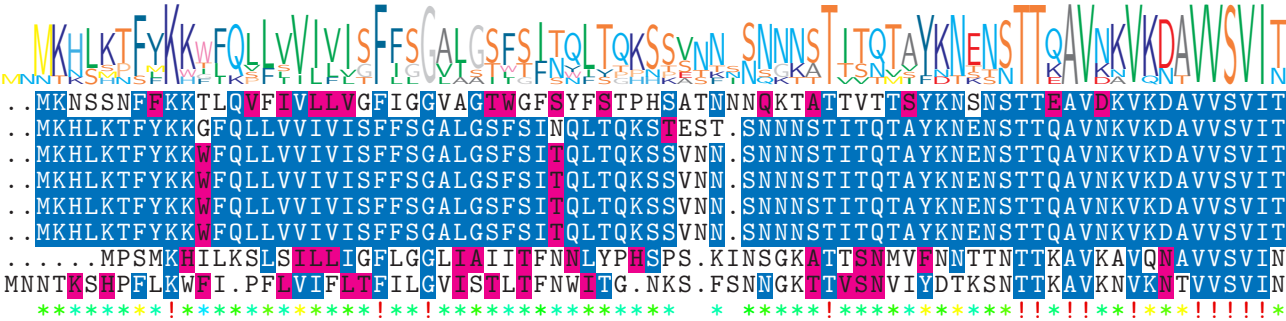


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NTDB id 539 SM12261 RS09395 WP 000681587.1
NTDB id 256 KZH43 RS10270 WP 000681597.1
NTDB id 215 SPD RS10945 WP 000681597.1
NTDB id 181 SPR RS10425 WP 000681597.1
NTDB id 145 SP RS11450 WP 000681597.1
NTDB id 323155 EB816 RS10090 WP 002994926.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

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consensus



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logo

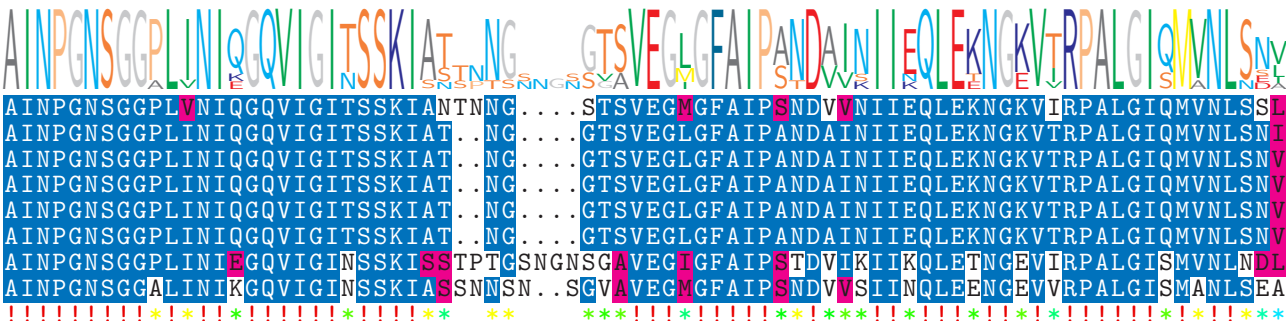
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consensus



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logo

NTDB id 425 SGO RS10525 WP 012131081.1
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consensus



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logo

NTDB id 425 SGO RS10525 WP 012131081.1	SSSSSDRLKLPD	NVKN	GVVVR	STQT	TGMPADG	KLEKYDVITK	IDDTEISSASDI	QSALYKHSIN	EEIKV	TYYRD	DGKEQ	TTT	384	
NTDB id 539 SM12261 RS09395 WP 000681587.1	NTSDIRRLNIP	SNVTSGV	VRSVQSN	MPANGHLE	KYDVITK	VDDKEIAS	STDLSALYNH	SIGDTIKI	TYRNGKE	ETTS			381	
NTDB id 256 KZH43 RS10270 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	VRSVQSN	MPANGHLE	KYDVITK	VDDKEIAS	STDLSALYNH	SIGDTIKI	TYRNGKE	ETTS			381	
NTDB id 215 SPD RS10945 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	VRSVQSN	MPANGHLE	KYDVITK	VDDKEIAS	STDLSALYNH	SIGDTIKI	TYRNGKE	ETTS			381	
NTDB id 181 SPR RS10425 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	VRSVQSN	MPANGHLE	KYDVITK	VDDKEIAS	STDLSALYNH	SIGDTIKI	TYRNGKE	ETTS			381	
NTDB id 145 SP RS11450 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	VRSVQSN	MPANGHLE	KYDVITK	VDDKEIAS	STDLSALYNH	SIGDTIKI	TYRNGKE	ETTS			381	
NTDB id 323155 EB816 RS10090 WP 002994926.1	STNAL	LSQINIP	TSVTGG	IVVAE	VKEGMP	ASGKLA	QYDVITE	IDGKT	VNSIS	SDLQSS	LYGHD	INDTIK	VT	393
NTDB id 381 SMU RS09890 WP 002262650.1	STSGRDT	LKIP	SDVTSG	IVVLST	QSGMP	ADGKL	KYDVITE	IDGKK	VASIS	DLQSILY	KHKKG	DKIKL	TFYREK	390
consensus	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	

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NTDB id 425 SGO RS10525 WP 012131081.1	IKLTK	ST	DDL	SSE.	397
NTDB id 539 SM12261 RS09395 WP 000681587.1	IKL	DKSSGD	LES..		393
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NTDB id 181 SPR RS10425 WP 000681597.1	IKLN	KSSGD	LES..		393
NTDB id 145 SP RS11450 WP 000681597.1	IKLN	KSSGD	LES..		393
NTDB id 323155 EB816 RS10090 WP 002994926.1	IKLTK	TTQDL	TKTQ		407
NTDB id 381 SMU RS09890 WP 002262650.1	IQLTK	TSQDL	NH..		402
consensus	!*!	!***!	!*		

- ⓧ

 non conserved
- ✖

 similar
- ⓧ

 ≥ 50% conserved