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NTDB id 1125602 FSA55 RS09410 WP 002287328.1	MADDRKAALDAALKKIEKSYGKCSIMKLCEK.IDQQISTIPSGSLALDVALGVGGYPRGR	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 1211 C694 RS00795 WP 000963128.1	MAIDEDKQKATSLATKQIDKVFVGKCALVRLCDK.QVEKIDISTGSLGLDLALCIGGVPRGR	61
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NTDB id 1334 RA0C RS04870 WP 004918236.1	MAKT..ETTSPPNDKKKALAALVLEKLDKTYGKCTVMKLCGDTVHDNIEVIPSLSLGLDLALGVGGYPKGR	68
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NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....K..KAASMSAEKQKALAAALAQIEKQFGKCSIMKMGDA.EVEPVQVVSTGSLGLDVALGVGGLPGR	68
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKSKALQAALSQIEKQFGKNTVMRLCDN.TVQAVEAVSTGSLTLDIALCIGGLPKGR	59
NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQFGKGAVMRMGDH.DRQAIPSTGSLGLDLALCIGGLPKGR	59
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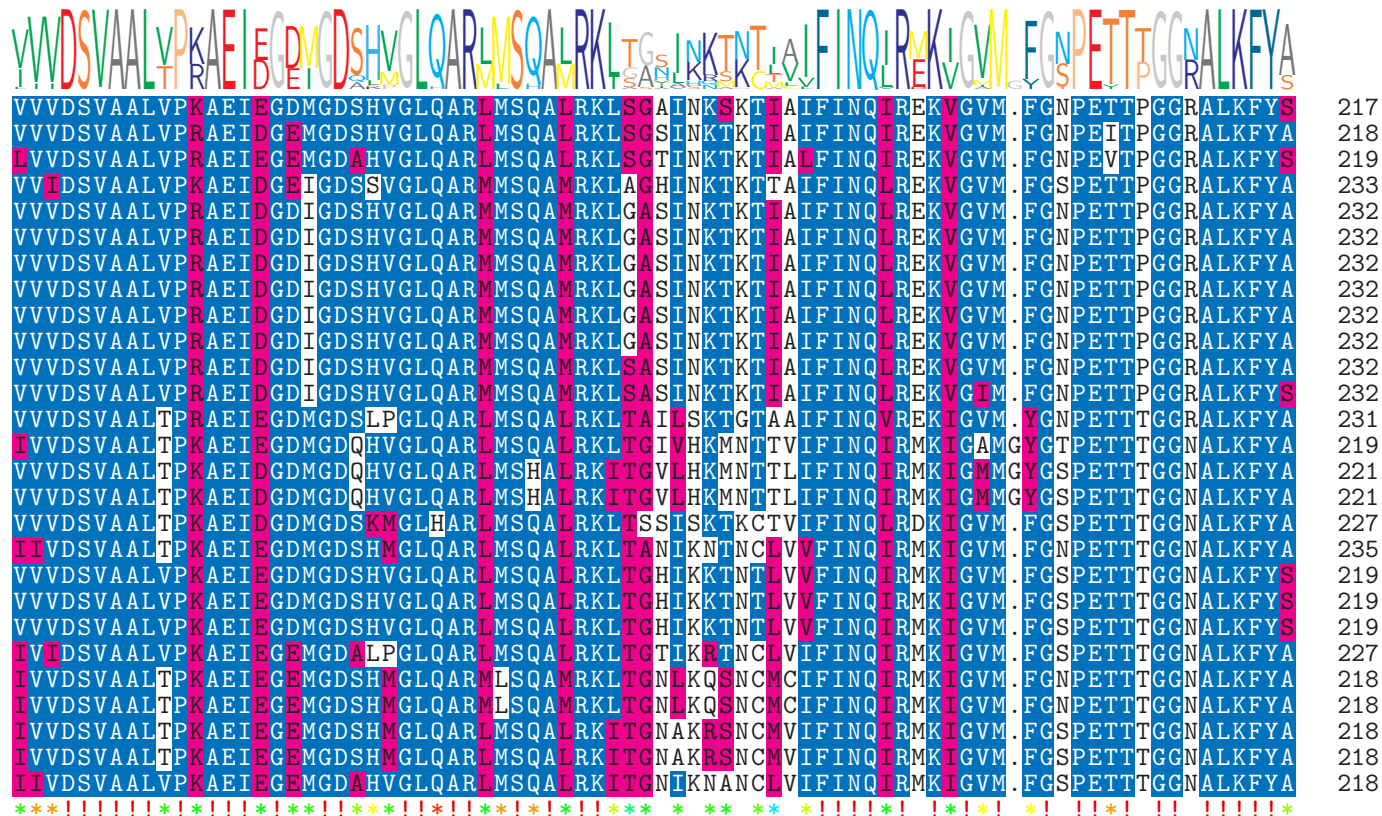
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NTDB id 625 LCA RS02525 WP 011374196.1	IVEIYGP	ESSGKTT	VALHATAEV	QKGGTA	YIDAEN	MDPKY	ATALGVN	IDDL	LLSQPD	TGEQ	GLEI	ADAL	VSSGAVD	I	140			
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NTDB id 1334 RA0C RS04870 WP 004918236.1	IVEIYGP	ESSGKTT	TLHATAEA	QKVGGT	AAFI	DAEHALD	FMHYAEK	LCIN	LEEL	IV	SQPD	N	GEQ	LEI	ADSL	TRSGG	VDI	148
NTDB id 1389 A4U84 RS07560 WP 021112712.1	IVEIYGP	ESSGKTT	TLSLIIAQA	QKGNKT	CAFI	DAEHALDP	VYARNLGVN	IDELL	SQPD	N	GEQ	LEI	CDAL	VRSGAVD	V	156		
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consensus	***!***	!!!!*!!!!*	!!	***!	!* **	!***!!!!*!* *	!!	!!*	*****!	***!!!!	!!!!*	!*!	* **	!***!	*			

