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NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKIIAIIIL..ISASISLMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1257 GC085 RS04855 WP 163621000.1	MRKIIAIIIL..ISASISLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 99299 QMG95 RS17265 WP 200766247.1		0
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 1193 PAKAF RS26665 WP 003161249.1	MNSGLSRLGIALLAA.MFAPALLAADLEKLDVAALPGDRVELKLQFDEPVA.APRGYTIEQPARIALDLPGVQNKLG..	76
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARLLIDFDKAGQKLAK..	77
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQGGLKQ..	77
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQGGLKQ..	77
consensus	* * * * *	

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NTDB id 1284 ACE17W RS12370 WP 010946666.1	STIITYTNEASVDVSDDDRMRLNINLKDAGSYTTRVEGNLFILYLNQSSNLKKEKALGSTRHYNTLAVKRSKQTELEKLSAFESTALITTYFKANSARTNDKEENSAPVKASPLPYOSVFTSAV	145
NTDB id 1257 GC085 RS04855 WP 163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGTAGSIY.SLVLNGKSNDLFENS.....KEVFITNQV	145
NTDB id 99299 QMG95 RS17265 WP 200766247.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGTAGSVY.SLVLNGKSNDLFENS.....KEVFITNQV	26
NTDB id 1186 GTF74 RS01320 WP 000788426.1	MAEFC...RAH...IYRCALLIFIGL.....FSTTGV	32
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGL...KT...YVAQTWLTWVGLAL.....C.....ASSMVFSAE	32
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKNGL...KT...YVAQTWLTWVGLAL.....C.....ASSMVFSAE	33
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MKMRHGL...TLTPRKLTGLFSLWFSLLL.....A.....P...FVAEE	156
NTDB id 1009 ACIAD RS15175 WP 004923730.1	KNRELISVGNTRSVTVVEAKDRTRLINLTTLSSTYTRVEGNLFFVVGNSPAGASVASTAPVKASPAPASYAQPIKPKPY	138
NTDB id 1052 ABD1 RS15850 WP 001017033.1	SLIPVTTNEASSIDVSSDDKRARVVVNLKDAGAFTRRVEGNVFILKINNIQSTAT.....PVAT..SA.....	138
NTDB id 1039 HON27 RS01460 WP 031993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDAGAFTRRVEGNTFILKINSAQTSNK.....PLPV..VS.....	138
consensus	* * * * *	

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NTDB id 1284 ACE17W RS12370 WP 010946666.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	225
NTDB id 1257 GC085 RS04855 WP 163621000.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	106
NTDB id 99299 QMG95 RS17265 WP 200766247.1	HAAQSNFVGVIDFRLSKEKHALLVVELASNAAAVDIQMPDGLNISLHETSIVVDEKLFVVVDVSDFATSVENVEVFRKESNT	112
NTDB id 1186 GTF74 RS01320 WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
NTDB id 1175 A1552VC RS12105 WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	113
NTDB id 1408 DSB67 RS13980 WP 010643500.1	EVQSNLSKSIDFRTNKKDAIVIVVELVSPA AIVDIKRVQEGLSIDLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSTT	236
NTDB id 1193 PAKAF RS26665 WP 003161249.1	VPAGRAIRNIDFQRGEKGEKNVVIDLSDPTLSPDIEQEGGKIRLDFAKTQLPDALRVRDLVDKDFATPVQFVNASDAQSDRT	218
NTDB id 1009 ACIAD RS15175 WP 004923730.1	PAPQGGIGNIGFQRGAQGEGLVVIDLQGTINTPVDVQQQGSKVVIRFPGKIPHTLARRLNTTDFATPVASIDSYNDGSGNG	217
NTDB id 1052 ABD1 RS15850 WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
NTDB id 1039 HON27 RS01460 WP 031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	
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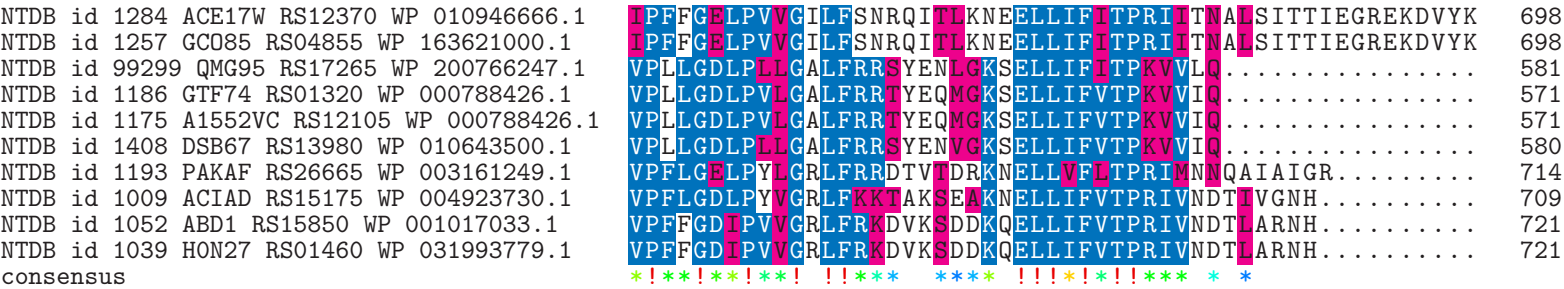
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logo



☐ non conserved
☒ similar
☒ ≥ 50% conserved