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NTDB id 92474 MFE95 RS08080 WP 016504529.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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MFHENPLCTV.....SAMTGEVFEIQQALW25
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consensus



Sequence	Position	Conservation
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ERNQQQQSLLLRYFALPLKEENNRLWLGVDSLSNLSACETIAFITGKPVPEPILLESSQLKELLQQLTPNQMVEEVK..	105	High
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CQQLGLRELITRHNALPLHRTPTSTLLLAVADPTNQAEEDFRFATGLQVELVLADFRELSTAIRRLYGRSLGHEKSGLK..	151	High
CQTLGLRDLITRHNALPLNRSTSTLILAVADPTNLQAEEDFRFATGLQVELVLADFRELTAAIRRLYGRSLGHEKSGLK..	151	High
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logo

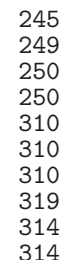
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consensus

[illegible]

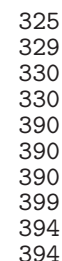
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consensus

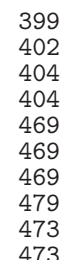
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consensus



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NTDB id 1384 A4U84 RS08525 WP 010787006.1	CDECYQGYRGRIGVYQCLSR	TAKK.....FDKQTAYLDFSSSLRESAKQK	443
NTDB id 92474 MFE95 RS08080 WP 016504529.1	QSACRCYQGYKGRIGVYQFLQKDKT	GCFVTDFFSSSLRESAKEK	444
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	SLLEMGFTEQDLAHPDFRVFQPVGC	LECREGYKGRVGIYEVMMKVTPETSKIIMEDGNALEIAAAASEKLGFNNLRRSGLKK	553
NTDB id 1018 ACIAD RS01685 WP 004920473.1	SLLEMGFTEPDLAQPEFQIFEPVGCH	DCREGYKGRVGIYEVMMKITPETSKIIMEDGNALEIAAATAETLGFNNLRRSGLKK	553
consensus	***	*****!*****	

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	IEQQLTNETEVDRLVGYE.	461
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	VMQGVTSLQEINRVTSSE..	570
consensus	*****!*!!!!*	

- [X] non conserved
- [X] similar
- [X] ≥ 50% conserved