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NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKIIAILII..ISASISMAYSKDNSIISVKVLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFISTDMQIPSDQ	78
NTDB id 1257 GC085 RS04855 WP 163621000.1	MRKIIAILIL..ISASISLAYSKDNSIISVKVLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFISTDMQIPSDQ	78
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 1009 ACIAD RS15175 WP 004923730.1	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGQGTEIRVMFNGLPP.QPQAYQLEQPARLIIDFDKAGQKLA..K	77
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQAAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLIIDFDKAGQGLK..Q	77
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consensus	* * * * *	

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NTDB id 1284 ACE17W RS12370 WP 010946666.1	STIITYTNEASVDVSDDDRRRLIYNLKRSLSYTRVEGNFSLVYNLSTKSKKKEKELALGTHRYNITLXKRYGATHEMPYVYRTSSSTAGSYTLKIKNSFALALXAPVKASAXXASYKESFFENQV	145
NTDB id 1257 GC085 RS04855 WP 163621000.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGTAGSIYSLVLNGKSNDLFENS.....KEVFITNQV	145
NTDB id 1186 GTF74 RS01320 WP 000788426.1	KTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGTAGSVYSLVLNGKSNDLFENS.....KEVFITNQV	32
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTWLTWLWGLALC.....AS.....SMVFSAE	32
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKNGLKT..YVAQTWLTWLWGLALC.....AS.....SMVFSAE	33
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKMRHGLTLTPRKLTGLFSLWFSLLLA.....P.....FVAEE	138
NTDB id 1039 HON27 RS01460 WP 031993779.1	SLIPVTTNEASSIDVSSDDKRRRVVNLKDAAGATTTRVEGNVFILKINNIQSTAT.....PVAT..SA.....	138
NTDB id 1052 ABD1 RS15850 WP 001017033.1	SKISVATNEASVSDVTSDDQRSRLTVNLKDTGATTTTRVEGNTFILKINSAQTSNK.....PLPV..VS.....	138
NTDB id 383293 FY041 RS25860 WP 082426354.1	SKISVATNEASVSDVTSDDQRSRLTVNLKDAAGATTTRVEGNTFILKINSAQTSNK.....PLPV..VS.....	0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	KNREL SVGNTRSVTVVEAKDRTRLIINLTTLSSTYTTTRVEGNLFFVVGNSPAGASVASTAPVKASPAPASYAQPIKPKPY	156
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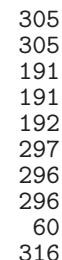
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPST	112
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NTDB id 1009 ACIAD RS15175 WP 004923730.1	PAPQGGIGNIGFQRGAQGEGLVVIDLQGTNTPVDVQQQGSKVVRFPGKIPKIPHTLARRLNTTDFATPVASIDSYNDGSNG	218
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NTDB id 1052 ABD1 RS15850 WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVG	217
NTDB id 383293 FY041 RS25860 WP 082426354.1		0
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consensus	* * * * *	

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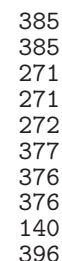
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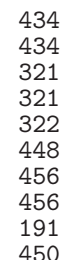
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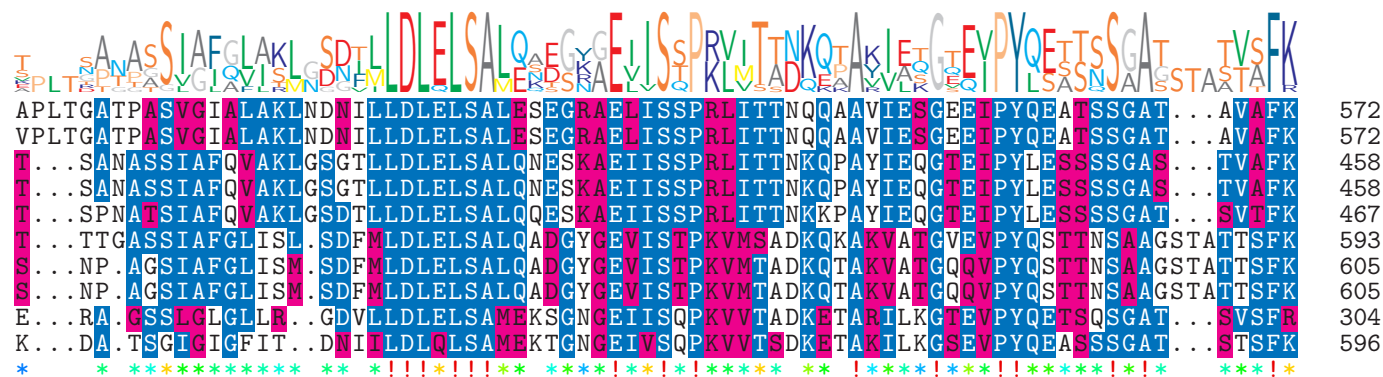
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consensus

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consensus

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NTDB id 1175 A1552VC RS12105 WP 000788426.1	GDLPVVGALFRRTYEQMGKSELLIFVTPKVVIQ.....	571
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NTDB id 1193 PAKAF RS26665 WP 003161249.1	GELPYLGRFLFRRTDTVTDRKNELLVFLTPRIMNNQAIAIGR.....	714
consensus	!***!***!***!*** * * !!!*!***!*** * *	

- ⓧ

 non conserved
- ✖

 similar
- ⓧ

 ≥ 50% conserved