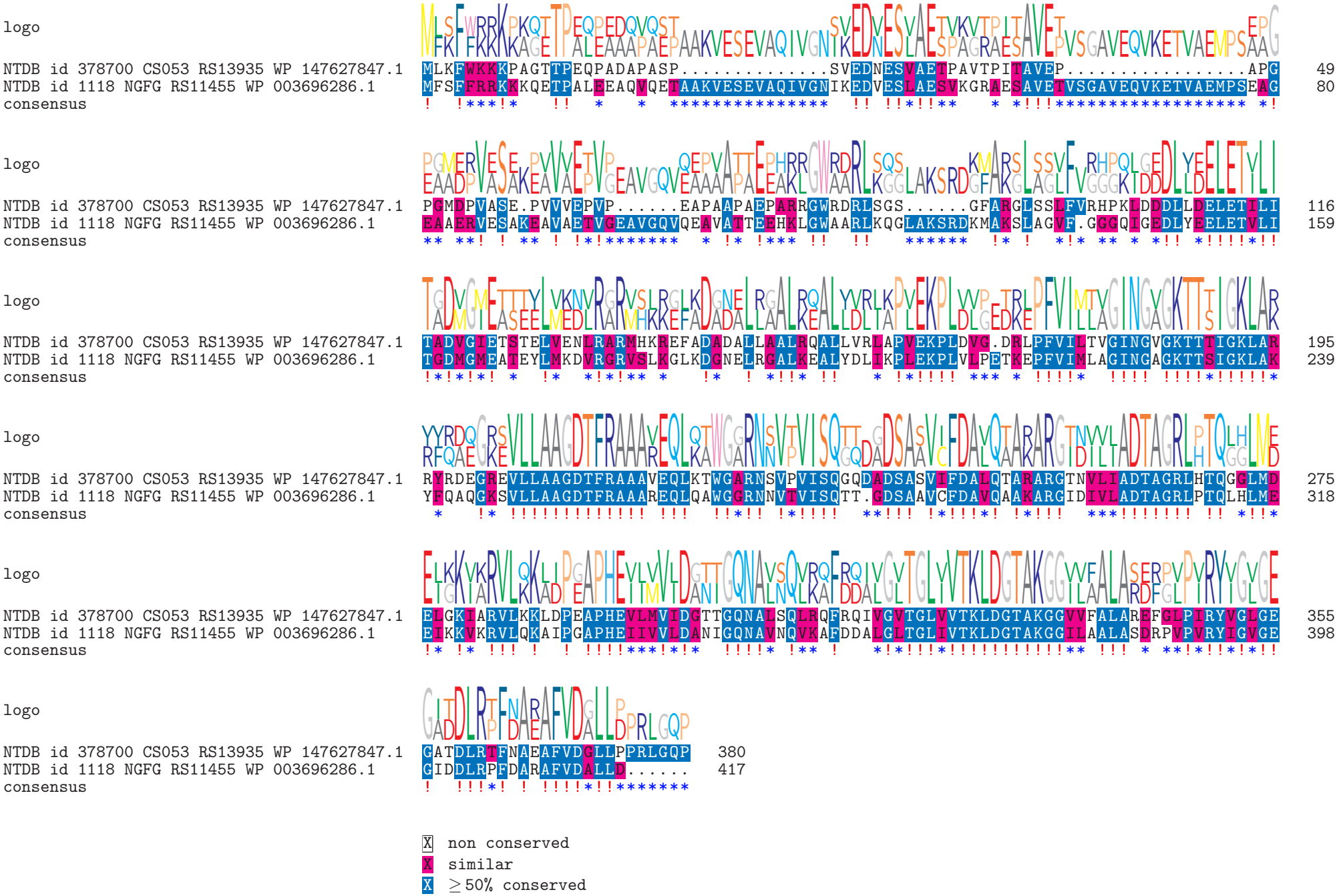
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NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
NTDB id 378700 CS053 RS13935 WP 147627847.1 PGMDPVASE.PVVVEPVP.....EAPAAPAEPARRGWRDRLSGS.....GFARGLSSLFVRHPKLDDDLLDELETILI 116
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consensus
NTDB id 378700 CS053 RS13935 WP 147627847.1 TADVGIETSTELVENLRARMHKREFADADALLAALRQALLVRLAPVEKPLDVG.DRLPFVILTVGINGVGKTTTIGKLAR 195
NTDB id 1118 NGFG RS11455 WP 003696286.1 TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK 239
consensus
NTDB id 378700 CS053 RS13935 WP 147627847.1 RYRDEGREVLLAAGDTFRAAAVEQLKTWGARNSVPVISQGQDADSASVIFDALQTARARGTNVLIADTAGRLHTQGGLMD 275
NTDB id 1118 NGFG RS11455 WP 003696286.1 YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTT.GDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLME 318
consensus
NTDB id 378700 CS053 RS13935 WP 147627847.1 ELGKIARVLKKLDPEAPHEVLMVIDGTTGQNALSQLRQFRQIVGVTGLVVTKLDGTAKGGVVFALAREFGLPIRYVGLGE 355
NTDB id 1118 NGFG RS11455 WP 003696286.1 EIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGE 398
consensus
NTDB id 378700 CS053 RS13935 WP 147627847.1 GATDLRTFNAEAFVDGLLPPRLGQP 380
NTDB id 1118 NGFG RS11455 WP 003696286.1 GIDDLRPFDARAFVDALLD..... 417
consensus
X non conserved
X similar
X ≥ 50% conserved