

logo

NTDB id 1284	ACE17W	RS12370	WP 010946666.1	MRKIIAILII..ISASISLMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFISTDMQIPSDQ	78
NTDB id 1257	GCO85	RS04855	WP 163621000.1	MRKIIAILIL..ISASISLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFISTDMQIPSDQ	78
NTDB id 1186	GTF74	RS01320	WP 000788426.1		0
NTDB id 1175	A1552VC	RS12105	WP 000788426.1		0
NTDB id 22760	VP	RS13485	WP 024699327.1		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	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFNGLPP.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
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NTDB id 1284	ACE17W	RS12370	WP 010946666.1	STITVINEASSVDVSDDDRRRLINLKLGLSYTRVEGNFSLWNLLEASNVKKEKALGSTRHYNITLXKRQSAUTTEMDPZYKESKSLQSYLKTENASNDNFONKEHVKASPLPYVSFLITSAE	145
NTDB id 1257	GCO85	RS04855	WP 163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSLVLNGKSNDLFENSKEV.....FITNQV	145
NTDB id 1186	GTF74	RS01320	WP 000788426.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSLVLNGKSNDLFENSKEV.....FITNQV	32
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	MKNGLKTY..VAQTWLTWVGL...ALCASSMV.....F..SAE	32
NTDB id 22760	VP	RS13485	WP 024699327.1	MKMRQGLTRTPRMLTGLFSLWLFL...LLAP...L.....A..AAE	33
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTLGLFSLWFL...LLAP...F.....V..AAE	33
NTDB id 1193	PAKAF	RS26665	WP 003161249.1	KNRELISVGNTRSVTVVEAKDRTRLIINLTTLSSYTTTVEGNLNFVVGNSPAGASVASTAPVKASPAPASYAQPIKPKPY	156
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	SLIPVTTNEASSIDVSSDDKRARVVVNLKDGAGFTTRVEGNVFILKINNIQSTAT.....PVAT..SA.....	138
NTDB id 1052	ABD1	RS15850	WP 001017033.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDGAGFTTRVEGNTFILKINSAQTSNK.....PLPV..VS.....	138
NTDB id 1039	HON27	RS01460	WP 031993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDGAGFTTRVEGNTFILKINSAQTSNK.....PLPV..VS.....	138
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NTDB id 1284	ACE17W	RS12370	WP 010946666.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	225
NTDB id 1257	GCO85	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 22760	VP	RS13485	WP 024699327.1	DVQSNALKSIDFRTNKS KDAVIVVELASPA AIVDVKRVQEGLSIDL MNTSVKDEQLYLLDVKDFATVVESVEVFRDTSTT	113
NTDB id 1408	DSB67	RS13980	WP 010643500.1	EVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDKRVQEGLSIDL LNTSVKDDQLYLLDVKDFSTTVVESVEVFRDTSTT	113
NTDB id 1193	PAKAF	RS26665	WP 003161249.1	VPAGRAIRNIDFQRGEKGE GNVVIDLSDPITLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASASQSDRT	236
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	PAPQGGIGNIGFQRGAQGEGLVVIDLQGTNTPVDVQQQGSKVVRFPGKIPKIPHTLARRLNTTDFATPVASIDSYNDGSNG	218
NTDB id 1052	ABD1	RS15850	WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
NTDB id 1039	HON27	RS01460	WP 031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
consensus				* * * * *	

