

NTDB id	Accession	Length	Consensus
1284	ACE17W RS12370 WP 010946666.1	78	MRKTIAILIL...ISASISMAYSKDNSLSISKVLPPEPENVRRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIIPSDQ
1257	GC085 RS04855 WP 163621000.1	78	MRKTIAILIL...ISASISLAYSKDNSLSISKVLPPEPENVRRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIIPSDQ
1066323	ACG4I1 RS07865 WP 000788426.1	0	
1186	GTF74 RS01320 WP 000788426.1	0	
1175	A1552VC RS12105 WP 000788426.1	0	
1408	DSB67 RS13980 WP 010643500.1	0	
1193	PAKAF RS26665 WP 003161249.1	76	MNSGLSRLGIALAA.MFAPALLAADTEKDVAAPGDRVEIKLQFDEPVA.APRGYTIEQPARIALDLPGVQNKIG..T
1009	ACIAD RS15175 WP 004923730.1	77	MKQIRQFSIGTVAYAIMQVASAQVAITNVPMNIPGIGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKAGQKLA..K
1039	HON27 RS01460 WP 031993779.1	77	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLK..Q
1052	ABD1 RS15850 WP 001017033.1	77	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQQGLK..Q
consensus			* * * * * * * * * * * * * * * * * *

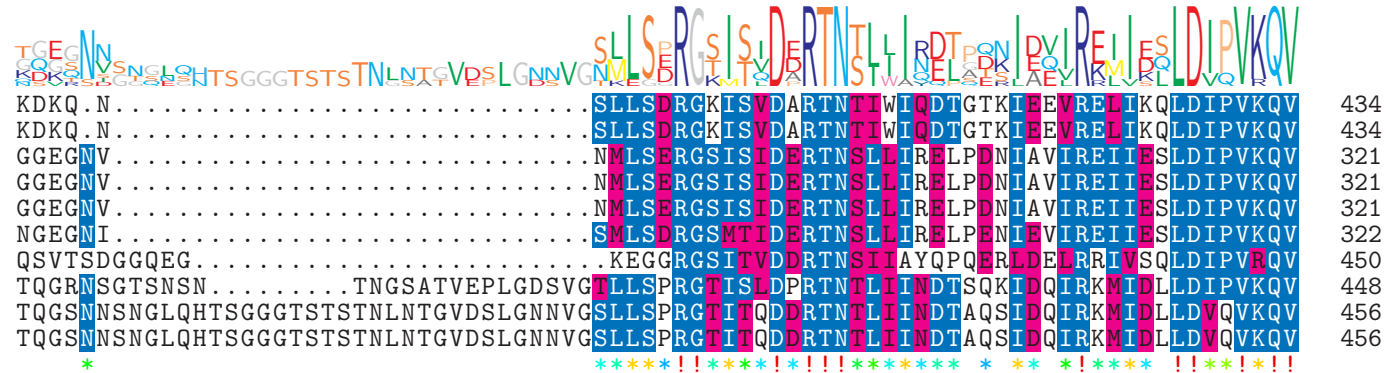
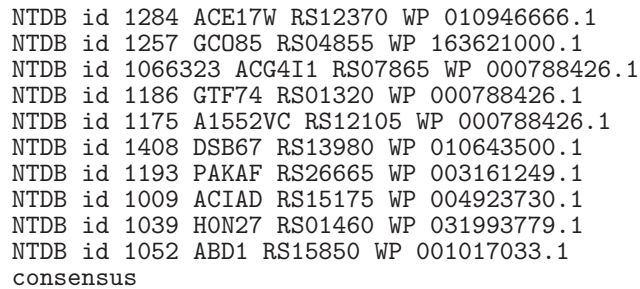
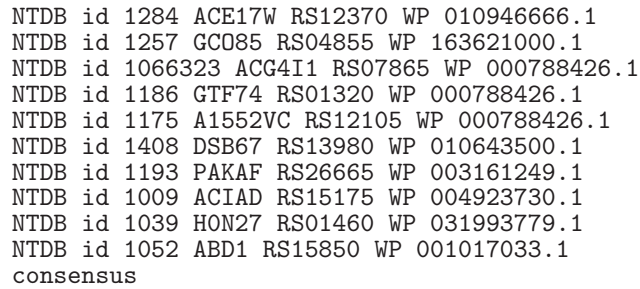
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NTDB id	1257	GC085	RS04855	WP	I63621000.1	KTKTIQLGSLHSYNIVAVGDVRRAILELDRSVSYSGSTAGSVYSVLVNGKSNDFENSKE.....VFITNQV	145
NTDB id	1066323	ACG4I1	RS07865	WP	O00788426.1	MKNGLKT..YVAQTLWLTLWG....LAICASS.....MVFSA.E	32
NTDB id	1186	GTF74	RS01320	WP	O00788426.1	MKNGLKT..YVAQTLWLTLWG....LAICASS.....MVFSA.E	32
NTDB id	1175	A1552VC	RS12105	WP	O00788426.1	MKNGLKT..YVAQTLWLTLWG....LAICASS.....MVFSA.E	32
NTDB id	1408	DSB67	RS13980	WP	O10643500.1	MKMRHGLTLTPRKLTGLFSLWFS....LLLAP.....FVAA.E	33
NTDB id	1193	PAKAF	RS26665	WP	O03161249.1	KNREISVGNTRSVTVVEAKDRTLLINLTTLSSYTTRVEGNLFVVVGNSPAGASVASTAPVKASPAPASYAQPIKPYP	156
NTDB id	1009	ACTAD	RS15175	WP	O04923730.1	SLIPVTNEASSIDVSSDDKRARVVNLKDGAFTTRVEGVNFILKINNIIQSTAT.....PVAT..SA.....	138
NTDB id	1039	HON27	RS01460	WP	O31993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDTGAFSTRVEGNTFILKINSQAQTSNK.....PLPV..VS.....	138
NTDB id	1052	ABD1	RS15850	WP	O01017033.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDTGAFSTRVEGNTFILKINSQAQTSNK.....PLPV..VS.....	138
consensus						* * * * *	

