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NTDB id 1284	ACE17W	RS12370	WP 010946666.1	MRKIIAILII..ISASISLMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1257	GCO85	RS04855	WP 163621000.1	MRKIIAILIL..ISASISLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1186	GTF74	RS01320	WP 000788426.1		0
NTDB id 1175	A1552VC	RS12105	WP 000788426.1		0
NTDB id 83741	VA249	RS12950	WP 420864276.1		0
NTDB id 1408	DSB67	RS13980	WP 010643500.1		0
NTDB id 1193	PAKAF	RS26665	WP 003161249.1	MNSGLSRLGIALLAA.MFAPALLAADLEKLDVAALPGDRVELKLQFDEPVA.APRGYTIEQPARTIALDLPGVQNKLG..T	76
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	MKQIIRQFSIGTVAYAIMQVSAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARLLIDFDKAGQKLA..K	77
NTDB id 1052	ABD1	RS15850	WP 001017033.1	MNHVFRQFSMGAVAIAIMQAASQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLLIDFDKAQQGLK..Q	77
NTDB id 1039	HON27	RS01460	WP 031993779.1	MNHVFRQFSMGAVAIAIMQAASQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLLIDFDKAQQGLK..Q	77
consensus				* * * * *	

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NTDB id 1284	ACE17W	RS12370	WP 010946666.1	STIITYTNEASVDVSDDDVRRLIYNLKRSLSYTRVEGNFSLWNNLELASSSKPKKSLGTHLYNITLXKRQSGTHTMPLTAVRTSSSKTSSYLYVKTENASNDAAECAKEPVKASPLPYVSVFISAE	145
NTDB id 1257	GCO85	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	MKNGLKT..YVAQTLWTLWVG....LALCASS.....MV.FSAE	32
NTDB id 83741	VA249	RS12950	WP 420864276.1	MKNGLKT..YVAQTLWTLWVG....LALCASS.....MV.FSAE	21
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKNGLKT..YVAQTLWTLWVG....LALCASS.....MV.FSAE	33
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NTDB id 1009	ACIAD	RS15175	WP 004923730.1	KNREL SVGNTRSVTVVEAKDRTRLIINLTTLSSYTTRVEGNLNFVVGNSPAGASVASTAPVKASPASYAQPIKPKPY	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
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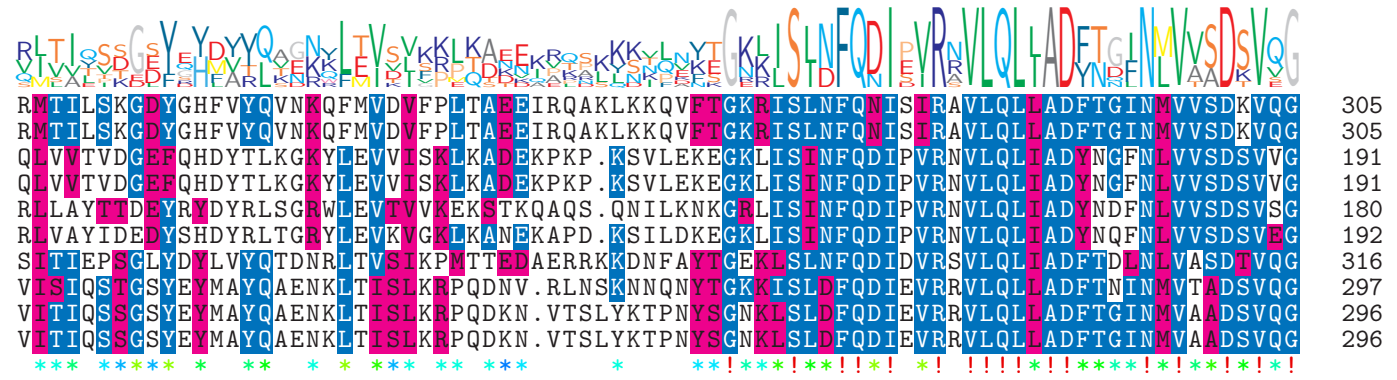
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NTDB id 1257	GCO85	RS04855	WP 163621000.1	VNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	225
NTDB id 1186	GTF74	RS01320	WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
NTDB id 83741	VA249	RS12950	WP 420864276.1	ASPPNELKSIDFRADNNT EAVIMIELTSSSVVVDIQRERDGLSIELLNSTSVSDNQLHLLDVRDFSTVIKQIDVSRYHSAT	101
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NTDB id 1193	PAKAF	RS26665	WP 003161249.1	VPAGRAIRNIDFQRGEKGEKNVVIDLSDPTLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASDAQSDRT	236
NTDB id 1009	ACIAD	RS15175	WP 004923730.1	PAPQGGIGNIGFQRGAQGEGLVVIDLQGTINTPVDVQQQGSKVVRFPGIKIPHTLARRLNTTDFATPVASIDSYNDGSNG	218
NTDB id 1052	ABD1	RS15850	WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
NTDB id 1039	HON27	RS01460	WP 031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
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consensus

