

Accession	Protein	Length	Score
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NTDB id 281 KZH43 RS03550 WP 000882517.1	MLCQNCKINDS.....TI	13	13
NTDB id 245 SPD RS03860 WP 000882517.1	MLCQNCKINDS.....TI	13	13
NTDB id 243 SPR RS03635 WP 000882517.1	MLCQNCKINDS.....TI	13	13
NTDB id 241 SP RS04015 WP 000882523.1	MLCQNCKINDS.....TI	13	13
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLI.....GRGQEMS	75	75
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYTPTIDRILEKAEYEAHQYQYGTIESAHLILAAMATTSGSIAYSLLAGMNVDSDDLII.....DLEDLSSHVKVKR	77	77
NTDB id 36346 SMI RS10310 WP 001109716.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE....EVALELTETNYSQDETF	75	75
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE....EVALELTETNYSQDETF	75	75
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE....EVALELTETNYSQDETF	75	75
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE....EVALELTETNYSQDETF	75	75
NTDB id 377 SMU RS09275 WP 002262344.1	...MTDYSLKMQEVFRLAQFEAAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFADVRVFEDYQAAAILAIGKKPKSSSN..DI	79	79
NTDB id 297 STER RS00545 WP 011680614.1	...MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVTGSAVYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDII	81	81
NTDB id 329 STU RS10020 WP 011225298.1	...MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVPGSAVYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDII	81	81
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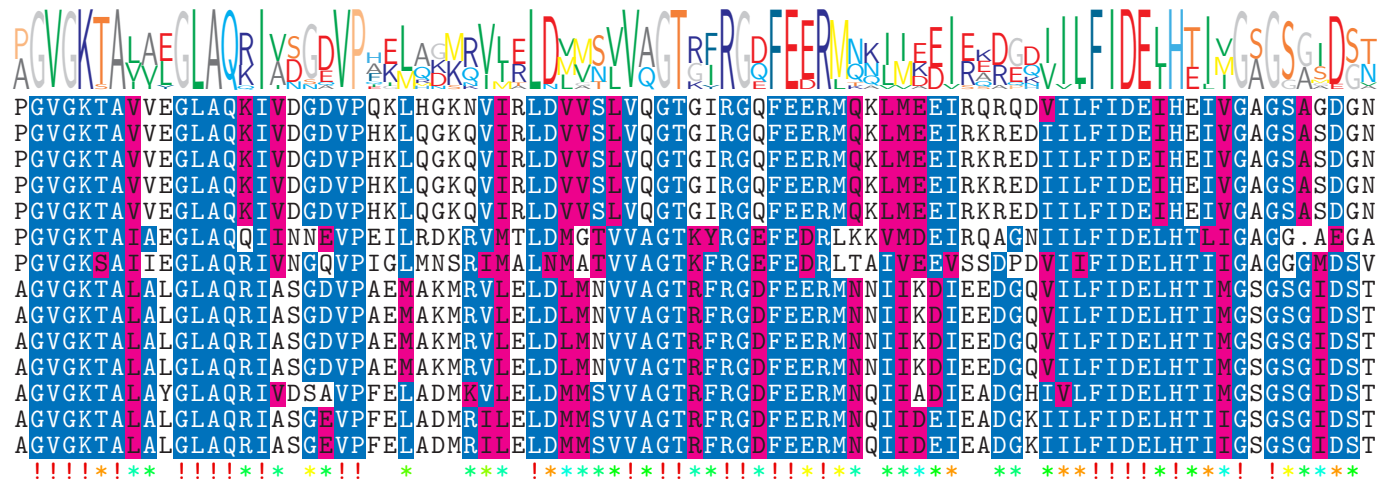
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NTDB id	297	STER RS00545 WP 011680614.1	DLRAQSPALEAMLQEAEIASVTGAVEVGSEHVLMFAFLH.K.DLMVCRLLEVAGFYKYDDSDKPRIIDLRRSLERNAGLSKQDL	164
NTDB id	329	STU RS10020 WP 011225298.1	DLRAQSPALEAMLQEAGGIASVTGAVEVGSEHVLMFAFLH.K.DLMVCRLLEVAGFYKYDDSDKPRIIDLRRSLERNAGLSKQDL	164
consensus			* * * * *	

