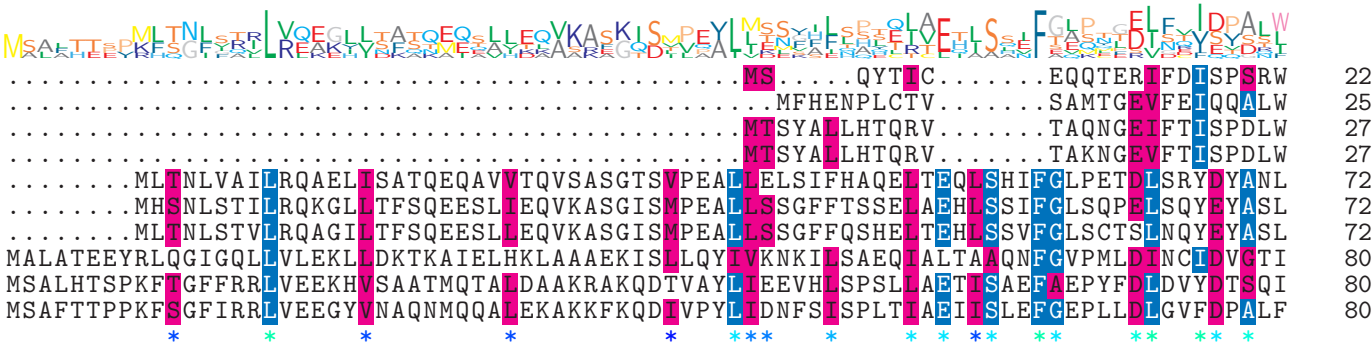


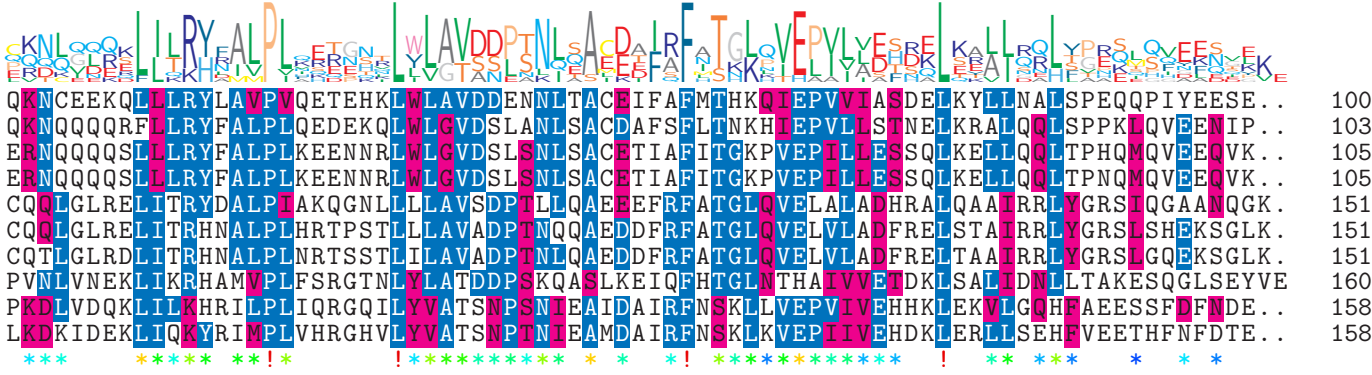
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



logo

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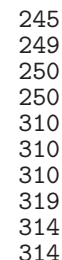


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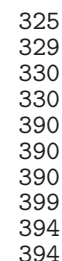
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consensus



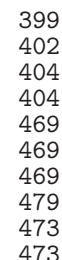
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consensus



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NTDB id 396363 GDY38 RS06070 WP 016504529.1	QSACRCYQGYKGRIGVYQFLQKDKT.....GCFVTDFFSSSLRESAKEK	444
NTDB id 1366 NTHI RS01950 WP 011271972.1	INSCDCHQGYRGRIGVYQFLHWQQN.....G.YQTD FENLRESGLEK	445
NTDB id 1355 HI 0298 AAC21962.1	ANSCDCHQGYRGRIGVYQFLHWQQN.....D.YQTD FKNLRRASGLEK	445
NTDB id 1168 A1552VC RS11070 WP 000957200.1	LQSQFAFQPNIEI...LYEANAAGCNECTGYSGRVGIYEVMAFNTELAEATMQRASIHQIERLAKANGMQTLQESGLEK	545
NTDB id 1293 VP RS12240 WP 005479695.1	L.QHLGIQT T DN...IFRANPDGCNECTHGYSGRGTGIYEV MRFD E SLSEALIKGASVHELEKLA IANGMSTLQMSGIEK	544
NTDB id 1401 DSB67 RS12665 WP 010643259.1	L.QHIGIANTEQ...IFQANPDGCNECTHGYSGRGTGIYEV MKFD E SLSEALIKGASVHELEKLA IANGMSTLQMSGIEK	544
NTDB id 1251 GC085 RS07725 WP 011946523.1	GLIELGFKEADLV..NLKLYKAVGCEQCTSGYRGRVGLFEVLPMTKELGQLIMSGGNSLDILKLAQSEGMLTIFQSGIEK	557
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SLLEMGFTEQDLAHPDFRVFQPVGCLECREGYKGRVGIYEV MKVTPEISKIIMEDGNALEIAAAASEKLGFN NLRRLRRSGLKK	553
NTDB id 1018 ACIAD RS01685 WP 004920473.1	SLLEMGFTEPDLAQPEFQIFEPVGCHDCREGYKGRVGIYEV MKITPEISKIIMEDGNALEIAAATAETLGFNNLRRLRRSGLKK	553
consensus	*** * ** *! !!!!!!!***** * * ** * * ** * * ** * * ** * * ** * *	

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	IEQQLTNETEVDRLVGYE.	461
NTDB id 396363 GDY38 RS06070 WP 016504529.1	VRLGMTDDNEVQRVLGE..	461
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NTDB id 1251 GC085 RS07725 WP 011946523.1	VKEGITTIEEVNRVTVD..	574
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	VMQGV TSLQEINRV TSE..	570
consensus	*****! * !!!!! * *	

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