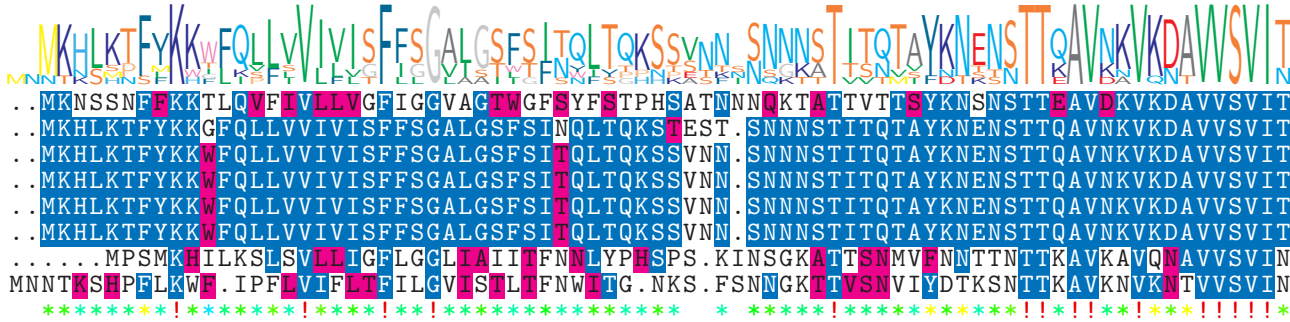


logo

NTDB id 425 SGO RS10525 WP 012131081.1
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 1136332 DQL31 RS10050 WP 002981947.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

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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 1136332 DQL31 RS10050 WP 002981947.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

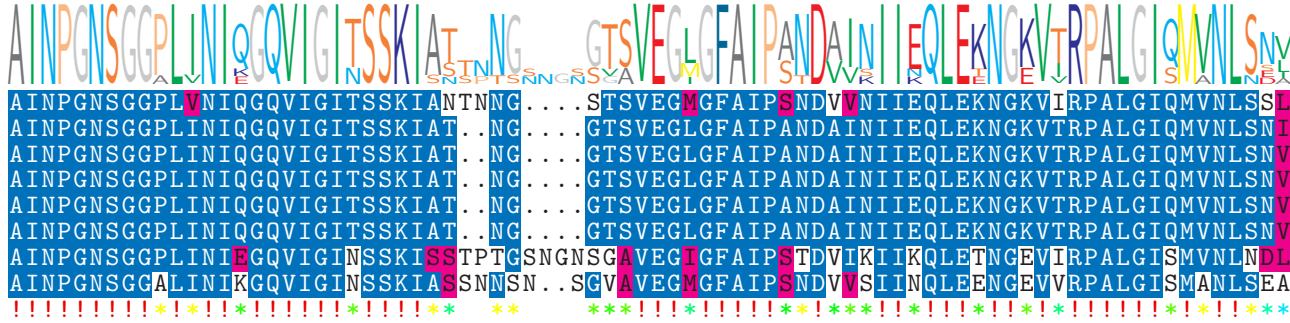
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NTDB id 181 SPR RS10425 WP 000681597.1
NTDB id 145 SP RS11450 WP 000681597.1
NTDB id 1136332 DQL31 RS10050 WP 002981947.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

NTDB id 425 SGO RS10525 WP 012131081.1
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NTDB id 181 SPR RS10425 WP 000681597.1
NTDB id 145 SP RS11450 WP 000681597.1
NTDB id 1136332 DQL31 RS10050 WP 002981947.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

NTDB id 425 SGO RS10525 WP 012131081.1	SSSSSDRLKLPD	NNVKN	GVVVR	STQT	IGMPAD	CKLEKYDVITK	DDTEISSASDI	QSALYKHSIN	EEIKV	TYYRD	GKEQT	TTT	384
NTDB id 539 SM12261 RS09395 WP 000681587.1	NTSDIRRLNIP	SNVTSGV	VRSVQSNMP	PANGHLEKYDVITKVDDKEIASSTDLQSALYNH	SIGDTIKITYYRNGKEETS								381
NTDB id 256 KZH43 RS10270 WP 000681597.1	STSDIRRLNIP	SNVTSGV	VRSVQSNMP	PANGHLEKYDVITKVDDKEIASSTDLQSALYNH	SIGDTIKITYYRNGKEETS								381
NTDB id 215 SPD RS10945 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	VRSVQSNMP	PANGHLEKYDVITKVDDKEIASSTDLQSALYNH	SIGDTIKITYYRNGKEETS								381
NTDB id 181 SPR RS10425 WP 000681597.1	STSDIRRLNIP	SNVTSGV	VRSVQSNMP	PANGHLEKYDVITKVDDKEIASSTDLQSALYNH	SIGDTIKITYYRNGKEETS								381
NTDB id 145 SP RS11450 WP 000681597.1	STSDIRRLNIP	SNVTSGV	VRSVQSNMP	PANGHLEKYDVITKVDDKEIASSTDLQSALYNH	SIGDTIKITYYRNGKEETS								381
NTDB id 1136332 DQL31 RS10050 WP 002981947.1	STNALSQINIP	TSVTGGIV	VAEVKEGMPAS	GKLAQYDVITE	DGKIVNST	SDLQSSLYGHD	INDTIKV	TFYRG	TTKKK	KAD			393
NTDB id 381 SMU RS09890 WP 002262650.1	STSGRDTL	LKIPSDVTSGIV	VLSTQSGMPAD	CKLKKYDVITE	DGKKVASI	SDLQSTLYKHK	GDKIKL	TFYRE	KDK	QIVE			390
consensus	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	

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NTDB id 425 SGO RS10525 WP 012131081.1	IKLTKSTDDLSSE.	397
NTDB id 539 SM12261 RS09395 WP 000681587.1	IKLTKSSGDLES..	393
NTDB id 256 KZH43 RS10270 WP 000681597.1	IKLTKSSGDLES..	393
NTDB id 215 SPD RS10945 WP 000681597.1	IKLTKSSGDLES..	393
NTDB id 181 SPR RS10425 WP 000681597.1	IKLTKSSGDLES..	393
NTDB id 145 SP RS11450 WP 000681597.1	IKLTKSSGDLES..	393
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NTDB id 381 SMU RS09890 WP 002262650.1	IKLTKTSQDLNH..	402
consensus	!*! !***! !**	

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved