

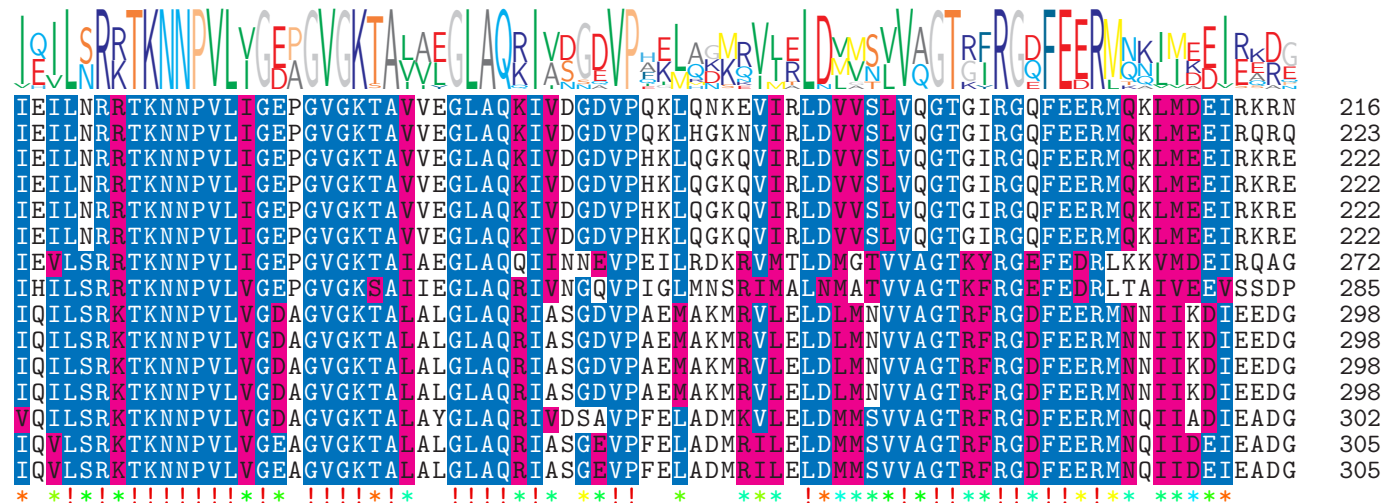
NTDB id 601 KW2 RS02615 WP 021036805.1	MLCQNCNINEA.....	11
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NTDB id 281 KZH43 RS03550 WP 000882517.1	MLCQNCNINDS.....	11
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NTDB id 243 SPR RS03635 WP 000882517.1	MLCQNCNINDS.....	11
NTDB id 241 SP RS04015 WP 000882523.1	MLCQNCNINDS.....	11
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLI.....GR	70
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYPTPLDRILEKAE EYAHQYQYGTIESAHLLAAMATTSGSIAYSLLAGMNV DSSDLLI.....DLEDLSSH	72
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
NTDB id 208208 BSL41 RS02145 WP 001109705.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
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NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
NTDB id 377 SMU RS09275 WP 002262344.1	MTDYS LKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEF EADVRFEDYQAAAILAIGKKPKSSS	76
NTDB id 297 STER RS00545 WP 011680614.1	MTIYSRKMQAIFHRAQLEAERFESPFL ETWHVLLAMVEVTGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLS	76
NTDB id 329 STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFL ETWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLS	76
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NTDB id 281 KZH43 RS03550 WP 000882517.1	...TTHLYTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDLNNRD.....FDP.FG	DFFNDLNNF	69
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NTDB id 85 BSU 00860 NP 387967.1	GQEMSQTIHYTPRAKKVLELSMDEARKLGHSYVGTETILLGLIREGEGVAARVLNNLGVSLNKARQQVLQL.....		141
NTDB id 610 V4T04 RS10165 WP 012897346.1	VKVKRSTLRFSPRAEVEVMTAASFLAIHNNSEAVGTETILLYALLQVEDGFGFLQLLKL.....QKINIVS	LRKELEKR	143
NTDB id 287 SP RS11210 WP 001109712.1	QDETFTELPFSRRLQVLFD EAEYVASV VHA KVLGTEHVLYAILHDSNALATRILERAGFSYEDKKDQVKIAALRRN	LEER	150
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consensus	*** * * * * *	***** * * *	** *

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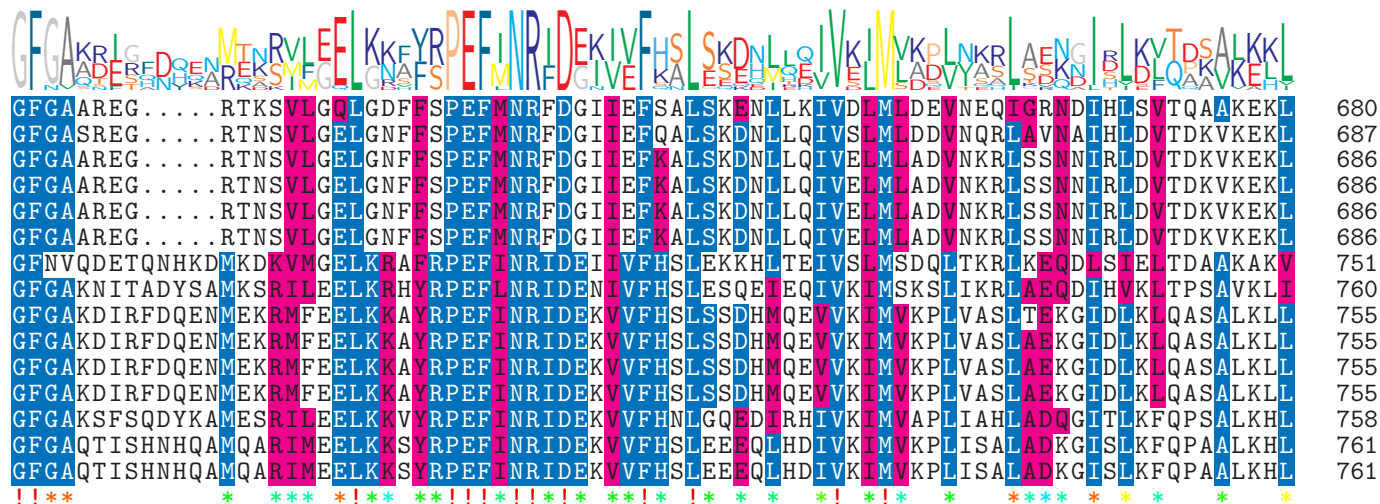
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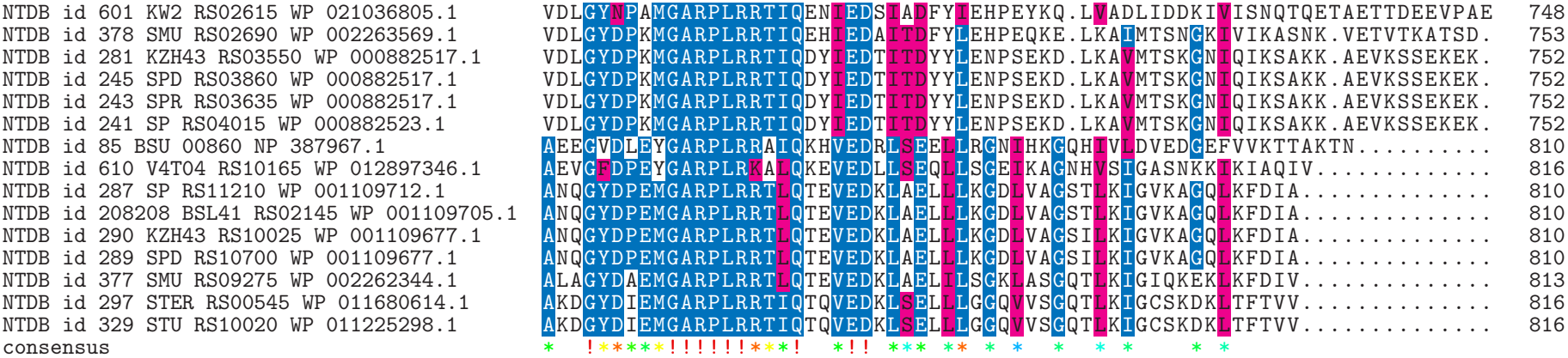
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consensus

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logo



 non conserved
 similar
 ≥ 50% conserved