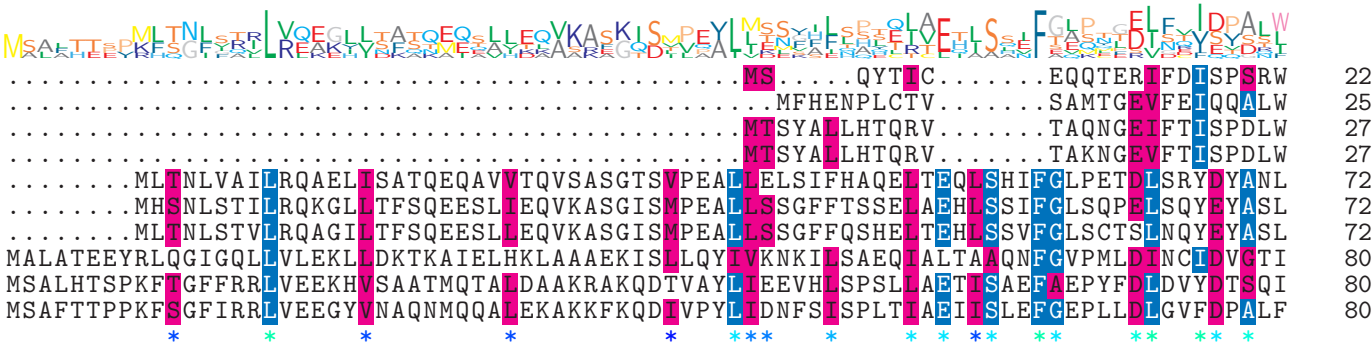


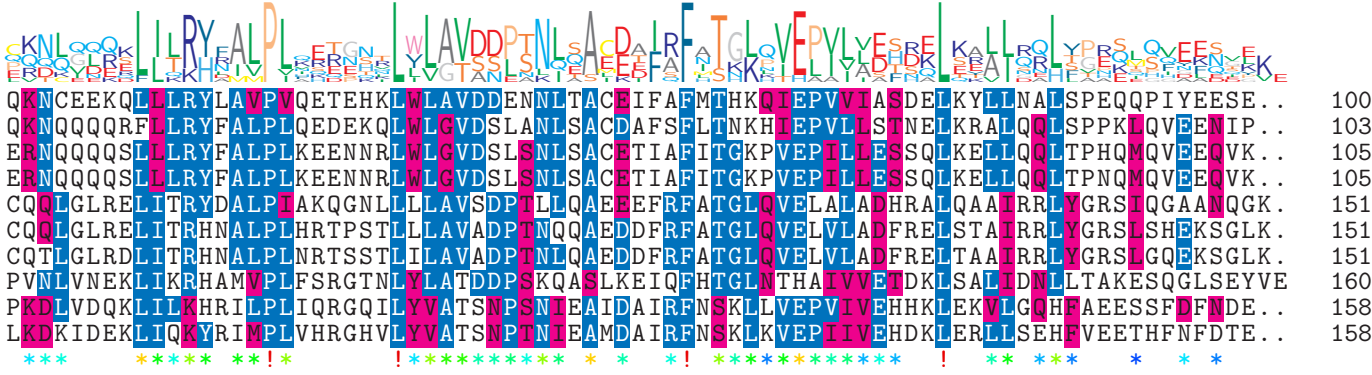
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



logo

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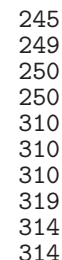


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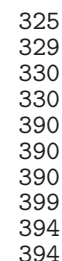
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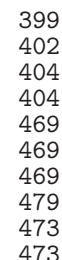
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consensus



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NTDB id 221525 BTV67 RS07910 WP 010906472.1	QSACRCYQGYKGRIGVYQFLQ	KDKT.....GCFVTD	FSSSLRESAKEK	444								
NTDB id 1366 NTHI RS01950 WP 011271972.1	INSCDCHQGYRGRIGVYQFLH	WQQN.....G.YQTD	FENLRESGLEK	445								
NTDB id 1355 HI 0298 AAC21962.1	ANSCDCHQGYRGRIGVYQFLH	WQQN.....D.YQTD	FKNLRASGLEK	445								
NTDB id 1168 A1552VC RS11070 WP 000957200.1	LQSQFAFQPNIEI...LYEANAAGCNE	CTGYSGRVGIYEVMAFNTELAEA	IMQRASIHQIERLAKANGMQTLQESGLEK	545								
NTDB id 1293 VP RS12240 WP 005479695.1	L.QHLGIQTDDN...IFRANPDGCNE	CTHGYSGRTGIYEVMRFDSEAL	LIKGASVHELEKLAIANGMSTLQMSGIEK	544								
NTDB id 1401 DSB67 RS12665 WP 010643259.1	L.QHIGIANTEQ...IFQANPDGCNE	CTHGYSGRTGIYEVMMKFDES	SEALIKGASVHELEKLAIANGMQTLQMSGIEK	544								
NTDB id 1251 GC085 RS07725 WP 011946523.1	GLIELGFKEADLV..NLKLYKAVGCE	QCTSGYRGRVGLFEVLPMTKEL	GQLIMSGGNSLDILKLAQSEGMLTIFQSGIEK	557								
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SLLEMGFTEQDLAHPDFRVFQPV	GCLEREGRYKGRVGIYEVMMKVTP	ETISKIIMEDGNALEIAAAASEKLG	FNNLRRSGLKK	553							
NTDB id 1018 ACIAD RS01685 WP 004920473.1	SLLEMGFTEPDLAQPEFQIFEPV	GCHDCREGRYKGRVGIYEVMMKIT	PEISKIIMEDGNALEIAAATAETL	GFNNLRRSGLKK	553							
consensus	***	*	** *! !!!!!!!***** *	*	** *	** *	** *	** *	** *	***	***	!!!!

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	IEQQLTNETEVDRLVGYE.	461
NTDB id 221525 BTV67 RS07910 WP 010906472.1	VRLGMTDDNEVQRVLGE..	461
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	VMQGVTSLQEVNRVTSE..	570
consensus	*****! *	!!!!!! *

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