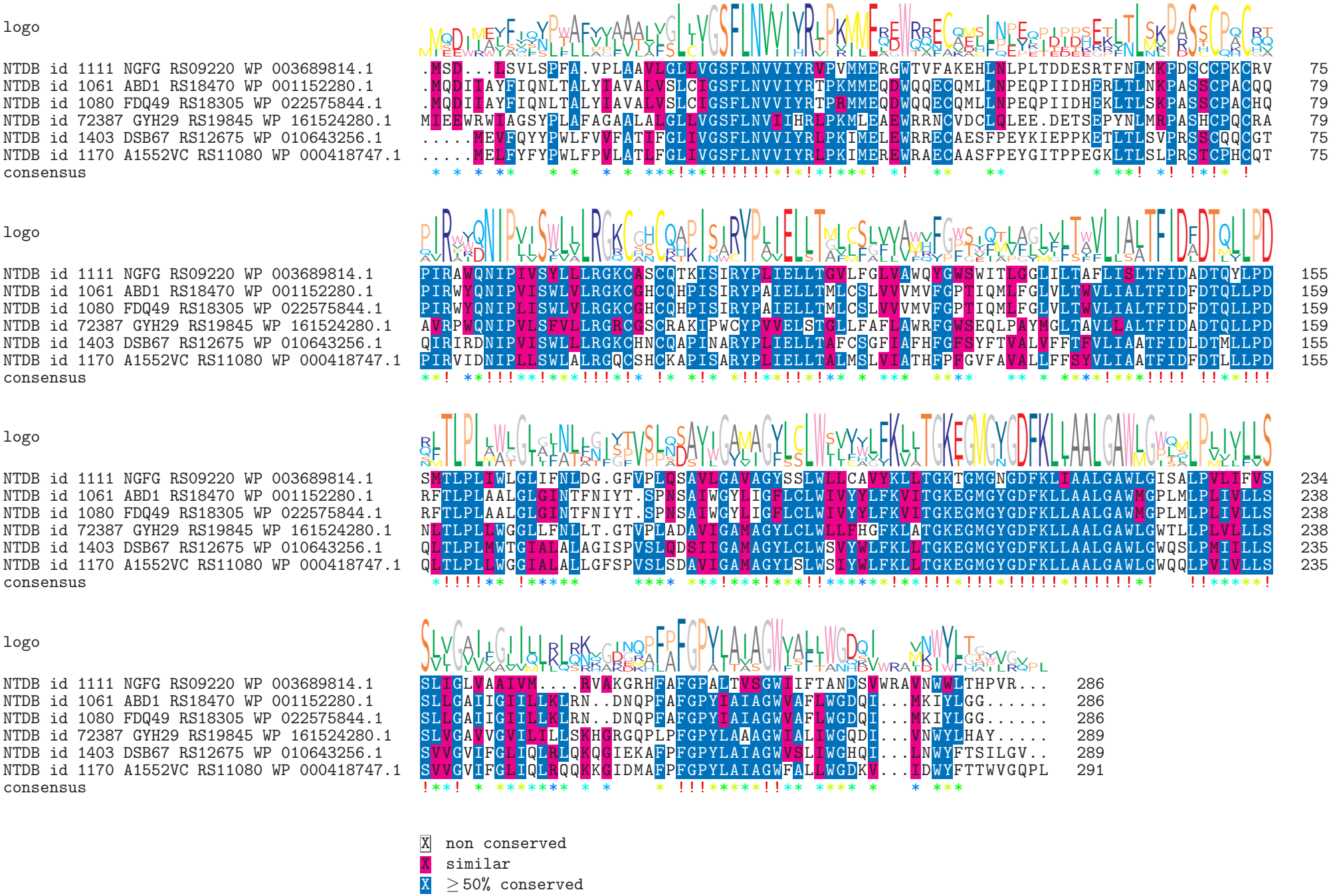
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 .MSD...LSVLSPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCRV 75
NTDB id 1061 ABD1 RS18470 WP 001152280.1 .MQDIIAYFIQNLTALYTAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQ 79
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 .MQDIIAYFIQNLTALYTAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQ 79
NTDB id 72387 GYH29 RS19845 WP 161524280.1 MIEEWRWIAGSYPLAFAGAALALGLLVGSFLNVIIHRLPKMLEAEWRRNCVDCLQLEE.DETSEPYNLMRPASHCPQCRA 79
NTDB id 1403 DSB67 RS12675 WP 010643256.1 .....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGT 75
NTDB id 1170 A1552VC RS11080 WP 000418747.1 .....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQT 75
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 PIRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPD 155
NTDB id 1061 ABD1 RS18470 WP 001152280.1 PIRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPD 159
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 PIRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPD 159
NTDB id 72387 GYH29 RS19845 WP 161524280.1 AVRPWQNIPVLSFVLLRGRCGSCRAKIPWCYPVVELSTGLLFAFLAWRFGWSEQLPAYMGLTAVLLALTFIDADTQLLPD 159
NTDB id 1403 DSB67 RS12675 WP 010643256.1 QIRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFTAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLPD 155
NTDB id 1170 A1552VC RS11080 WP 000418747.1 PIRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPD 155
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 SMTLPLIWLGLIFNLDG.GFVPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFVS 234
NTDB id 1061 ABD1 RS18470 WP 001152280.1 RFTLPLAALGLGINTFNIYT.SPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLS 238
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 RFTLPLAALGLGINTFNIYT.SPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLS 238
NTDB id 72387 GYH29 RS19845 WP 161524280.1 NLTLPLLWGGLLFNLLT.GTVPLADAVIGAMAGYLCLWLLFHGFKLATGKEGMGYGDFKLLAALGAWLGWTLLPLVLLLS 238
NTDB id 1403 DSB67 RS12675 WP 010643256.1 QLTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLS 235
NTDB id 1170 A1552VC RS11080 WP 000418747.1 QLTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLS 235
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 SLIGLVAAIVM....RVAKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 SLLGAIIGIILLKLRN..DNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 SLLGAIIGIILLKLRN..DNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 72387 GYH29 RS19845 WP 161524280.1 SLVGAVVGVILILLSKHGRGQPLPFGPYLAAAGWIALIWGQDI...VNWYLHAY..... 289
NTDB id 1403 DSB67 RS12675 WP 010643256.1 SVVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
NTDB id 1170 A1552VC RS11080 WP 000418747.1 SVVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
consensus
X non conserved
X similar
X ≥ 50% conserved