

logo

Accession	Sequence	Position
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NTDB id 241 SP RS04015 WP 000882523.1	MLCQNCNKINDST.....	12
NTDB id 58167 SPN034183 RS03955 WP 000882516.1	MLCQNCNKINDST.....	12
NTDB id 281 KZH43 RS03550 WP 000882517.1	MLCQNCNKINDST.....	12
NTDB id 245 SPD RS03860 WP 000882517.1	MLCQNCNKINDST.....	12
NTDB id 243 SPR RS03635 WP 000882517.1	MLCQNCNKINDST.....	12
NTDB id 378 SMU RS02690 WP 002263569.1	MLCQNCNKINEAT.....	12
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLIG.....R	70
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYPTPTLDRILEKAEYYAHQYQYGTIESAHLLEAMATTSGSIAYSLLAGMNVDDSSDLLIDL.....EDLSSH	72
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
NTDB id 377 SMU RS09275 WP 002262344.1	MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFEADVRFEDYQAAAAILAIGKKPKSSS	76
NTDB id 297 STER RS00545 WP 011680614.1	MTIYSRKMQAIFHRAQLEAERFESPFLFTWVLLAMVEVTGSAVYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
NTDB id 329 STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFLFTWVLLAMVEVPGSAVYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
consensus	*.....*	

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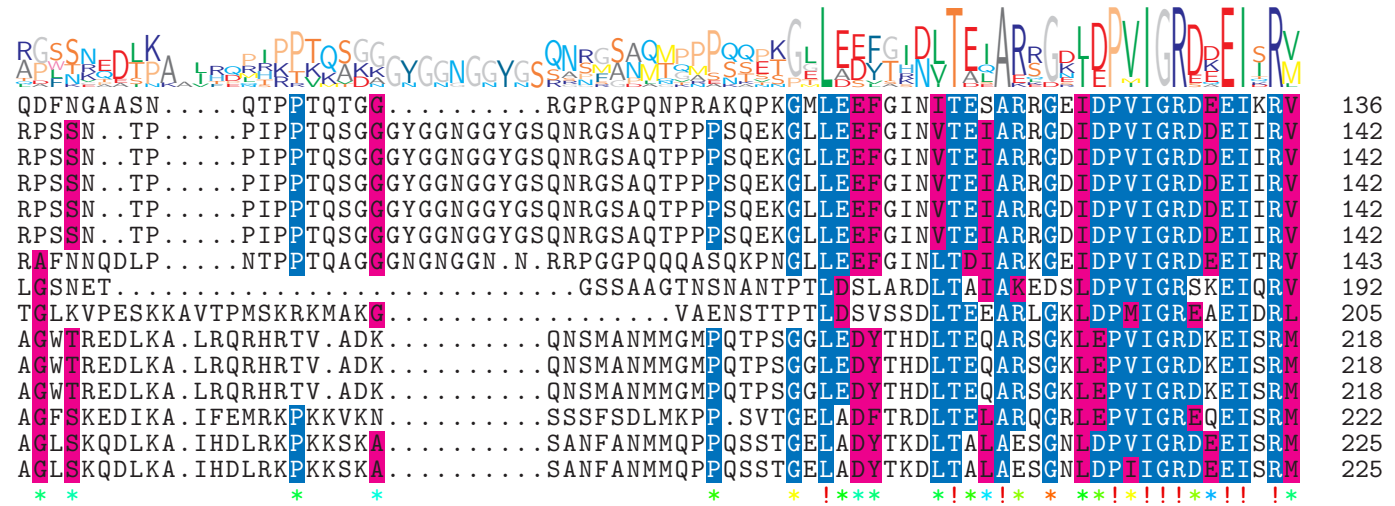
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NTDB id	245	SPD	RS03860	WP	000882517.1	IHL	YTNLNGKQKKQIDLC	QCNCYKT...	IKTDPNNSL	LFKGMTDLNNRD.....	FDP	FGDFFNDLNNF	69	
NTDB id	243	SPR	RS03635	WP	000882517.1	IHL	YTNLNGKQKKQIDLC	QCNCYKT...	IKTDPNNSL	LFKGMTDLNNRD.....	FDP	FGDFFNDLNNF	69	
NTDB id	378	SMU	RS02690	WP	002263569.1	IHL	YANVNGQKKQIDLC	QCNCYQT...	MKTDPE	NGPLNHLSQQNSSS.....	INP	FFDDFFGD LN NF	70	
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NTDB id	290	KZH43	RS10025	WP	001109677.1	QDETFT	ELPFSSRLQLIF	DEAEYVASVVHAKVLGT	EHVLYAILHD	GNALATRILERAGFSYEDKKDQVKIAALRRN	LEER		150	
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NTDB id	297	STER	RS00545	WP	011680614.1	ESDIIDL	RAGSPALEARMLQE	AQEIASVTGA	VEVGESEHVLM	MAFLHLHKDLMVCRLL	EVAGFQYKDSDKPRI	IDLRRSLERN	156	
NTDB id	329	STU	RS10020	WP	011225298.1	ESDIIDL	RAGSPALEARMLQE	AGIASVTGA	VEVGESEHVLM	AFLLHKDLMVCRLL	EVAGFQYKDSDKPRI	IDLRRSLERN	156	
consensus						*	*	*	*	*	*	*	*	*

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NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 241 SP RS04015 WP 000882523.1
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NTDB id 287 SP RS11210 WP 001109712.1
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NTDB id 329 STU RS10020 WP 011225298.1
consensus

