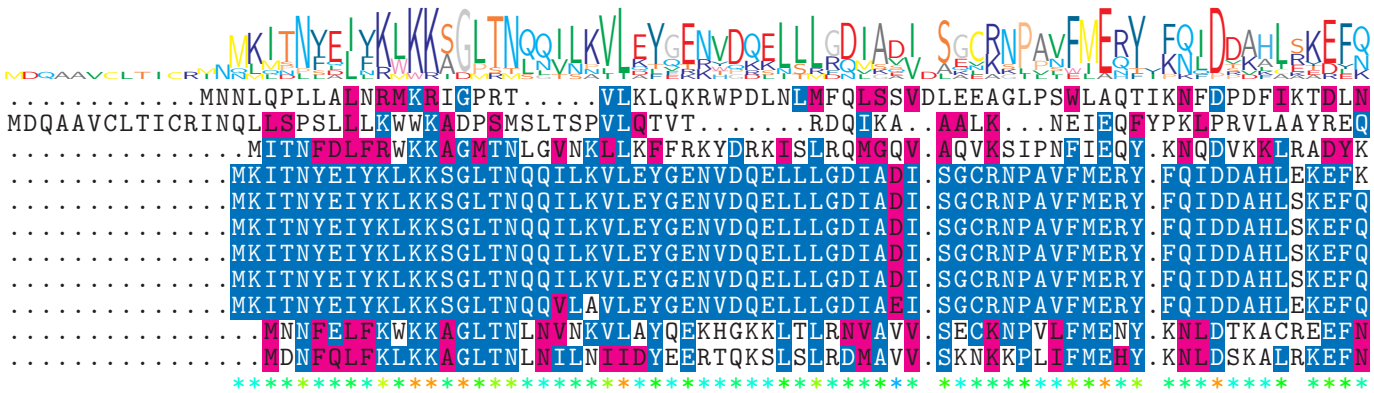


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NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 49834 SSUA7 RS04315 WP 014638936.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus



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logo

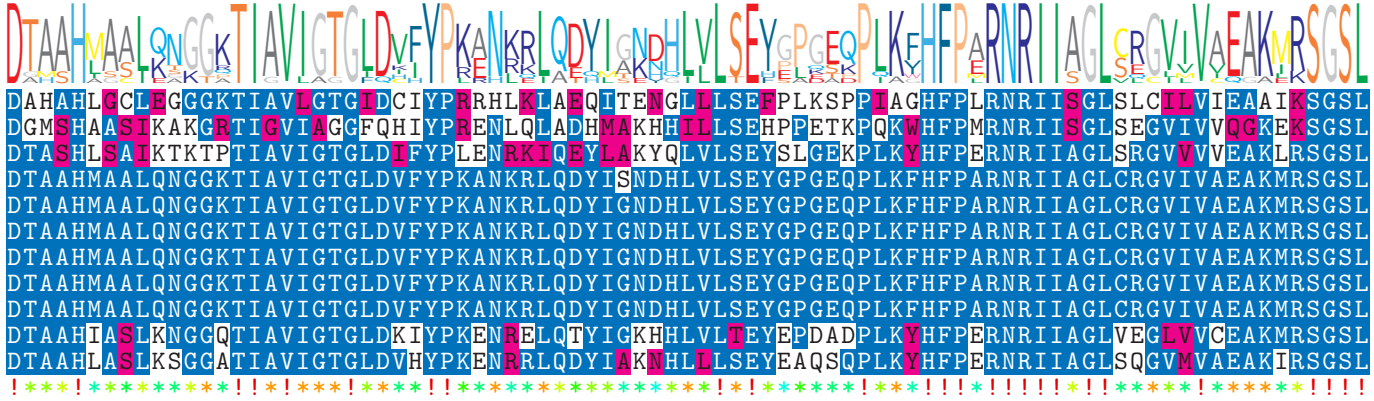
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NTDB id 497 SM12261 RS05565 WP 000705318.1
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NTDB id 407 SMU RS04605 WP 002262865.1
consensus