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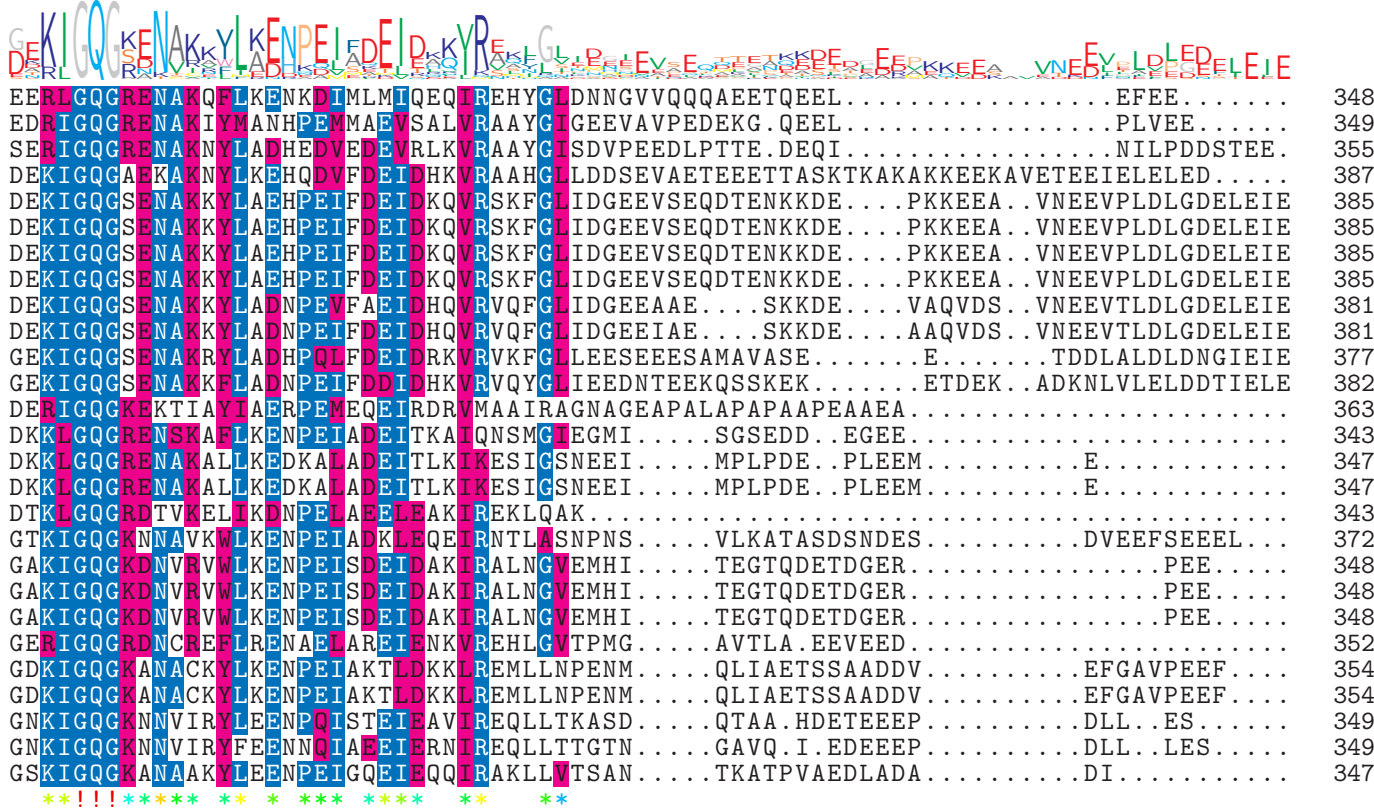


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consensus	**!* **!* *	* * * * * ! * * ! ! ! * ! * ! ! *	* * * * * ! * ! * * * * * * * * * * * * * * ! ! * * * * !		

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consensus



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consensus			

- non conserved
- similar
- ≥ 50% conserved