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NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK....KTNFDDITKKYGAERDKALADALALIEKD	FGKGSIMRLGERAE.AAN.QKVS	VSSGSLALDI	ALGAGGYPKGR	74
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NTDB id 625 LCA RS02525 WP 011374196.1MAKDERQAALDAALKKIEKN	FGKGSIMRMGEKVD.TQVSTVSSGSLALDEALGVGGYPRGR			60
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NTDB id 1113618 LFA RS08725 WP 045095853.1MEENKQKALAAALSQIERQ	FGKGSVMRMG	DSTVSRDIEAISTGSLGLDI	ALGIGGLPKGR	60
NTDB id 1205 PSJM300 05545 AFN77183.1MDENKKRALAAALGQIEKQ	FGKGVMLGERAE.QAIP	SISTGSLGLDI	ALGIGGLPKGR	59
consensus	*** ** *	*****	*****	*****	*****

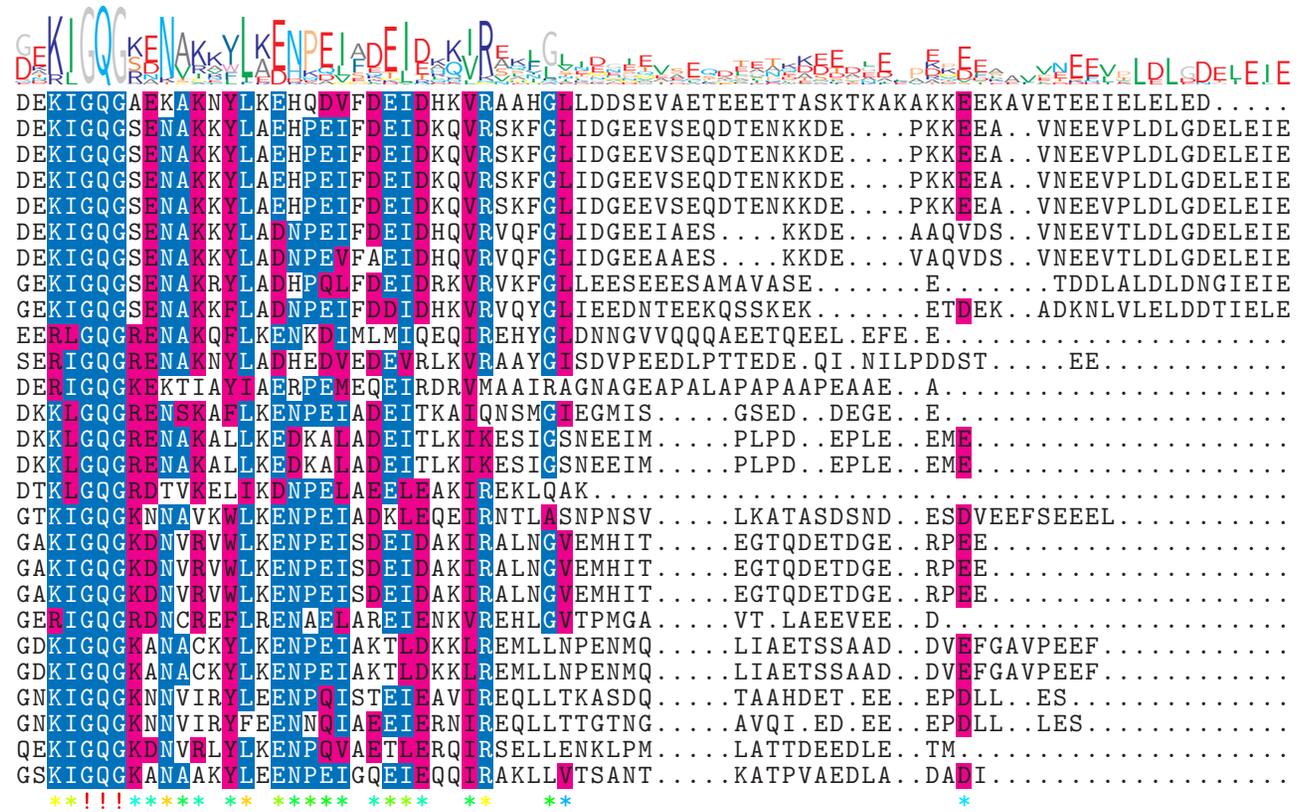
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NTDB id 269 KZH43 RS08665 WP 001085462.1	V	V	V	D	S	V	A	A	L	V	P	R	A	E	I	D	G	I	G	D	S	H	V	G	L	Q	A	R	M	S	Q	A	M	R	K	L	G	A	S	I	N	K	T	K	T	I	A	I	F	I	N	Q	L	R	E	K	V	G	V	M	.	F	G	N	P	E	T	T	P	G	G	R	A	L	K	F	Y	A	232		
NTDB id 228 SPD RS09265 WP 001085462.1	V	V	V	D	S	V	A	A	L	V	P	R	A	E	I	D	G	I	G	D	S	H	V	G	L	Q	A	R	M	S	Q	A	M	R	K	L	G	A	S	I	N	K	T	K	T	I	A	I	F	I	N	Q	L	R	E	K	V	G	V	M	.	F	G	N	P	E	T	T	P	G	G	R	A	L	K	F	Y	A	232		
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consensus	*	*	*	*	!	!	!	!																																																																									

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NTDB id 600 KW2 RS01765 WP 011675413.1
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 NTDB id 228 SPD RS09265 WP 001085462.1
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NTDB id 159 SP RS09750 WP 001085462.1	IEE	388
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NTDB id 1211 C694 RS00795 WP 000963128.1	...	347
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	...	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	...	348
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consensus

 non conserved
 similar
 ≥ 50% conserved