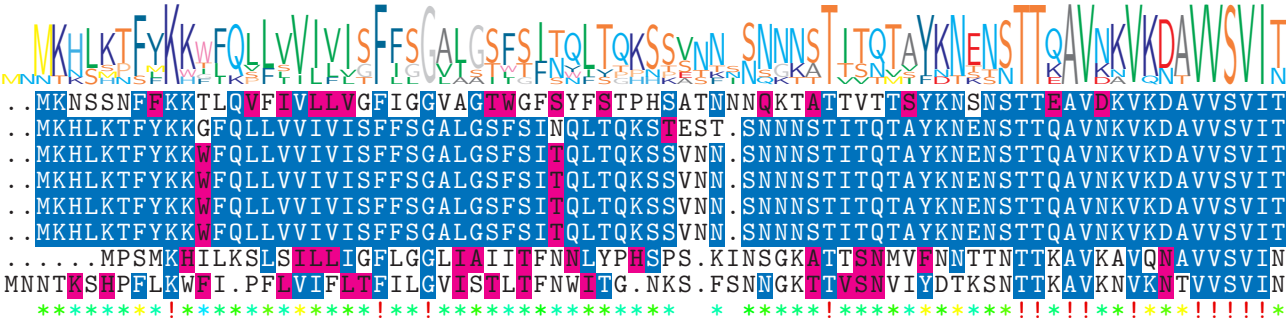


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NTDB id 539 SM12261 RS09395 WP 000681587.1
NTDB id 256 KZH43 RS10270 WP 000681597.1
NTDB id 215 SPD RS10945 WP 000681597.1
NTDB id 181 SPR RS10425 WP 000681597.1
NTDB id 145 SP RS11450 WP 000681597.1
NTDB id 281614 C5P47 RS09065 WP 002994926.1
NTDB 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
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NTDB id 281614 C5P47 RS09065 WP 002994926.1
NTDB id 381 SMU RS09890 WP 002262650.1
consensus



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logo

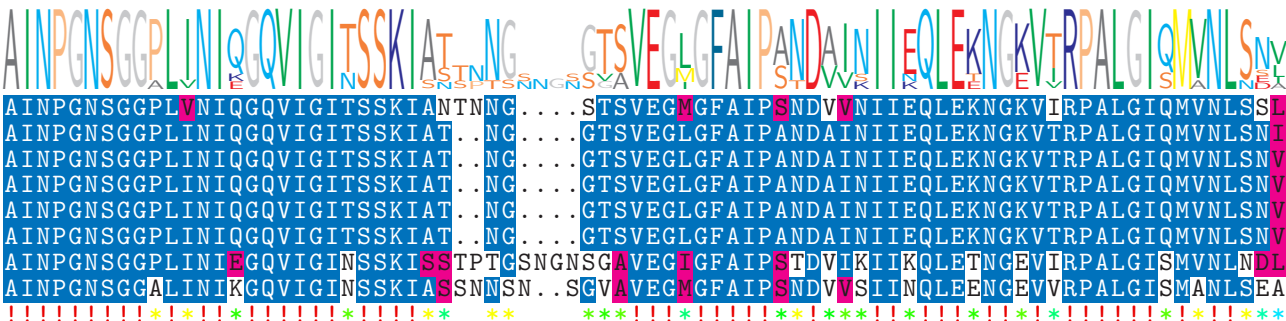
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consensus



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logo

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consensus



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logo

NTDB id 425 SGO RS10525 WP 012131081.1	SSSSSSDRLKLPD	NTDKVKN	GVVVR	STQT	IGMPADGKLEKYDVITKID	TEISSASDI	QSALYKHSIN	EEIKV	TYYRD	GKEQTTT	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	NTSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS				381
NTDB id 215 SPD RS10945 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS				381
NTDB id 181 SPR RS10425 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS				381
NTDB id 145 SP RS11450 WP 000681597.1	NTSDIRRLNIP	SNVTSGV	IVRSVQSNMP	PANGHLEKYDVITKVDDKEIAS	STDLQSALYNH	SIGDTIKITYYR	NGKEETTS				381
NTDB id 281614 C5P47 RS09065 WP 002994926.1	STNAL	LSQINIP	TSVTGGIV	VAEVKEGMPASG	KLAQYDVITEIDGKT	VNSIS	DLQSSLYGHD	INDTIK	VTFYR	GTTKKKAD	393
NTDB id 381 SMU RS09890 WP 002262650.1	STSGRDT	LKIPSD	VTS	GIVVLST	QSGMPADGKLK	KYDVITEIDGKKVASI	SDLQSILYKHKKGDKIKL	TFYREKDK	QTVE		390
consensus	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	

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NTDB id 425 SGO RS10525 WP 012131081.1	IKLTKSTDDL	SE.	397
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NTDB id 256 KZH43 RS10270 WP 000681597.1	IKLN	KSSGDLES..	393
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NTDB id 381 SMU RS09890 WP 002262650.1	IQLTKTSQDL	NH..	402
consensus	!*!	!*!!*	!*!

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved