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NTDB id 1284	ACE17W	RS12370	WP	010946666.1	MRKIIAILII..ISASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1257	GC085	RS04855	WP	163621000.1	MRKIIAILII..ISASLSLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 75112	Vgang	RS11065	WP	211294085.1		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	MNSGLSRLGIALIAA.MFAPALLAADLEKIDVAALPGDRVELKLQFDEPVA.APRGYTIEQPARIALDLPGVQNKIGT..	76
NTDB id 1009	ACIAD	RS15175	WP	004923730.1	MKQIRQFSIGTVAYAIMQVSAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKAGQKLAK..	77
NTDB id 1052	ABD1	RS15850	WP	001017033.1	MNHVFRQFSMGAVAIATMQAASQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLKQ..	77
NTDB id 1039	HON27	RS01460	WP	031993779.1	MNHVFRQFSMGAVAIATMQAASQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLKQ..	77
consensus					* * * * *	

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[illegible]

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NTDB	id	1284	ACE17W	RS12370	WP	010946666.1	VNAKHEIKRIDFRGIEKGQGRVLIDVSDTGIPIDVTQVGKEIVVNFLNLTKIPLNLMKRYDVSDFHSPVQIITMQEGKKV	225
NTDB	id	1257	GCO85	RS04855	WP	163621000.1	VNAKHEIKRIDFRGIEKGQGRVLIDVSDTGIPIDVTQVGKEIVVNFLNLTKIPLNLMKRYDVSDFHSPVQIITMQEGKKV	225
NTDB	id	75112	Vgang	RS11065	WP	211294085.1	QAISNQFLGLDFRLNKQKEGMIVVKMASSASAVDIKKNAAGGLNIDLILQTSVTDDNLYVIDVKDFS TAVSTVEVFREENT	127
NTDB	id	1186	GTF74	RS01320	WP	000788426.1	SATANQLLENIDFRVNKEKA AVLVELASPASAVVDVQKVQEGLSIELLKTADVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB	id	1175	A1552VC	RS12105	WP	000788426.1	SATANQLLENIDFRVNKEKA AVLVELASPASAVVDVQKVQEGLSIELLKTADVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB	id	1408	DSB67	RS13980	WP	010643500.1	EVQSANLSKSIDFRTNKDKDAVIVVELVSPA AIVDTKRVRQEGLSIDLLNTSVKDDQLYLLDVKDFS TVVESVEVERDTSTT	113
NTDB	id	1193	PAKAF	RS26665	WP	003161249.1	VPAGRATRINIDFRQRGEKGE GNVVITDLSDPTLSPDIQEQQGGKIRLDFAKTQLPDALRVRLDVKDFA TPVQFVNASAQSDRT	236
NTDB	id	1009	ACIAD	RS15175	WP	004923730.1	PAPQGIGIGNIGFQRGAQGEGLVVITDLQGTNTPV DVQQGSGKVVVRFPGIKIPHTLARRLNTTDFATPVA SIDSYNDGSNG	218
NTDB	id	1052	ABD1	RS15850	WP	001017033.1	JAPQGVSNIGFQRSQGEGLVVVDLLGSNTPV DVQQGSGKVVIRTIGTKIPHTLARRLNVNDFATPV SIDA YNDKGVG	217
NTDB	id	1039	HON27	RS01460	WP	031993779.1	JAPQGVSNIGFQRSQGEGLVVVDLLGSNTPV DVQQGSGKVVIRTIGTKIPHTLARRLNVNDFATPV SIDA YNDKGVG	217
consensus								* * * * *

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NTDB id 1193 PAKAF RS26665 WP 003161249.1
NTDB id 1009 ACIAD RS15175 WP 004923730.1
NTDB id 1052 ABD1 RS15850 WP 001017033.1
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consensus

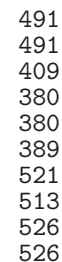
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consensus

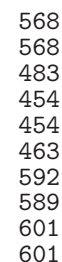
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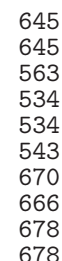
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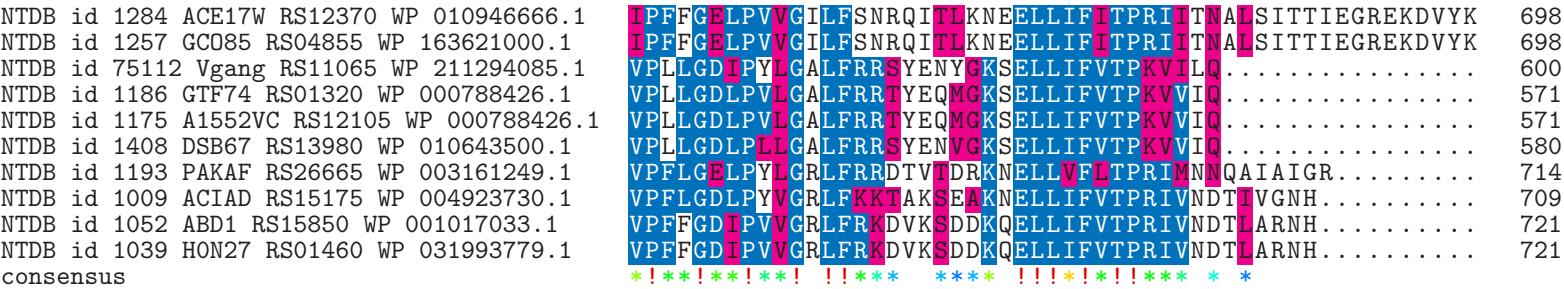
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non conserved
 similar
 ≥ 50% conserved