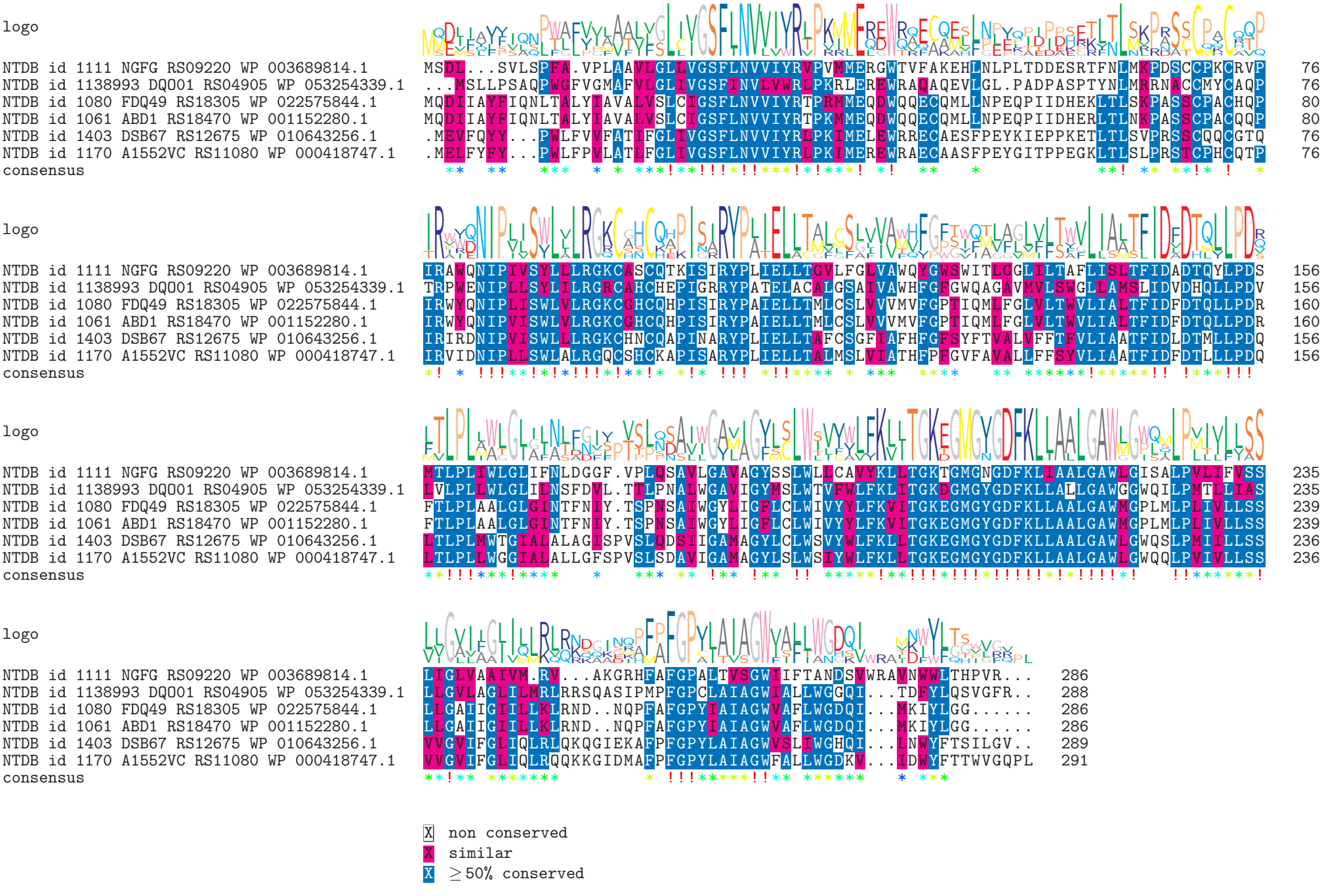
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NTDB id 1138993 DQ001 RS04905 WP 053254339.1
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NTDB id 1061 ABD1 RS18470 WP 001152280.1
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consensus
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...MSLLPSAQPWGFVGMAFVLGLIVGSFINVLVWRLPKRLERERWRAQAQEVLGL.PADPASPTYNLMRRNACCMYCAQP 76
MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
.MEVFQYY...PWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
.MELFYFY...PWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
logo
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consensus
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TRPWENIPLLSYLILRGRCAHCHEPIGRRYPATELACALGSAIVAWHFGFGWQAGAVMVLSWGLLAMSLIDVDHQLLPDV 156
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IRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPDQ 156
logo
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consensus
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LVLPLLWLGLILNSFDVL.TTLPNALWGAVIGYMSLWTVFWLFKLITGKDGMGYGDFKLLALLGAWGGWQILPMTLLIAS 235
FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
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LTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLSS 236
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logo
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NTDB id 1061 ABD1 RS18470 WP 001152280.1
NTDB id 1403 DSB67 RS12675 WP 010643256.1
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consensus
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LLGVLAGLILMRLRRSQASIPMPFGPCLAIAGWIALLWGGQI...TDFYLQSVGFR.. 288
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VVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
VVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
X non conserved
X similar
X ≥ 50% conserved