

logo

NTDB id	1284	ACE17W	RS12370	WP	010946666.1	MRKIIAILILILI..SASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id	1257	GCO85	RS04855	WP	163621000.1	MRKIIAILILILI..SASLSLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id	1186	GTF74	RS01320	WP	000788426.1		0
NTDB id	1175	A1552VC	RS12105	WP	000788426.1		0
NTDB id	186894	FORC37	RS13930	WP	017419660.1		0
NTDB id	1408	DSB67	RS13980	WP	010643500.1		0
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	MNSGLSRLGIALLA..MFAPALLAADLEKLDVAALPGDRVELKLQFDEPVA..APRGYTIEQPARIALDLPGVQNKLGTK.	77
NTDB id	1009	ACIAD	RS15175	WP	004923730.1	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGQTEIRVMFNGLPP..QPQAYQLEQPARLILDFDKAGQKLAKS.	78
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NTDB id	1284	ACE17W	RS12370	WP	010946666.1	KTKTIQLGSLHSYNIVAVGDRVRAILEIDRSV..SYSGSTAGSIYSLVLNGKSNDFENSKE.....VFITNQ	144
NTDB id	1257	GCO85	RS04855	WP	163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILEIDRSV..SYSGSTAGSVYSLVLNGKSNDFENSKE.....VFITNQ	144
NTDB id	1186	GTF74	RS01320	WP	000788426.1	MKNGLKTYV..A.QTWLTLW.VGLALCA...SS.....M.VFSA	31
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	MKNGLKTYV..A.QTWLTLW.VGLALCA...SS.....M.VFSA	31
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NTDB id	1193	PAKAF	RS26665	WP	003161249.1	.NRELSVGNTRS SVTVVEAKDRTRLINLTLS.SYTRVEGNNLFVVVGNSPAGASVASTAPVKASPAPASYAQPIKPKP	155
NTDB id	1009	ACIAD	RS15175	WP	004923730.1	.LIPVTTNEASSIDVSSDDKRARVVNLK DAG.AHTTRVEGNVFIKINNIQSTATP.....VAT.SA.....	138
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NTDB id	1284	ACE17W	RS12370	WP	010946666.1	VVNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVCKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKK	224
NTDB id	1257	GCO85	RS04855	WP	163621000.1	VVNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVCKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKK	224
NTDB id	1186	GTF74	RS01320	WP	000788426.1	ESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTVDADDKLYLLDVKDFSTPVESVEVFRKEPS	111
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	ESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTVDADDKLYLLDVKDFSTPVESVEVFRKEPS	111
NTDB id	186894	FORC37	RS13930	WP	017419660.1	QSEAEANALKSIDFRNTAAKDAITIVELATPATVVDIKPVAEGLSIELINTSVKDEQLYLLDVKDFSTPVESIEVFRVSK	116
NTDB id	1408	DSB67	RS13980	WP	010643500.1	EEVQSNLSKSIDFRNTKDKDAVIVVELVSPA AIVDIKRVQEGLSIDLINTSVKDDQLYLLDVKDFSTPVESVEVFRDTST	112
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	YVPAGRAIRNIDFQRGEKGEGNVIDLSDP TLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASAQSDR	235
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NTDB id	1284	ACE17W	RS12370	WP	010946666.1	VRMTILSKCDYGHFVYQVNMQFMVDVFP LTAEEIRQAKLKKQVFTGKRISNFQNI STRAVLQLLADFTGINMVVSDKVQ	304
NTDB id	1257	GCO85	RS04855	WP	163621000.1	VRMTILSKCDYGHFVYQVNMQFMVDVFP LTAEEIRQAKLKKQVFTGKRISNFQNI STRAVLQLLADFTGINMVVSDKVQ	304
NTDB id	1186	GTF74	RS01320	WP	000788426.1	TQLVVTVDGEFHQHDYTLKGKYLEVVISKLADEKPK.PKSVLEKEBK LISNFQDIPVRNVQLLIADYNGFNLVVSDSVV	190
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	TQLVVTVDGEFHQHDYTLKGKYLEVVISKLADEKPK.PKSVLEKEBK LISNFQDIPVRNVQLLIADYNGFNLVVSDSVV	190
NTDB id	186894	FORC37	RS13930	WP	017419660.1	TRLVANVSCEFRHDYRLSGRYLEV TISRIKSEGEAR.EQSVLEKEBK LISNFQDIPVRNVQLLIADYNGFNLVVSDSVT	195
NTDB id	1408	DSB67	RS13980	WP	010643500.1	TRLVAYIDEDYSHDYRLTG RYLEVVKVGLKANEKAP.DKSILDKEBK LISNFQDIPVRNVQLLIADYNGFNLVVSDSVE	191
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	TSITIEPSGLYDYL VYQTDNRLTVS IKPMTTEDAERRKKDNFAYTCEKLSNFQDIDVR SVLQLIADFTDLNLVASDVQ	315
NTDB id	1009	ACIAD	RS15175	WP	004923730.1	GVLSIQSTGSYEYMAQAE NKLTISIKFPQ..DNVRLNSKNNQNYTCKKLSNFQDIEVRVQLLIADFTNINMTADSVQ	296
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consensus					



