

logo

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NTDB id 1125013 EQB74 RS02750 WP 002287328.1	MADDRKAALDAALKKIEKSYGKCSIMKLCEK.IDQQISTIPSLSALDVALGVGGYPRGR	59
NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERQAALDAALKKIEKNFGKCSIMRMCEK.VDTQVSTVSSGSLALDEALGVGGYPRGR	60
NTDB id 600 KW2 RS01765 WP 011675413.1	MATK.....KKTNFDDITKKYGAERDKALADALALIEKDFGKCSIMRLCEA.ANQKVSVSSGSLALDTALCAGGYPRGR	74
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NTDB id 1334 RA0C RS04870 WP 004918236.1	MAKT..ETTSPPNDKKKALAALVLEKLDKTYGKCTVMKLCGDTVHDNIEVIPSLSGLGLDLALGVGGYPKGR	68
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	MDENKQKALAAALGQIEKQFGKCSIMRLCDN.RAMDVETISTGSLSLDLALCAGGLPMGR	59
NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDENKSKALQAALSQIEKQFGKNTVMRLCDN.TVQAVEAVSTGSLTLDIALCIGGLPKGR	59
NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKSKALQAALSQIEKQFGKNTVMRLCDN.TVQAVEAVSTGSLTLDIALCIGGLPKGR	59
NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQFGKCAVMRMGDH.DRQAIPSTGSLGLDLALCIGGLPKGR	59
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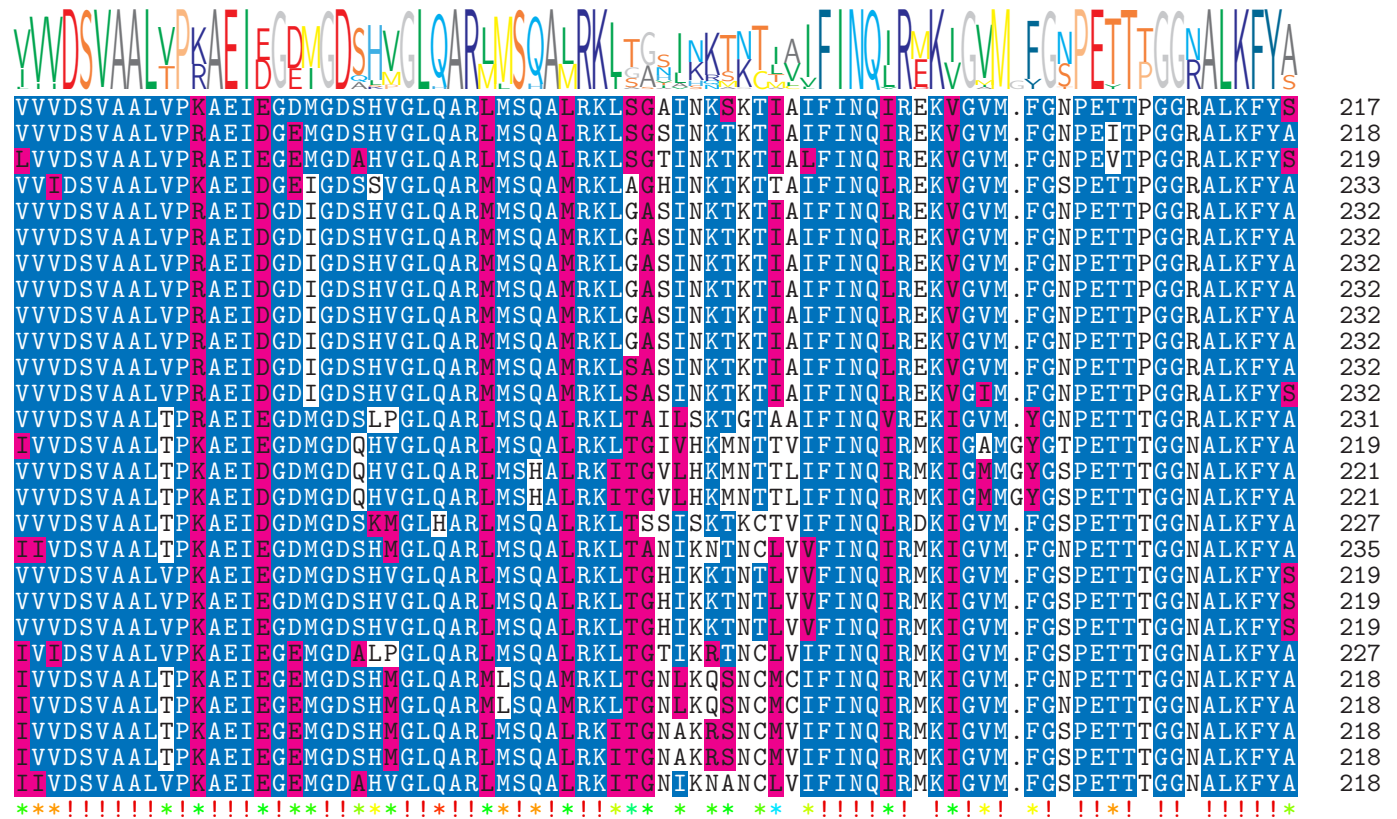
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	IVEIYGP	ESSGKTT	TLSLIAQA	QKGNKTC	AFIDAEHALDP	VYARNLGVM	IDELL	SQPD	NGEQ	ALEI	CDAL	VR	SGAVD	V	156	
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consensus	***!***	!!!!*!!!!*	!***!	!* **	!***!!!!*!* *	!!	!!****!!**!!!!	!!!!*	!*! *	***!***!	*					

