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NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKIIAILI...ISASISMAYSKDNSISVKVLPPEPVRVDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1257 GCO85 RS04855 WP 163621000.1	MRKIIAILI...ISASISLAYSKDNSISVKVLPPEPVRVDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 70388 VCA1004 RS01055 WP 408646887.1		0
NTDB id 1186 GTF74 RS01320 WP 000788426.1		0
NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNSGLSRLGIALIAA.MFAPALLAADIEKIDVAALPGDRVEIKLQFDEPVA.APRGYTIEQPARIALDLPGVQNKIG..T	76
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGQGTEIRVMFENGLPP.QPQAYQLEQPARLLDFDKAGQKIA..K	77
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFENGLPP.QPQAYQLENPSRLILDFDKAQQGK..Q	77
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFENGLPP.QPQAYQLENPSRLILDFDKAQQGK..Q	77
consensus	* * * * *	

logo

NTDB id 1284 ACE17W RS12370 WP 010946666.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTACSIYSVLVNGKSNDLFENSK.....EVFITNQV	145
NTDB id 1257 GCO85 RS04855 WP 163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTACSVYSVLVNGKSNDLFENSK.....EVFITNQV	145
NTDB id 70388 VCA1004 RS01055 WP 408646887.1	MAAPTT.....T.....STVS.....ITQ.TSVQ	18
NTDB id 1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQTWLTWVGLA....LCAS.....SMV.FSAE	32
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTWLTWVGLA....LCAS.....SMV.FSAE	32
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKMRHGILTLTPRKLTGLESLWFSLL....LAP.....F.VAAE	33
NTDB id 1193 PAKAF RS26665 WP 003161249.1	KNRELSVGNTRSVTVVEAKDRTRLIINLTTLSSYTTRVEGNLFFVVGNSPAGASVASTAPVKASPAPASYAQPIKPKPY	156
NTDB id 1009 ACIAD RS15175 WP 004923730.1	SLIPVTTNEASSIDVSSDDKRARVVVNLKDAGAFTRVEGNVFIILKINNISTAT.....PVAT..SA.....	138
NTDB id 1052 ABD1 RS15850 WP 001017033.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDAGAFTRVEGNTHILKINSAQTSNK.....PLPV..VS.....	138
NTDB id 1039 HON27 RS01460 WP 031993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDTGAFTRVEGNTHILKINSAQTSNK.....PLPV..VS.....	138
consensus	* * * * *	

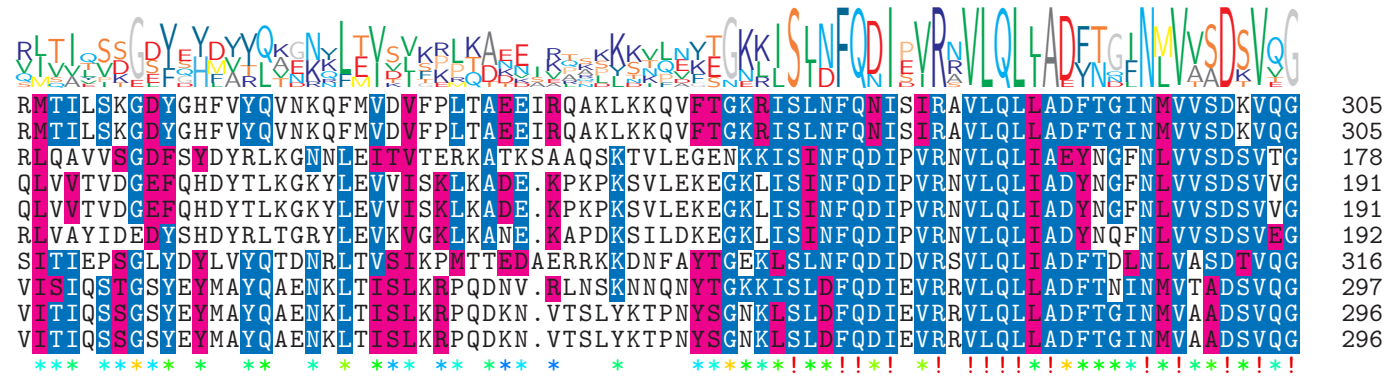
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NTDB id 1284 ACE17W RS12370 WP 010946666.1	VNAKHEIKRIDFRGIEKQGGRVLIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQEGKKV	225
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
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NTDB id 1193 PAKAF RS26665 WP 003161249.1	VPAGRAIRNIDFRQGEKGEENVVIDLSDPTLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASQSDRT	236
NTDB id 1009 ACIAD RS15175 WP 004923730.1	PAPQCGIGNIGFQRGAQGEGLVVVDLQGINTPVDVQQQGSKVVRFPGIKIPTHLARRLNTTDFATPVASIDSYNDGSNG	218
NTDB id 1052 ABD1 RS15850 WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVRTITGKIPTHLARRLNVNDFATPVSSIDAYNDKGVG	217
NTDB id 1039 HON27 RS01460 WP 031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVRTITGKIPTHLARRLNVNDFATPVSSIDAYNDKGVG	217
consensus	* * * * *	

