

NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKIIAILIL..ISASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1257 GCO85 RS04855 WP 163621000.1	MRKIIAILIL..ISASLSLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 408735 GPY07 RS09065 WP 000788426.1		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	MNSGLSRLGIALLA..MFAPALLAADLEKLDVAALPGDRVELKQLQFDEPVA.APRGYTIEQPARIALDLPGVQNKIG..T	76
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKQTIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKAGQKLA..K	77
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLK..Q	77
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLK..Q	77
consensus	* * * * * * * * * * * * * * * *	

[illegible]

NTDB id	1284	ACE17W	RS12370	WP	010946666.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLN	TKIPLNLMKRYDVSDFHSVPVQIITMQEGKKV	225		
NTDB id	1257	GCO85	RS04855	WP	163621000.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLN	TKIPLNLMKRYDVSDFHSVPVQIITMQEGKKV	225		
NTDB id	408735	GPY07	RS09065	WP	000788426.1	SATANQLENIDFRVNKEKA	AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFS	TPVESVEVFRKEPST	112	
NTDB id	1186	GTF74	RS01320	WP	000788426.1	SATANQLENIDFRVNKEKA	AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFS	TPVESVEVFRKEPST	112	
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	SATANQLENIDFRVNKEKA	AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFS	TPVESVEVFRKEPST	112	
NTDB id	1408	DSB67	RS13980	WP	010643500.1	EVQSNSLKSIDFRTNKD	KDAVIVVELVSPAAIVDTKR	VQEGLSIDLNTSVKDDQLYLLDVKDFS	TVVESVEVFRDTSTT	113
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	VPAGRAIRNIDFQRGEKG	EKNVVIDLSDPTLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVN	ASAQSDRT	236	
NTDB id	1009	ACIAD	RS15175	WP	004923730.1	PAPQQGIGNIGFQRGAQGEGLVVIDLQGTNT	PVDVQQQGSKVVVRFPGIKIPHTHARRLNTDFATPVASIDSYNDGSNG	218		
NTDB id	1039	HON27	RS01460	WP	031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNT	PVDVQQQGSKVVIRTIGTKIPHTHARRLNVNDFATPVSSIDAYNDKGVG	217		
NTDB id	1052	ABD1	RS15850	WP	001017033.1	.AQPGQVSNIGFQRGSQGEGLVVVDLLGSNT	PVDVQQQGSKVVIRTIGTKIPHTHARRLNVNDFATPVSSIDAYNDKGVG	217		
consensus						* * * * *	***** * ** ! ***** **** * ***** ! * * ! ***** *			

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logo



X non conserved
X similar
X ≥ 50% conserved