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consensus

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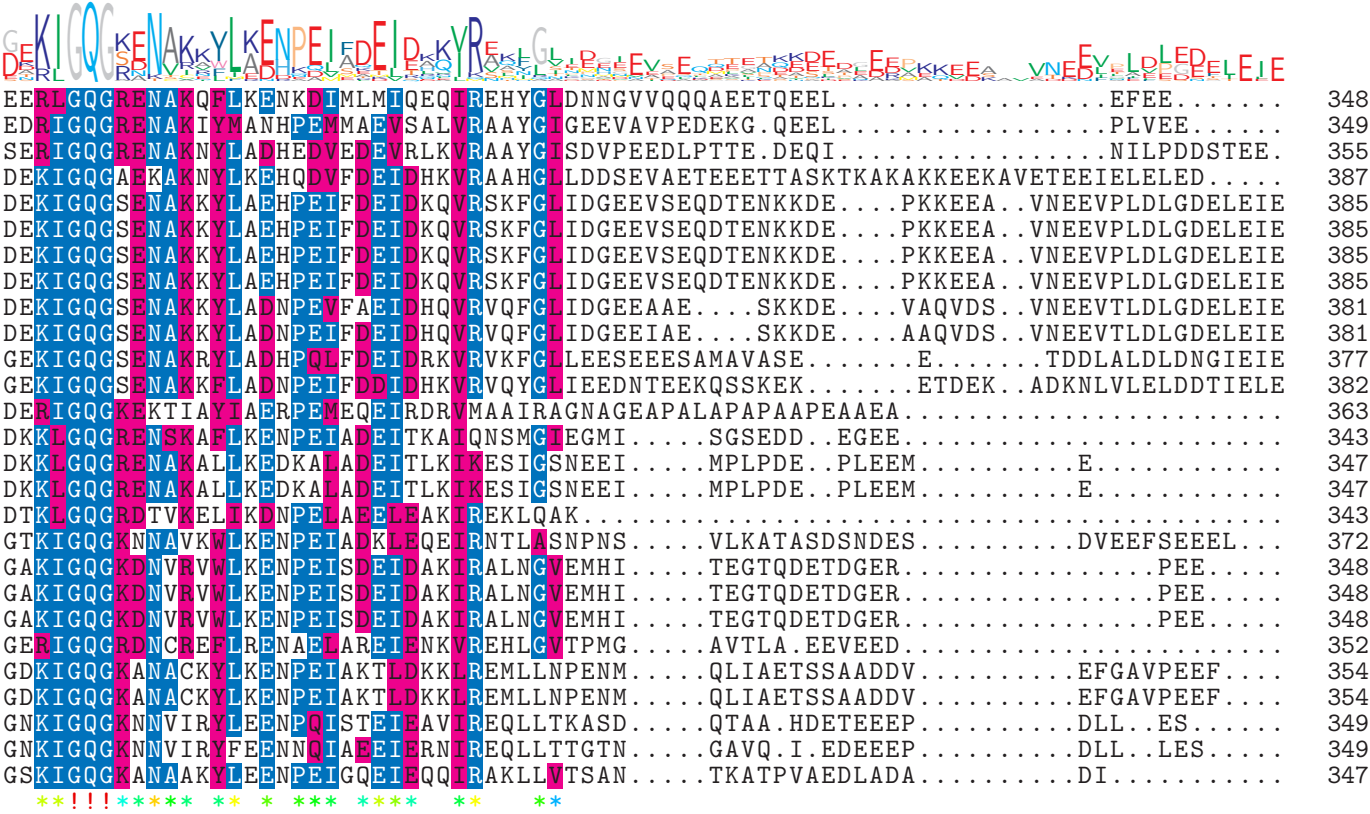


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NTDB id 499 SM12261 RS08315 WP 078228268.1	SVRLDVRGSTQIKGTG.DQKD	TNMGKETKIKVVKNKVAPPFKBAFVEIMYGEGISKTGELLKIASD	DI	IKKAGAWYSYK	311
NTDB id 453 Spy49 1753c ACI62003.1	SVRLDVRGTTQIKGTG.DQKD	SSICKETKIKVVKNKVAPPFKVAEVEIMYGEGISRTGELVKIASD	DI	IKKAGAWFSYN	311
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consensus	**!* **!* *	* * * * * ! * * ! ! ! * ! * ! * *	* * * * * ! * ! * * * * * * * * * * * * * ! ! * * * * !		

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consensus



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consensus			

- non conserved
- similar
- ≥ 50% conserved