

logo

NTDB id 1284	ACE17W	RS12370	WP 010946666.1	MRKIIAAILII..ISASISLMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFISTDMQIPSDQ	78
NTDB id 1257	GC085	RS04855	WP 163621000.1	MRKIIAAILIL..ISASISLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFISTDMQIPSDQ	78
NTDB id 1186	GTF74	RS01320	WP 000788426.1		0
NTDB id 1175	A1552VC	RS12105	WP 000788426.1		0
NTDB id 268082	C1N51	RS08665	WP 199770605.1		0
NTDB id 1408	DSB67	RS13980	WP 010643500.1		0
NTDB id 1193	PAKAF	RS26665	WP 003161249.1	MNSGLSRLGIALLAA.MFAPALLAADLEKLDVAALPGDRVELKQLQFDEPVA.APRGYTIEQPARIALDLPGVQNKLG..T	76
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	MKQIIRQFSIGTVAYAIMQVSAQVAITNVVPMNIPGQGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKAGQKLA..K	77
NTDB id 1052	ABD1	RS15850	WP 001017033.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQGGLK..Q	77
NTDB id 1039	HON27	RS01460	WP 031993779.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQGGLK..Q	77
consensus				* * * * *	

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NTDB id 1284	ACE17W	RS12370	WP 010946666.1	STIITYTNEASVDVSDDDRRRLINLNKGLSYTRVEGNFSLVYN..SKQTLAREASS..PVPYVSMVFANQV	145
NTDB id 1257	GC085	RS04855	WP 163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSLVLNGKSNDLFENSKE.....VFITNQV	145
NTDB id 1186	GTF74	RS01320	WP 000788426.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSLVLNGKSNDLFENSKE.....VFITNQV	32
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	MKNGLKTY..VAQTWLTWVG...LALCASS.....MVFS.AE	32
NTDB id 268082	C1N51	RS08665	WP 199770605.1	MKNGLKTY..VAQTWLTWVG...LALCASS.....MVFS.AE	33
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFF...LLLA.....PFVA.AE	33
NTDB id 1193	PAKAF	RS26665	WP 003161249.1	MKMRHGLTLTPRKLTGLFSLWFS...LLLA.....PFVA.AE	156
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	KNRELISVGNTRSVTVVEAKDRTRLIINLTTLSSYTTRVEGNLNFVVGNSPAGASVASTAPVKASPAPASYAQPIKPKPY	138
NTDB id 1052	ABD1	RS15850	WP 001017033.1	SLIPVTTNEASSIDVSSDDKRARVVVNLKDGAGFTTRVEGNVFILKINNIQSTAT.....PVAT..SA.....	138
NTDB id 1039	HON27	RS01460	WP 031993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDGAGFTTRVEGNVFILKINSAQTSNK.....PLPV..VS.....	138
consensus				* * * * *	

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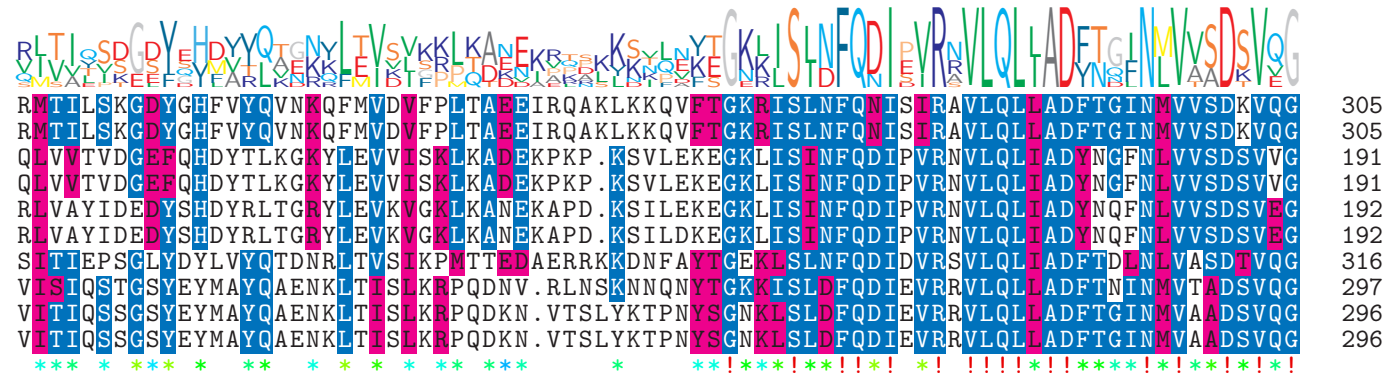
NTDB id 1284	ACE17W	RS12370	WP 010946666.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	225
NTDB id 1257	GC085	RS04855	WP 163621000.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	225
NTDB id 1186	GTF74	RS01320	WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
NTDB id 268082	C1N51	RS08665	WP 199770605.1	EVQSNLSKSIDFRTNKDKDAVIVVELASPA AIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSTT	113
NTDB id 1408	DSB67	RS13980	WP 010643500.1	EVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSTT	113
NTDB id 1193	PAKAF	RS26665	WP 003161249.1	VPAGRAIRNIDFQRGEKGEKNVVIDLSPTDIPIDVTQVGKEIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASDAQSDRT	236
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	PAPQGGIGNIGFQRGAQGEGLVVIDLQGTNTPVDVQQQGSKVVRFPGIKIPHTLARRLNTTDFATPVASIDSYNDGSNG	218
NTDB id 1052	ABD1	RS15850	WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
NTDB id 1039	HON27	RS01460	WP 031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
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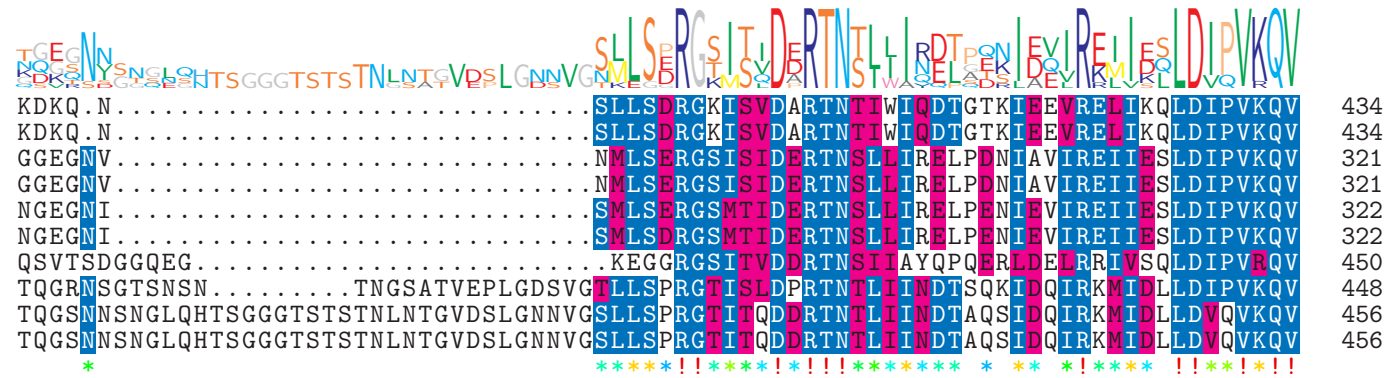


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consensus

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

698
698
571
571
582
580
714
709
721
721