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logo

NTDB id 1250 GC085 RS13065 WP 027219989.1
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NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 49834 SSUA7 RS04315 WP 014638936.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus



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logo

NTDB id 1250 GC085 RS13065 WP 027219989.1	ITARMAL	EQNRDVL	AIPGSI	HNPL	ARGCHYL	LQQGAKL	VTSV	ADVL	DELKIEHH	QFTSN	KPIFSL	ASGKEN	LVKFI	G	FETTT	IDQ	322
NTDB id 114 BSU 16110 NP 389493.1	ITAYQAL	EQGRDVF	AVPGSL	FD	PYAGGPI	KLIQQGAK	ATWSA	ED	I	FEEL	PERNV	QYTE	P	F			297
NTDB id 599 KW2 RS05940 WP 021037268.1	ITCERALE	EGRD	IFAIPG	NIADGT	SDGCNHL	IQGAKL	VYQAQ	DILEE	YLYN								282
NTDB id 525 SMSK321 RS06295 WP 000705298.1	ITCERAMEE	GRDVF	AIPGSI	LDGLSD	GCHHLI	QEGAKL	VTSGQ	QDVL	AEFEF								282
NTDB id 267 KZH43 RS05590 WP 000705306.1	ITCERAMEE	GRDVF	AIPGSI	LDGLSD	GCHHLI	QEGAKL	VTSGQ	QDVL	AEFEF								282
NTDB id 226 SPD RS05990 WP 000705306.1	ITCERAMEE	GRDVF	AIPGSI	LDGLSD	GCHHLI	QEGAKL	VTSGQ	QDVL	AEFEF								282
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NTDB id 157 SP RS06205 WP 000705306.1	ITCERAMEE	GRDVF	AIPGSI	LDGLSD	GCHHLI	QEGAKL	VTSGQ	QDVL	AEFEF								282
NTDB id 497 SM12261 RS05565 WP 000705318.1	ITCERAMEE	GRDVF	AIPGSI	LDGLSD	GCHHLI	QEGAKL	VTSGQ	QDVL	AEFEF								282
NTDB id 49834 SSUA7 RS04315 WP 014638936.1	ITCERALEE	GR	EVYAIP	GSIM	DGKSD	GCHHLI	QEGAKC	VTSG	L	D	I	I	S	E	Y	Q	280
NTDB id 407 SMU RS04605 WP 002262865.1	ITCERAMEE	GRDVF	VVPGN	ILDGQ	SEGCHHLI	QEGAKC	ITSG	F	D	I	L	N	E	F	N	F	280
consensus	!	!	!	!	!	!	!	!	!	!	!	!	!	!	!	!	

logo

NTDB id 1250 GC085 RS13065 WP 027219989.1	I	IDRSGY	GMEQ	VT	SGLA	ELEL	KGAV	I	AVPGGY	I	RCEYER	361
NTDB id 114 BSU 16110 NP 389493.1	I	IDRSGY	GMEQ	VT	SGLA	ELEL	KGAV	I	AVPGGY	I	RCEYER	297
NTDB id 599 KW2 RS05940 WP 021037268.1												282
NTDB id 525 SMSK321 RS06295 WP 000705298.1												282
NTDB id 267 KZH43 RS05590 WP 000705306.1												282
NTDB id 226 SPD RS05990 WP 000705306.1												282
NTDB id 192 SPR RS05715 WP 000705306.1												282
NTDB id 157 SP RS06205 WP 000705306.1												282
NTDB id 497 SM12261 RS05565 WP 000705318.1												282
NTDB id 49834 SSUA7 RS04315 WP 014638936.1												280
NTDB id 407 SMU RS04605 WP 002262865.1												280
consensus												

- non conserved
- similar
- ≥ 50% conserved