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NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 378 SMU RS02690 WP 002263569.1
NTDB id 241 SP RS04015 WP 000882523.1
NTDB id 281 KZH43 RS03550 WP 000882517.1
NTDB id 245 SPD RS03860 WP 000882517.1
NTDB id 243 SPR RS03635 WP 000882517.1
NTDB id 395308 GFH33 RS00525 WP 000971187.1
NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

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[illegible]

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

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[illegible]

logo

NTDB id 601 KW2 RS02615 WP 021036805.1	AGSSNDLKA	PPTQSGG	QNSGANQMPPOSTGL	EEFTDLTE	ARGGLDPVIGRD	EIRV	136
NTDB id 378 SMU RS02690 WP 002263569.1	DFNGAASNQ	TPPTQTG	RGPRGPQNPRAKQPK	MLEEFGIN	ITESARRGE	IDPVIGRDEE	IKRV
NTDB id 241 SP RS04015 WP 000882523.1	RAFNNQDLPN	TPPTQAG	GGNGNGGN.N	RRPGGPQQQASQ	KPNGLLEEFGIN	LTDIARKGE	IDPVIGRDEE
NTDB id 281 KZH43 RS03550 WP 000882517.1	RPSSN..TPP	IPPTQSG	GGYGGNGGYGS	QNRGSAQT	PPPSQEKGLLEEFGIN	VTEIARRGD	IDPVIGRDDE
NTDB id 245 SPD RS03860 WP 000882517.1	RPSSN..TPP	IPPTQSG	GGYGGNGGYGS	QNRGSAQT	PPPSQEKGLLEEFGIN	VTEIARRGD	IDPVIGRDDE
NTDB id 243 SPR RS03635 WP 000882517.1	RPSSN..TPP	IPPTQSG	GGYGGNGGYGS	QNRGSAQT	PPPSQEKGLLEEFGIN	VTEIARRGD	IDPVIGRDDE
NTDB id 395308 GFH33 RS00525 WP 000971187.1	ASSGHQGSST	NANTPT	LD	SLARDLT	VVARENR	LDPVIGRSKE	IQRV
NTDB id 85 BSU 00860 NP 387967.1	TGSSA	AGTNS	NANTPT	LD	SLARDLT	AIKEDSL	DPVIGRSKE
NTDB id 610 V4T04 RS10165 WP 012897346.1	TGLKVPESK	KAVTPMSKR	KMAK	VAENST	PTLDSV	SSDLTEEAR	LGLDPMIGRE
NTDB id 290 KZH43 RS10025 WP 001109677.1	AGWTRED	LKA.LRQRH	RTV.ADK	QNSMAN	MMGMPQT	PSGGLEDY	THDLTEQARS
NTDB id 289 SPD RS10700 WP 001109677.1	AGWTRED	LKA.LRQRH	RTV.ADK	QNSMAN	MMGMPQT	PSGGLEDY	THDLTEQARS
NTDB id 287 SP RS11210 WP 001109712.1	AGWTRED	LKA.LRQRH	RTV.ADK	QNSMAN	MMGMPQT	PSGGLEDY	THDLTEQARS
NTDB id 377 SMU RS09275 WP 002262344.1	AGFSKEDI	KA.IFEMR	KPKVK	NSSSF	SDLMKPP	SVT	TGELADFT
NTDB id 297 STER RS00545 WP 011680614.1	AGLSKQDL	KA.IHDLR	KPKKS	A	SANFAN	MMQPPQ	SS
NTDB id 329 STU RS10020 WP 011225298.1	AGLSKQDL	KA.IHDLR	KPKKS	A	SANFAN	MMQPPQ	SS
consensus	*	*	*	**	**	!***	**!***