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NTDB id 85	BSU 00860 NP 387967.1	GSSAAGTNSNANTPTLDSLARDLTAIAKEDSLDPVIGRSKEIQRVIEVLSRRRTKNNPVLIGE	209
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NTDB id 36346	SMI RS10310 WP 001109716.1	KA.LRQRHRTV.....ADKQNSMANMMGMPQTTPSGGLEDYTHDLTEQARSQKLEPVGIRGDKEISRMIIQLSRRTKNNPVLVGD	235
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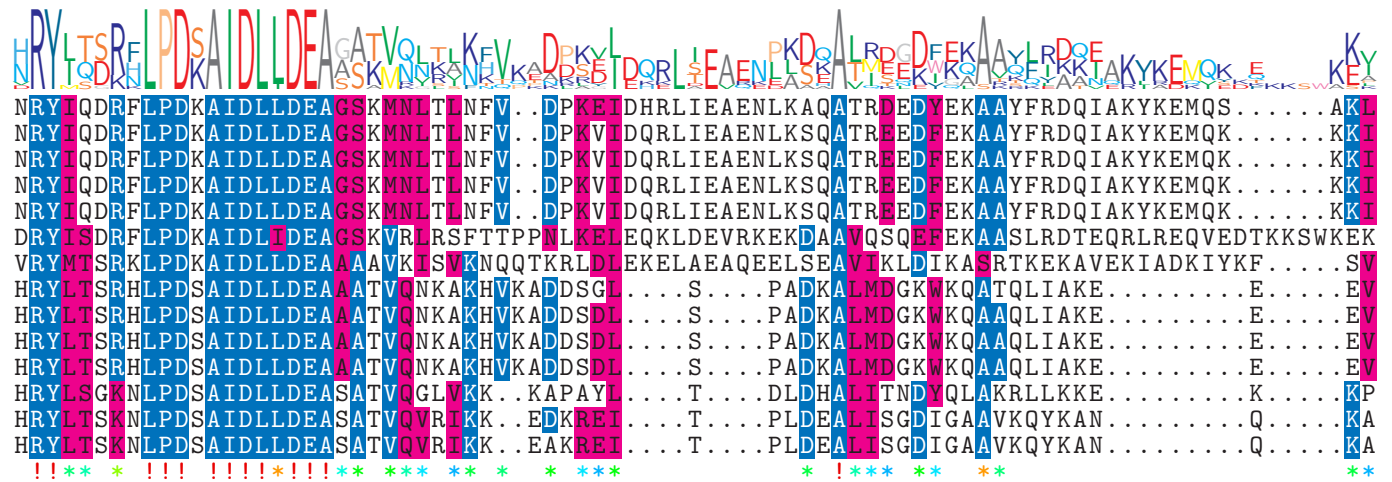
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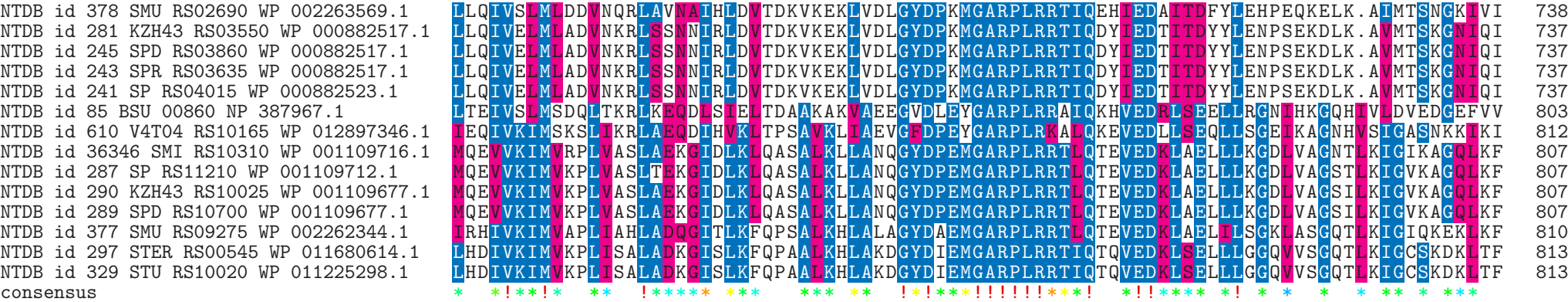
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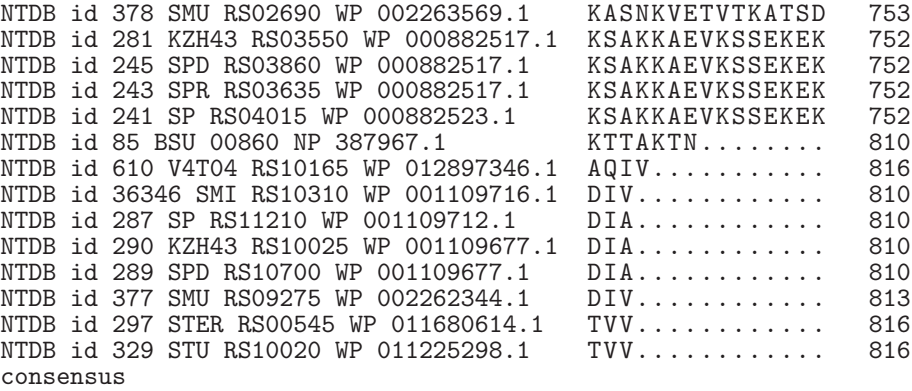


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☐ non conserved
☒ similar
☒ ≥ 50% conserved