTQDNRLTVS..TKPMTTEDAERRKKDNFAYTGEKLSLINFQDIPVRNVLQLIADFTDLNLVASDTVQGNIT	319
NTDB id 1186 GTF74 RS01320 WP 000788426.1	TVDGEFQHDYTLKGKYLEVV..ISKLK.ADEKPKPKSVLEKEGKLISINFQDIPVRNVLQLIADYNGFNLVVS	194
NTDB id 1175 A1552VC RS12105 WP 000788426.1	TVDGEFQHDYTLKGKYLEVV..ISKLK.ADEKPKPKSVLEKEGKLISINFQDIPVRNVLQLIADYNGFNLVVS	194
NTDB id 1408 DSB67 RS13980 WP 010643500.1	YIDEDYSHDYRLTGRYLEVK..VGKLIK.ANEKAPDKSILDKEGKLISINFQDIPVRNVLQLIADYNGFNLVVS	195
consensus	MK...TWIGLLAALSQP.....LWAATPKPVTLVLDVVPVQVLQAMVAQEDRNLVVSPDVSGSL	

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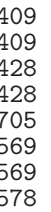
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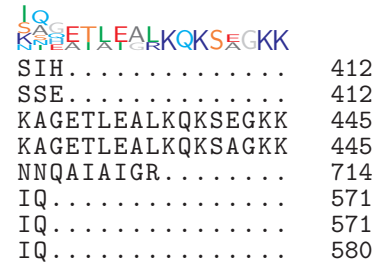
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-  non conserved
-  similar
-  $\geq 50\%$ conserved