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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 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKQKATSLATKQIDKVFVGKCALVRLCDK.QVEKIDISTGSLGLDLALCIGGVPRGR	61
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKSKALQAALSQIEKQFGKNTVMRLCDN.TVQAVEAVSTGSLTLDIALCIGGLPKGR	59
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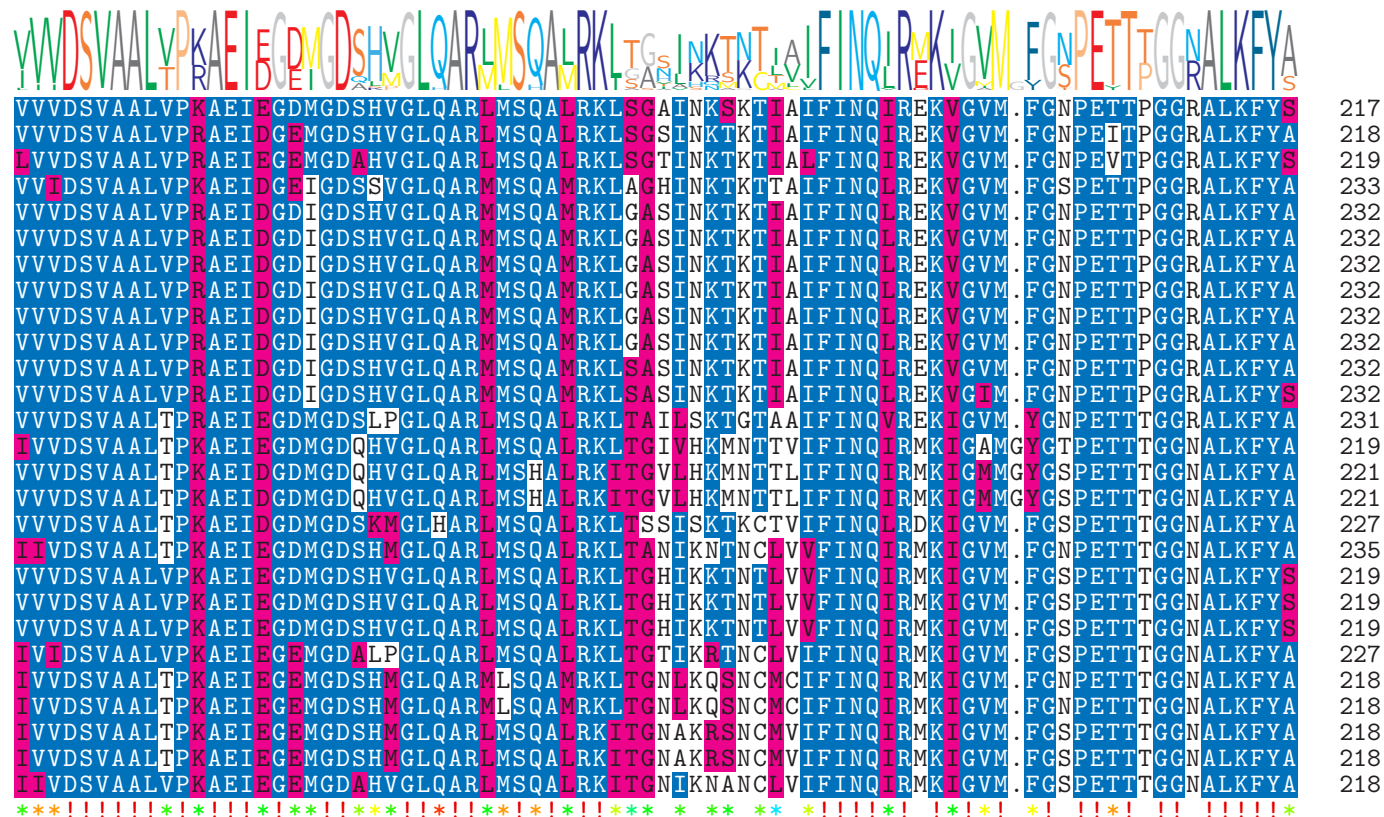
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NTDB id 228 SPD RS09265 WP 001085462.1	IIEIYGP	ESSGKTT	VALHATAQA	QKEGGIA	AFIDAE	HALDPAYAAAL	LGVM	IDELL	LSQPD	SGEQ	GLEI	AGKL	ID	SGAVD	L	153
NTDB id 194 SPR RS08825 WP 001085462.1	IIEIYGP	ESSGKTT	VALHATAQA	QKEGGIA	AFIDAE	HALDPAYAAAL	LGVM	IDELL	LSQPD	SGEQ	GLEI	AGKL	ID	SGAVD	L	153
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	IVEIYGP	ESSGKTT	TLSLIAQA	QKGNKT	CAFIDAE	HALDPVYARN	LGVM	IDELL	VSQPDN	GEQ	ALEI	CDAL	VRSGA	VDV		156
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consensus	***!***	!!!!*!!!!*	!***!	!* **	!***!!!!*!* *	!!	!!****	!* *!!!!	!!!!*!*!	* **	!***!	*				