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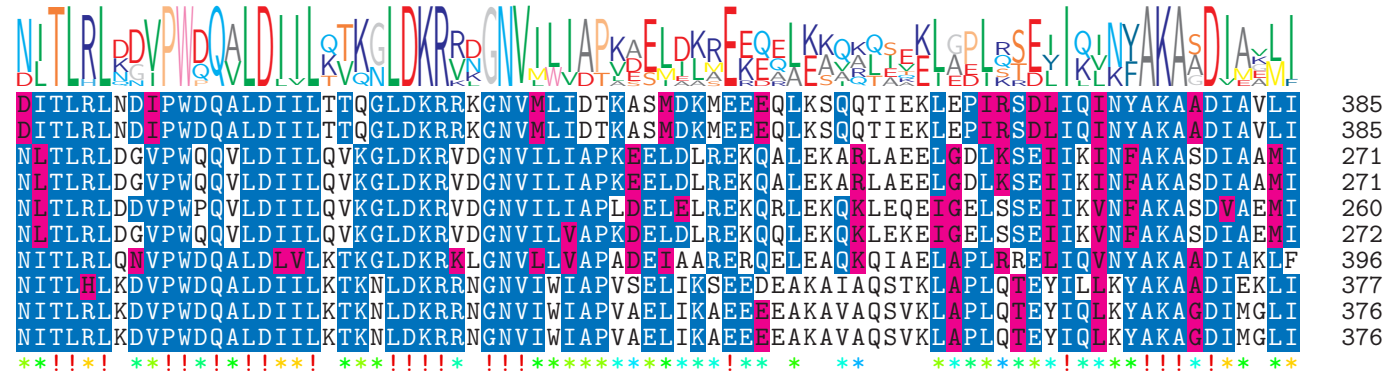


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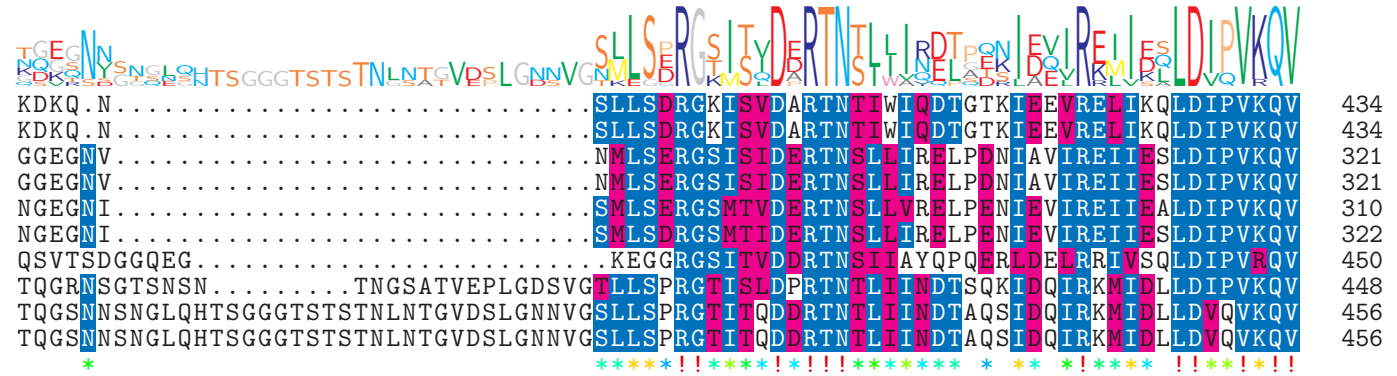