logo

NTDB id 601 KW2 RS02615 WP 021036805.1	IEILNRRTKNNPVL	IGEPGVGKTA	VEGLAQK	IVDGDV	PQKLQNK	EVIRLDV	VS
NTDB id 378 SMU RS02690 WP 002263569.1	IEILNRRTKNNPVL	IGEPGVGKTA	VEGLAQK	IVDGDV	PQKLQNK	EVIRLDV	VS
NTDB id 241 SP RS04015 WP 000882523.1	IEILNRRTKNNPVL	IGEPGVGKTA	VEGLAQK	IVDGDV	PHKLQGK	QVIRLDV	VS
NTDB id 281 KZH43 RS03550 WP 000882517.1	IEILNRRTKNNPVL	IGEPGVGKTA	VEGLAQK	IVDGDV	PHKLQGK	QVIRLDV	VS
NTDB id 245 SPD RS03860 WP 000882517.1	IEILNRRTKNNPVL	IGEPGVGKTA	VEGLAQK	IVDGDV	PHKLQGK	QVIRLDV	VS
NTDB id 243 SPR RS03635 WP 000882517.1	IEILNRRTKNNPVL	IGEPGVGKTA	VEGLAQK	IVDGDV	PHKLQGK	QVIRLDV	VS
NTDB id 395308 GFH33 RS00525 WP 000971187.1	IEVLSRRTKNNPVL	IGEPGVGKTA	IEGLAQ	QIVNNE	VEPET	LRDKR	VM
NTDB id 85 BSU 00860 NP 387967.1	IEVLSRRTKNNPVL	IGEPGVGKTA	IEGLAQ	QIVNNE	VEPET	LRDKR	VM
NTDB id 610 V4T04 RS10165 WP 012897346.1	IHILSRRTKNNPVL	VGEPGVGK	SAIEGLAQ	RIVNGQ	VP	IGLMNS	R
NTDB id 290 KZH43 RS10025 WP 001109677.1	IQILSRKTKNNPVL	VGDAGVGKTA	LALGLAQ	RIASGD	VPAEMAK	MRVLE	EL
NTDB id 289 SPD RS10700 WP 001109677.1	IQILSRKTKNNPVL	VGDAGVGKTA	LALGLAQ	RIASGD	VPAEMAK	MRVLE	EL
NTDB id 287 SP RS11210 WP 001109712.1	IQILSRKTKNNPVL	VGDAGVGKTA	LALGLAQ	RIASGD	VPAEMAK	MRVLE	EL
NTDB id 377 SMU RS09275 WP 002262344.1	VQILSRKTKNNPVL	VGDAGVGKTA	LALGLAQ	RIASGD	VPAEMAK	MRVLE	EL
NTDB id 297 STER RS00545 WP 011680614.1	IQVLSRRTKNNPVL	VGEAGVGKTA	LALGLAQ	RIASGE	VVPFEL	ADMRI	EL
NTDB id 329 STU RS10020 WP 011225298.1	IQVLSRRTKNNPVL	VGEAGVGKTA	LALGLAQ	RIASGE	VVPFEL	ADMRI	EL
consensus	*	*!***	*!***	*!***	*!***	*!***	*!***

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NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

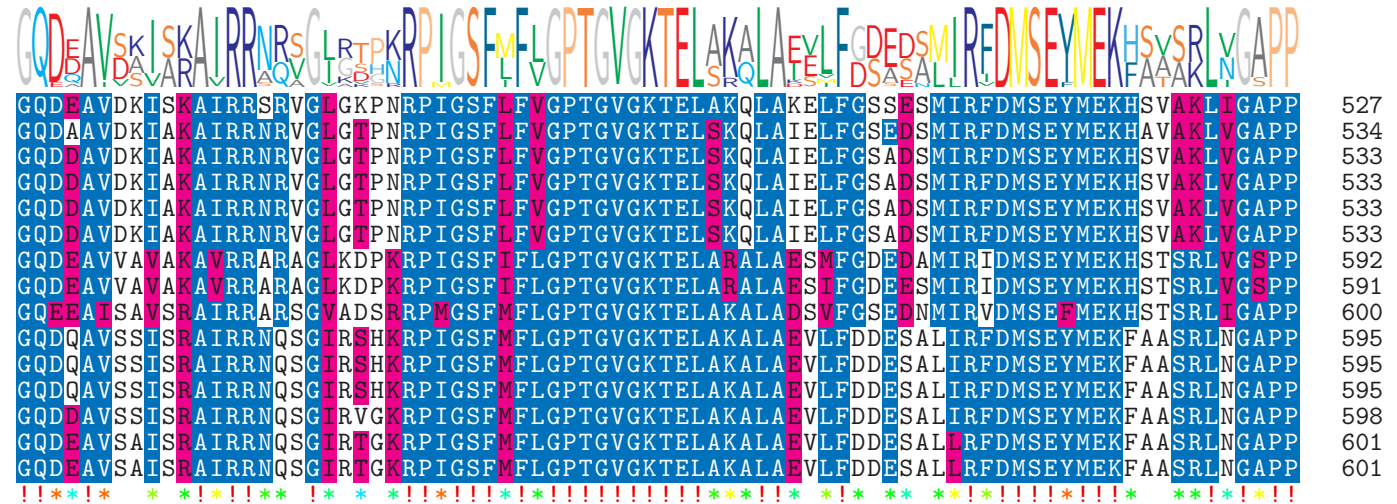
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consensus