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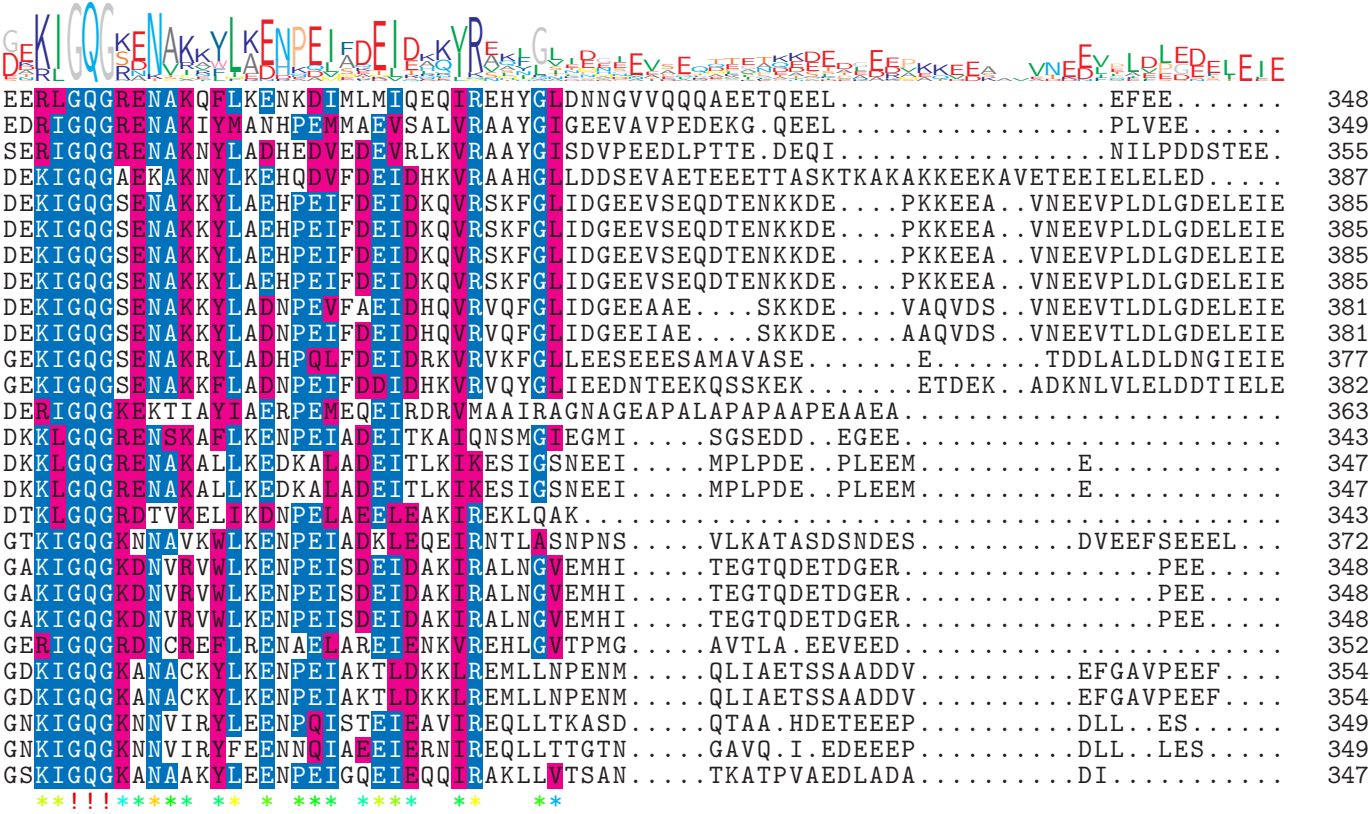


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consensus	**!* **!* *	* * * * * ! * * ! ! ! * ! * ! * *	* * * * * ! * ! * * * * * * * * * * ! ! * * * * !		

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consensus



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consensus			

- non conserved
- similar
- ≥ 50% conserved