NTDB id	1284	ACE17W	RS12370	WP	O10946666.1	VNAKHEIKRIDFRGIEKKQGGRLIDVSDTGIPIDVTQVGKEIVVNFLNNTKIPLNLMKRYDVSDFHSVPVQIITMQQEGKKV	225
NTDB id	1257	GC085	RS04855	WP	163621000.1	VNAKHEIKRIDFRGIEKKQGGRLIDVSDTGIPIDVTQVGKEIVVNFLNNTKIPLNLMKRYDVSDFHSVPVQIITMQQEGKKV	225
NTDB id	1066323	ACG4I1	RS07865	WP	000788426.1	SATANQLENIDFRVNKEKA AVLIVE LASPSAVVDVQKVQEGLSTIELLKTVDVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB id	1186	GTF74	RS01320	WP	000788426.1	SATANQLENIDFRVNKEKA AVLIVE LASPSAVVDVQKVQEGLSTIELLKTVDVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	SATANQLENIDFRVNKEKA AVLIVE LASPSAVVDVQKVQEGLSTIELLKTVDVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB id	1408	DSB67	RS13980	WP	010643500.1	EVQSNSLSKSIDFRTNKD KDAVI VVELVSPAA IVD IKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTPVESVEVFRDTSTT	113
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	VPAGRAIRNIDFQRGEKGEGNVVIDLSDPPTLSPDIQQGGKIRLDFAKTQLPDALRVRDLVKDFATPVQFVNASAQSDRT	236
NTDB id	1009	ACTAD	RS15175	WP	004923730.1	PAPQGIGIGNIGFQRGAQGEGLVVIDLQGTNPVDVQQQGSKWVRFPGIKIPTHLARRLNTTD FATPVASIDS YND GSNG	218
NTDB id	1039	HON27	RS01460	WP	031993779.1	.AQPQGVSNI GFQRGSQGEGLVVVDL LGSNT PVDVQQQGSKWVRT IGTKIPTHLARRLNVNDF ATPVSS IDAYND KGVG	217
NTDB id	1052	ABD1	RS15850	WP	001017033.1	.AQPQGVSNI GFQRGSQGEGLVVVDL LGSNT PVDVQQQGSKWVRT IGTKIPTHLARRLNVNDF ATPVSS IDAYND KGVG	217
consensus						* * * * * ! * ! * * ***** * !! * ! * ! * ! * ! * ! * ! * ! * ! * ! * ! *	