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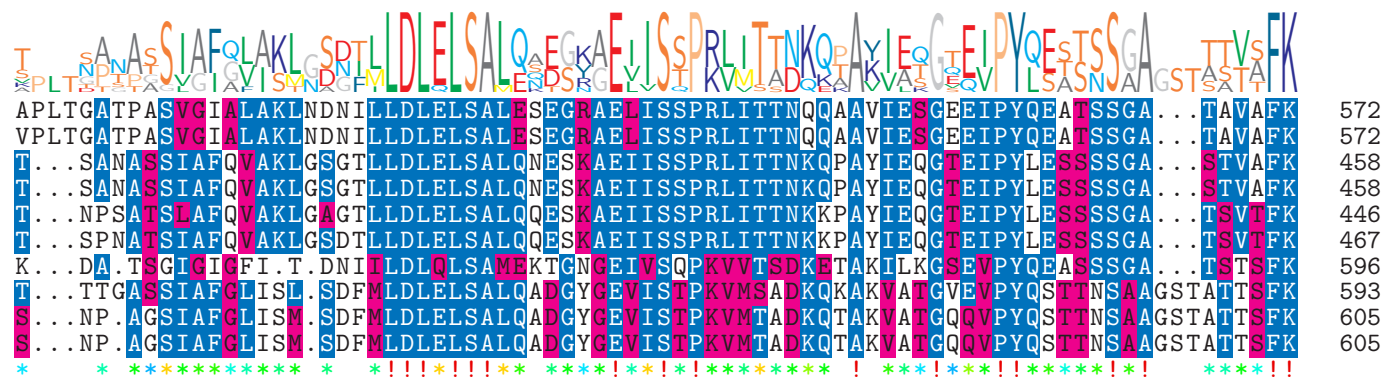
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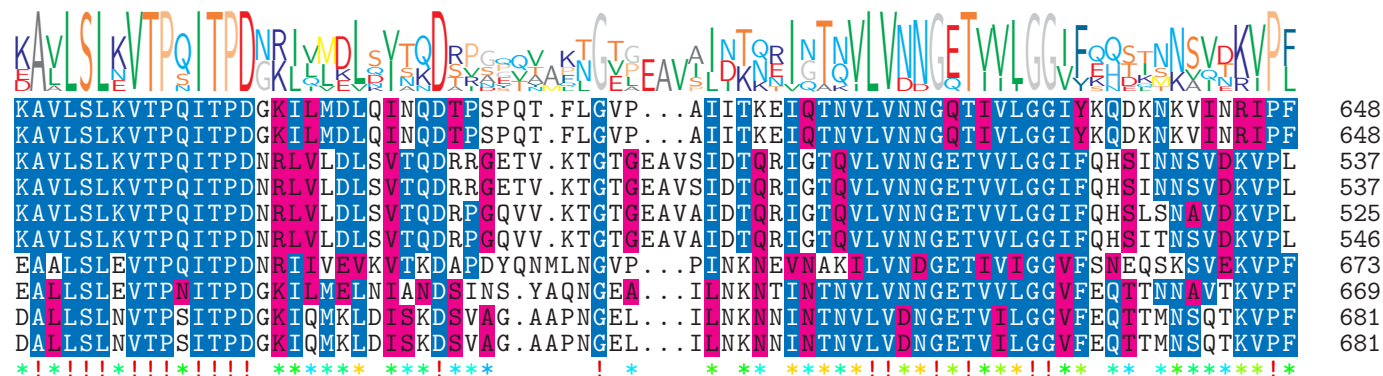
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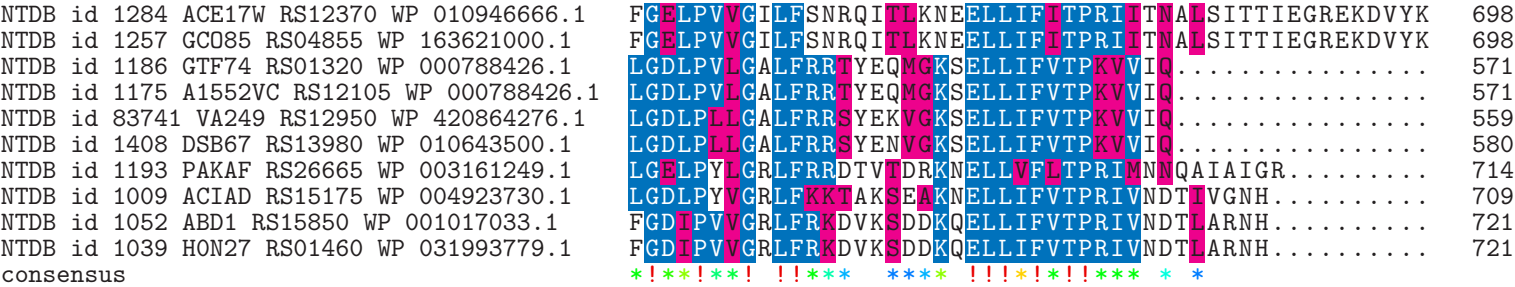
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NTDB id 1039 HON27 RS01460 WP 031993779.1
consensus
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logo



X non conserved
X similar
X ≥ 50% conserved