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NTDB id 1052 ABD1 RS15850 WP 001017033.1
NTDB id 1039 HON27 RS01460 WP 031993779.1
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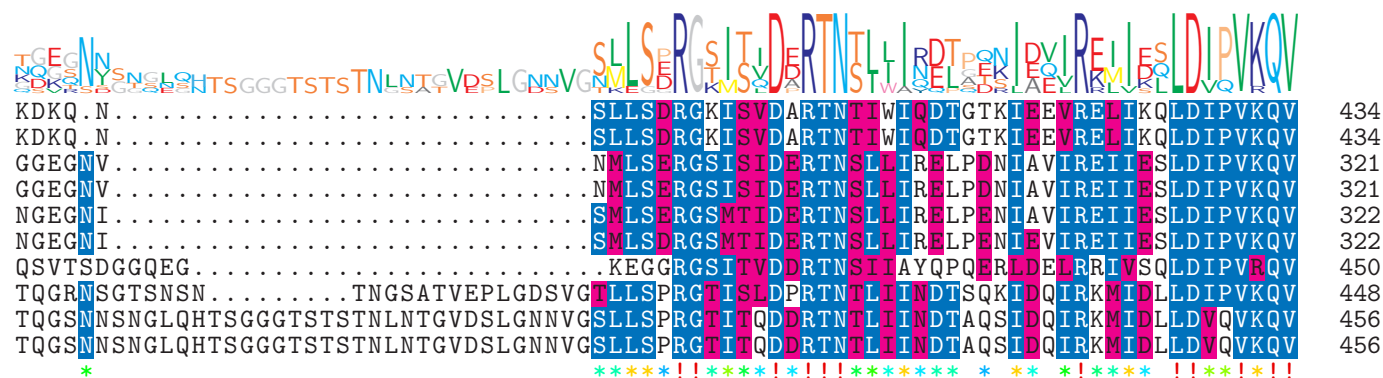
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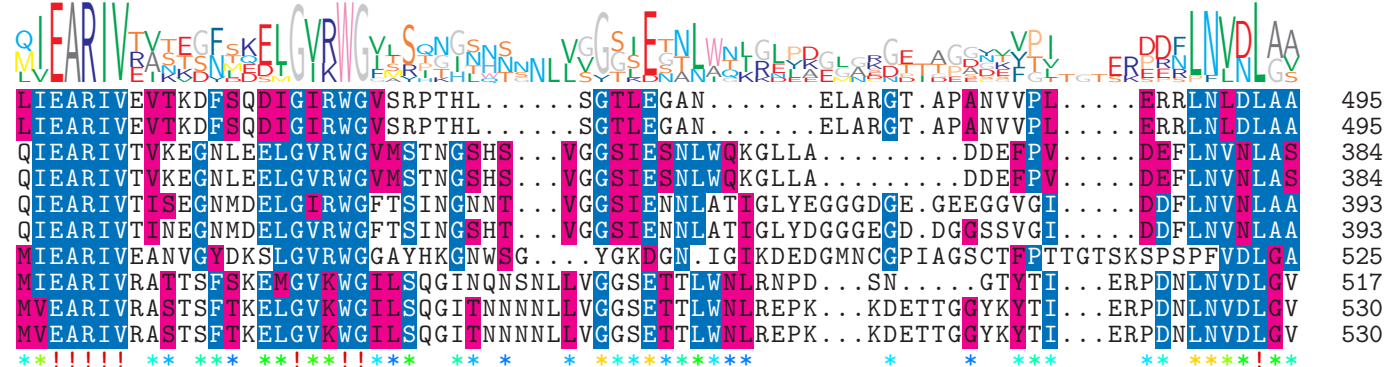
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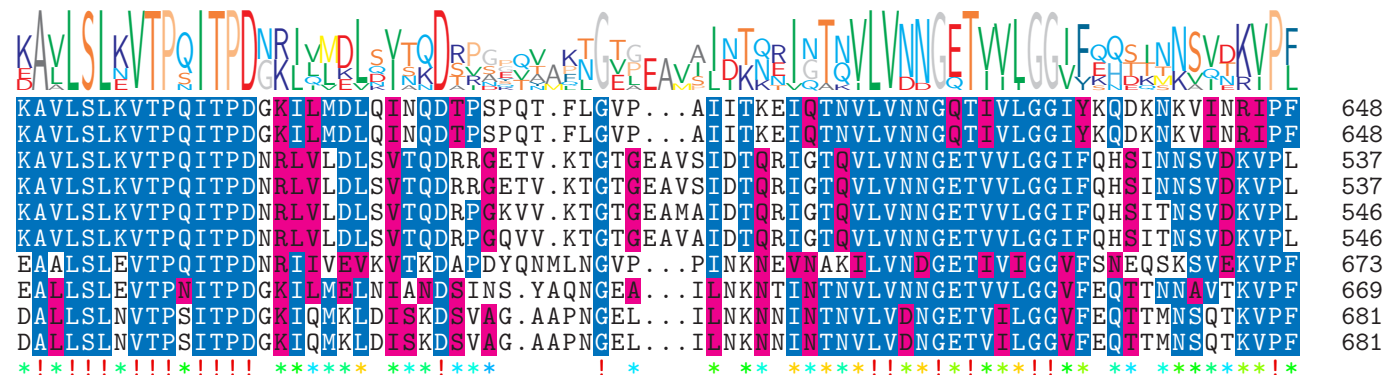
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consensus

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logo



☐ non conserved
☒ similar
☒ ≥ 50% conserved