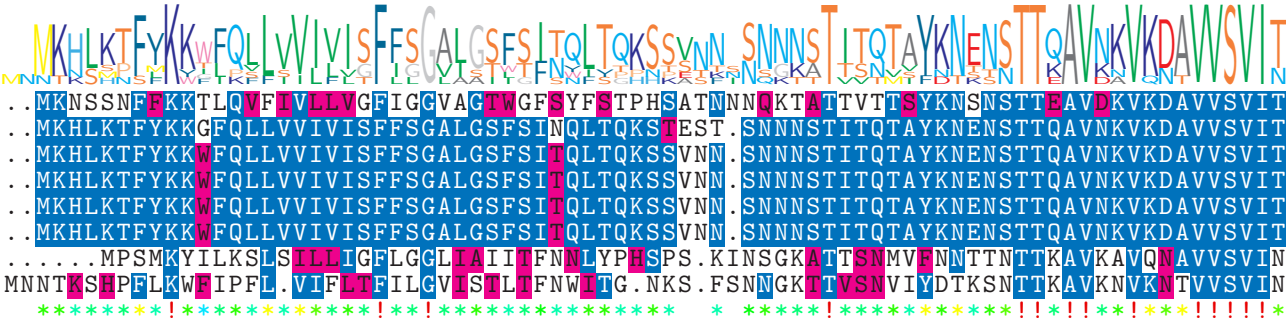


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NTDB id 539 SM12261 RS09395 WP 000681587.1
NTDB id 256 KZH43 RS10270 WP 000681597.1
NTDB id 215 SPD RS10945 WP 000681597.1
NTDB id 181 SPR RS10425 WP 000681597.1
NTDB id 145 SP RS11450 WP 000681597.1
NTDB id 308921 DZN89 RS09355 WP 011285340.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

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NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

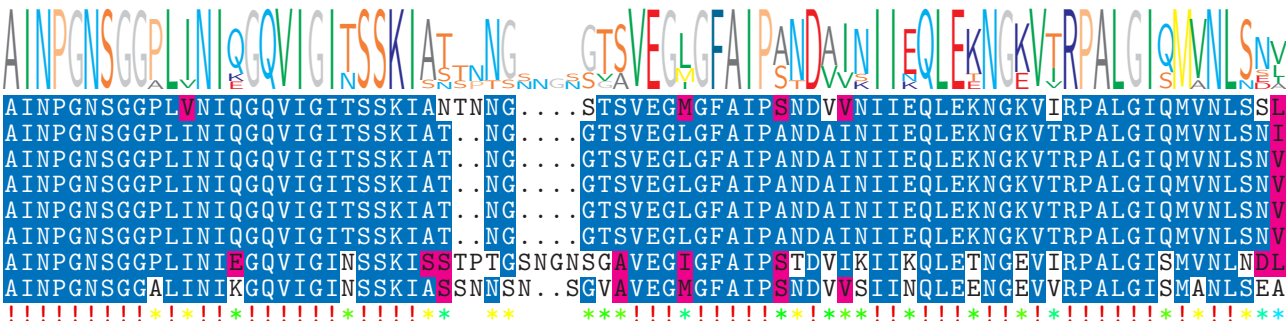
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NTDB id 181 SPR RS10425 WP 000681597.1
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consensus



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logo

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consensus



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logo

NTDB id 425 SGO RS10525 WP 012131081.1	SSSSSDRLKLPD	NVKN	GVVVR	STQT	GM	PADG	KLEKYDVITKID	TEISSASDI	QSALYKHSIN	EEIKV	TYYR	DGKEQ	TTT	384			
NTDB id 539 SM12261 RS09395 WP 000681587.1	NTSDIRRLNIP	SNVTSGV	VRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS							381			
NTDB id 256 KZH43 RS10270 WP 000681597.1	STSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS							381			
NTDB id 215 SPD RS10945 WP 000681597.1	STSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS							381			
NTDB id 181 SPR RS10425 WP 000681597.1	STSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS							381			
NTDB id 145 SP RS11450 WP 000681597.1	STSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS							381			
NTDB id 308921 DZN89 RS09355 WP 011285340.1	STNA	LSQ	INIP	TSVTGG	IVVAEVKEGMPASG	KL	AQYDVITEIDGKT	VNSI	SDLQSS	LYGHD	IND	TIK	VTFYRG	TTKKK	AD	393	
NTDB id 381 SMU RS09890 WP 002262650.1	STSG	RDT	LKIP	SDVTSG	IVVLSTQSGMPADG	KL	KYDVITEIDGKK	VASIS	DLQS	ILYKHK	KGDK	IKL	TFYRE	KDK	Q	TVE	390
consensus	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	

logo

NTDB id 425 SGO RS10525 WP 012131081.1	IKLTKSTDDL	SSE.	397
NTDB id 539 SM12261 RS09395 WP 000681587.1	IKL	DKSSGDLES..	393
NTDB id 256 KZH43 RS10270 WP 000681597.1	IKLN	KSSGDLES..	393
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NTDB id 145 SP RS11450 WP 000681597.1	IKLN	KSSGDLES..	393
NTDB id 308921 DZN89 RS09355 WP 011285340.1	IKLTK	TTQDLTKTQ	407
NTDB id 381 SMU RS09890 WP 002262650.1	IQLTK	TSQDLNH..	402
consensus	!*!	!***!*	

- ⓧ

 non conserved
- ✖

 similar
- ⓧ

 ≥ 50% conserved