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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
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NTDB id 1193 PAKAF RS26665 WP 003161249.1
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consensus

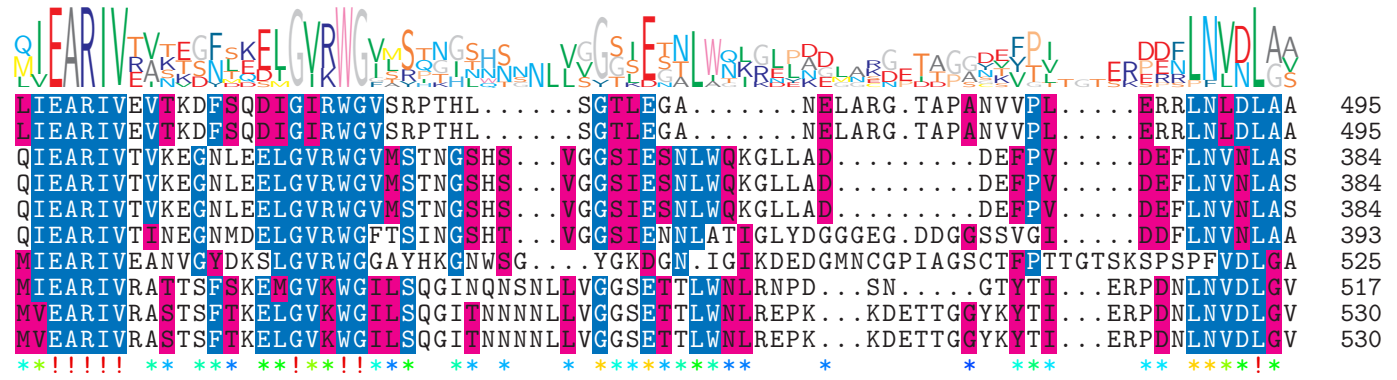
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consensus

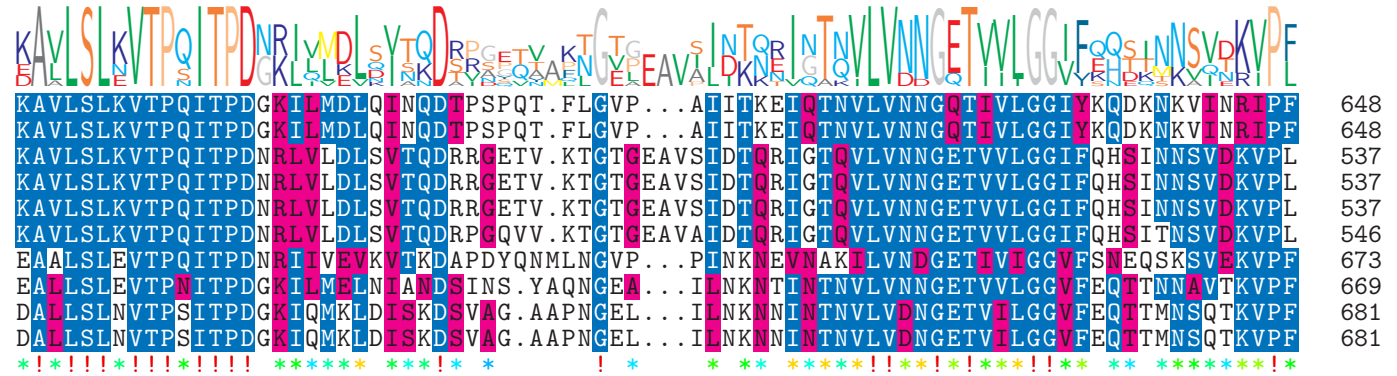
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NTDB id 1039 HON27 RS01460 WP 031993779.1
NTDB id 1052 ABD1 RS15850 WP 001017033.1
consensus

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logo



X non conserved
X similar
X ≥50% conserved