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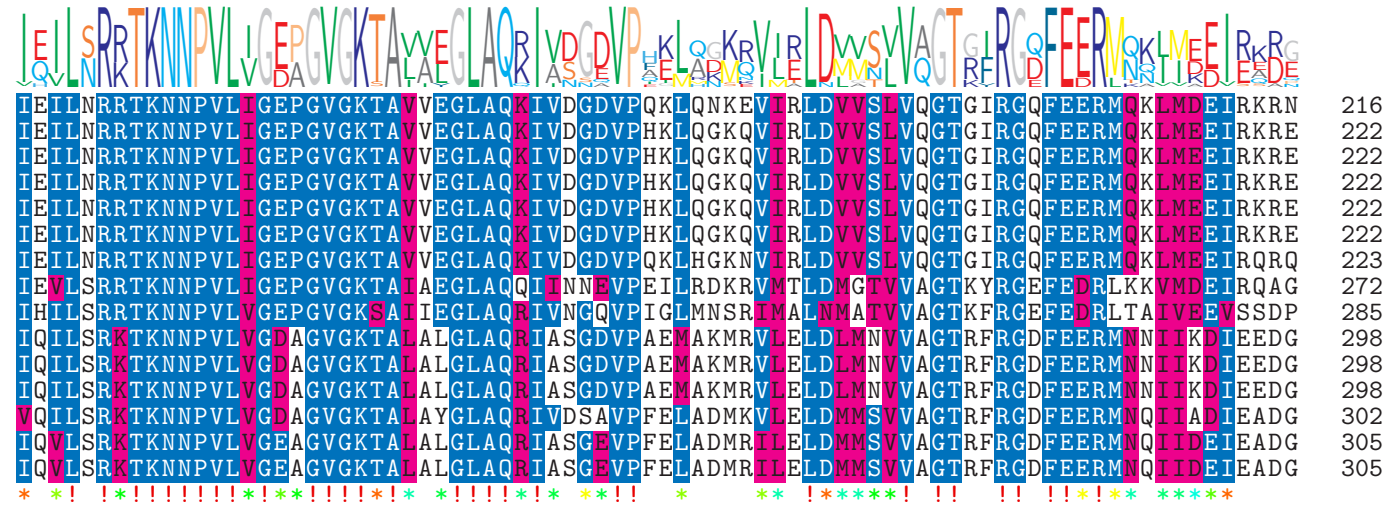


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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

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NTDB id 610 V4T04 RS10165 WP 012897346.1	IKLDIKASRTKEKAVEKIADKIYKFS.....VKEDKRQEVTDQAVAVASTLTGVPITQMTKES	SDRLINLEKELHKRVI	520	
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consensus	* * * * *	* * * * *		

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consensus							!!*!* *	* *!*!* *	! *	*	!!*																																																																									

logo

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NTDB id 601 KW2 RS02615 WP 021036805.1
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consensus

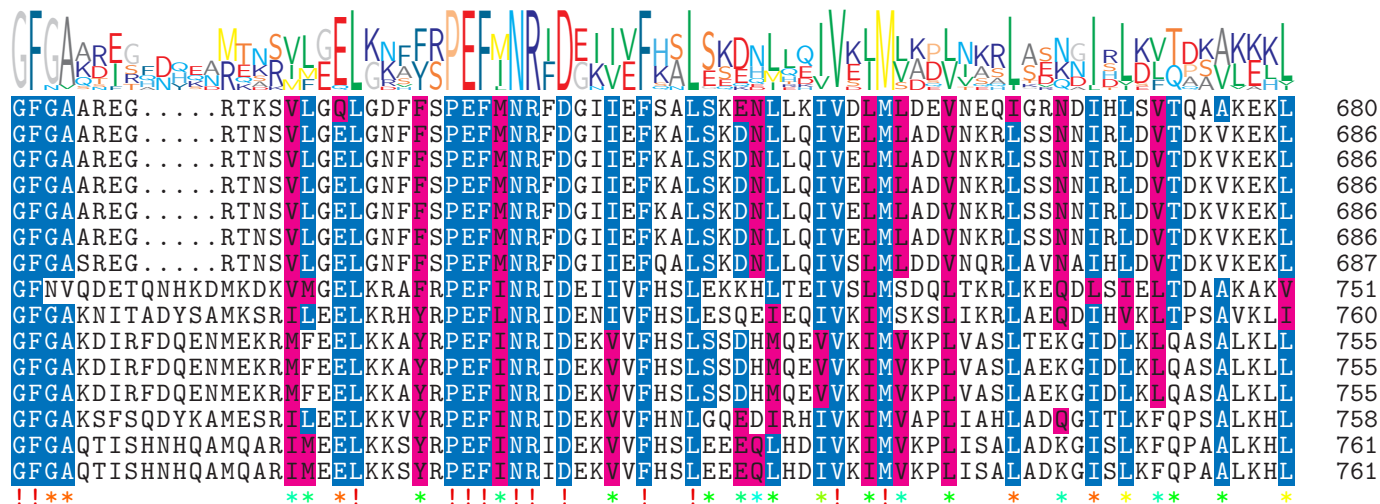
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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

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NTDB id	85	BSU	00860	NP	387967.1	AEEGVLDLEYGARPLRRAIQ	KHVEDRLSEELLRGNIHKGGHIVLDVEDGEFVVKTAKTN.....	810				
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consensus						* ! * * *	* ! ! ! ! ! * * * !	* ! ! * * *	*	** * * *	* * *	* * *

-  non conserved
-  similar
-  $\geq 50\%$ conserved