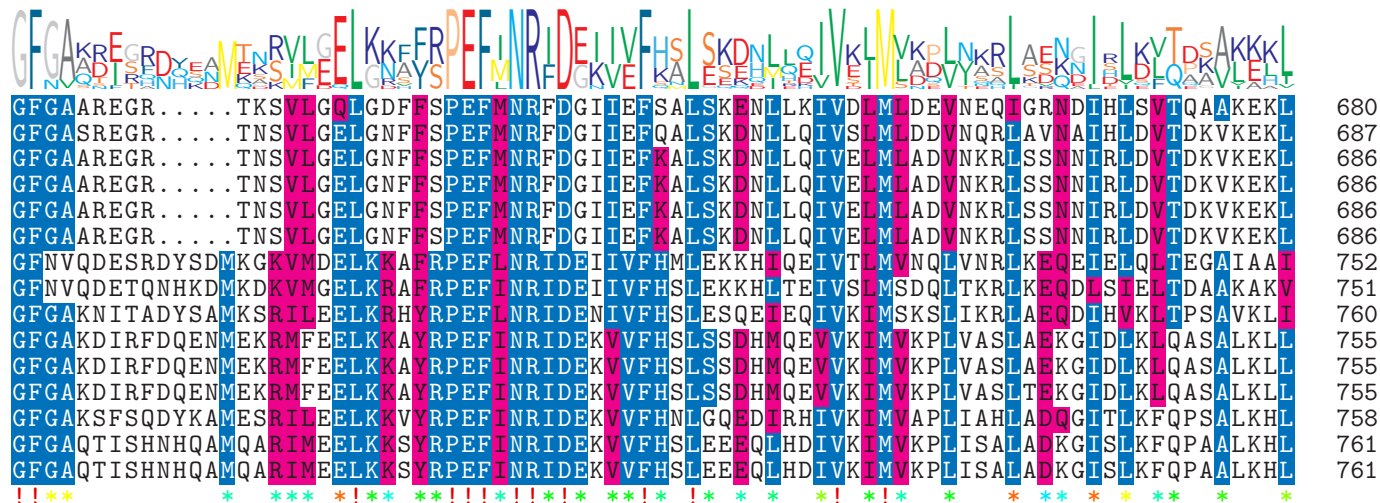
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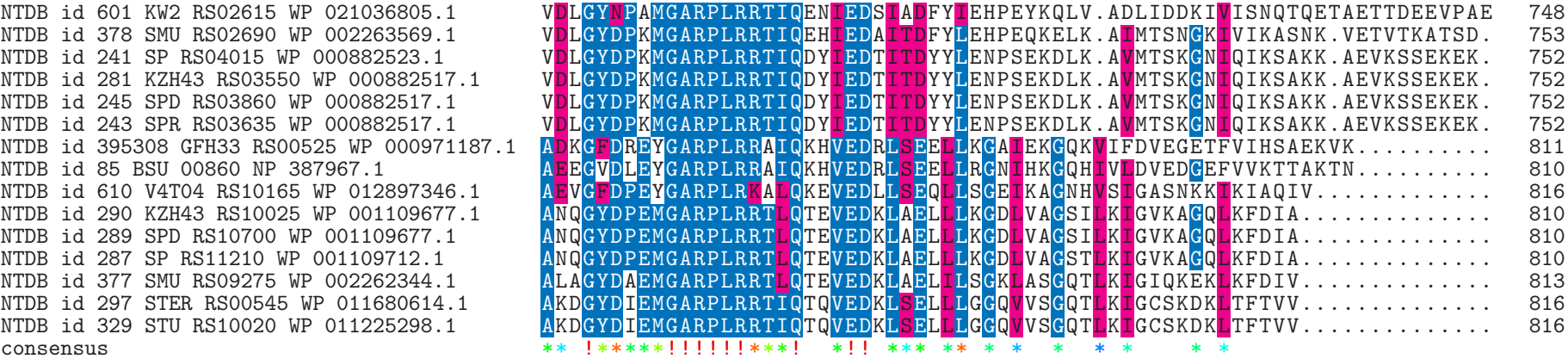
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NTDB id 329 STU RS10020 WP 011225298.1
consensus

```



logo



X non conserved
X similar
X ≥ 50% conserved