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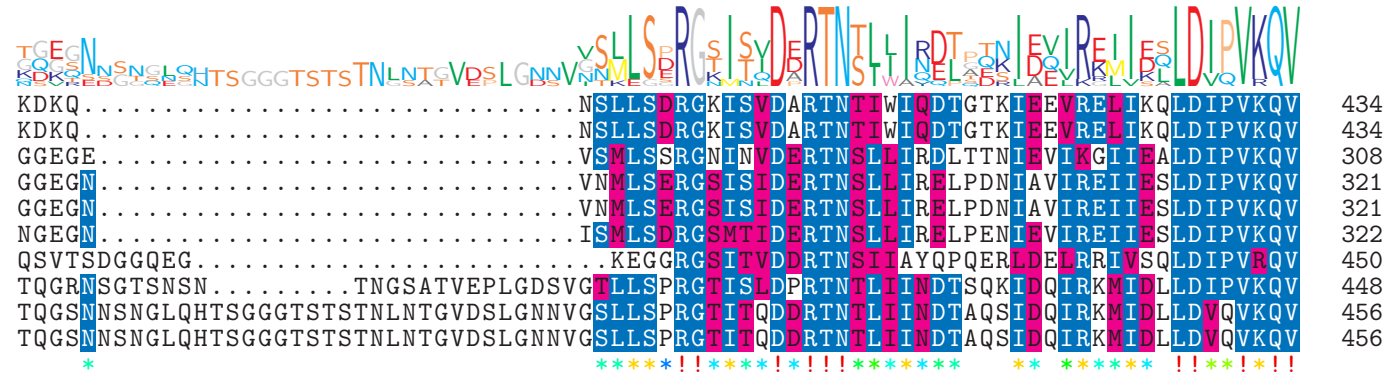
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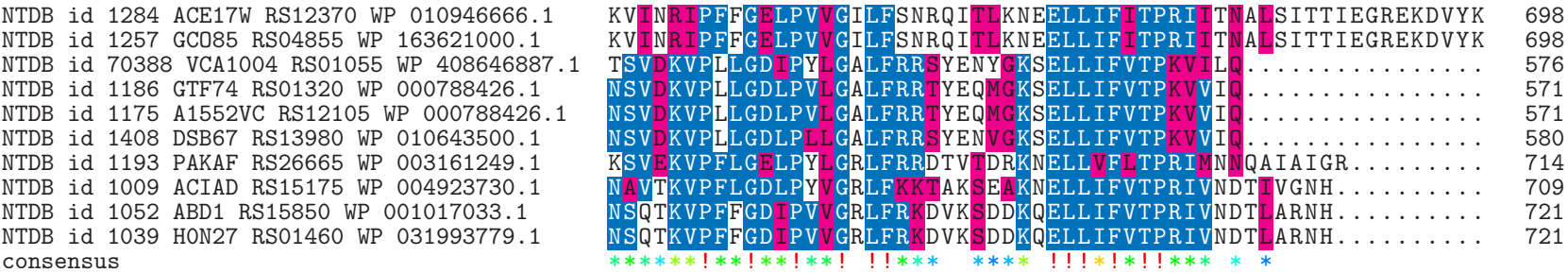
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NTDB id 1039 HON27 RS01460 WP 031993779.1
consensus

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logo



 non conserved
 similar
 ≥ 50% conserved