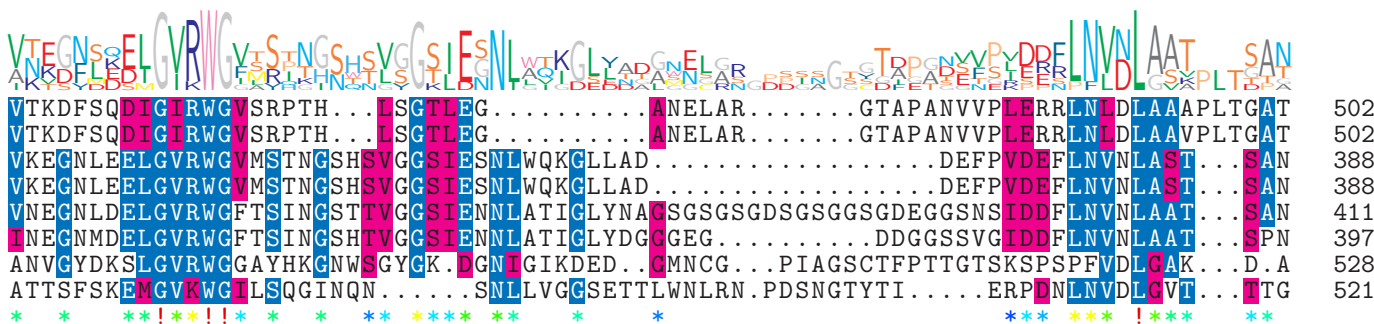
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consensus					



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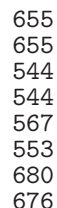
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NTDB id 1284 ACE17W RS12370 WP 010946666.1
NTDB id 1257 GCO85 RS04855 WP 163621000.1
NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 186894 FORC37 RS13930 WP 017419660.1
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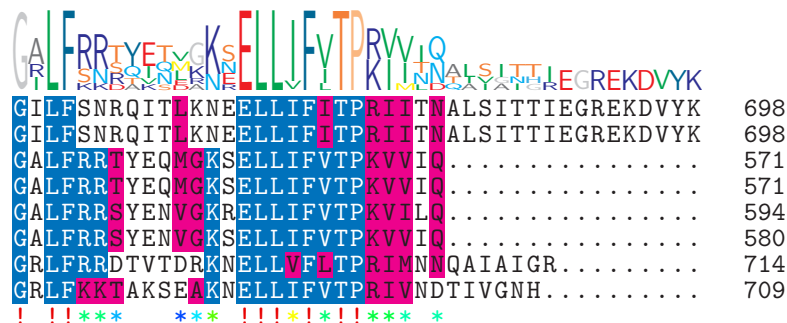
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NTDB id 1284 ACE17W RS12370 WP 010946666.1
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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 186894 FORC37 RS13930 WP 017419660.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1193 PAKAF RS26665 WP 003161249.1
NTDB id 1009 ACIAD RS15175 WP 004923730.1
consensus

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