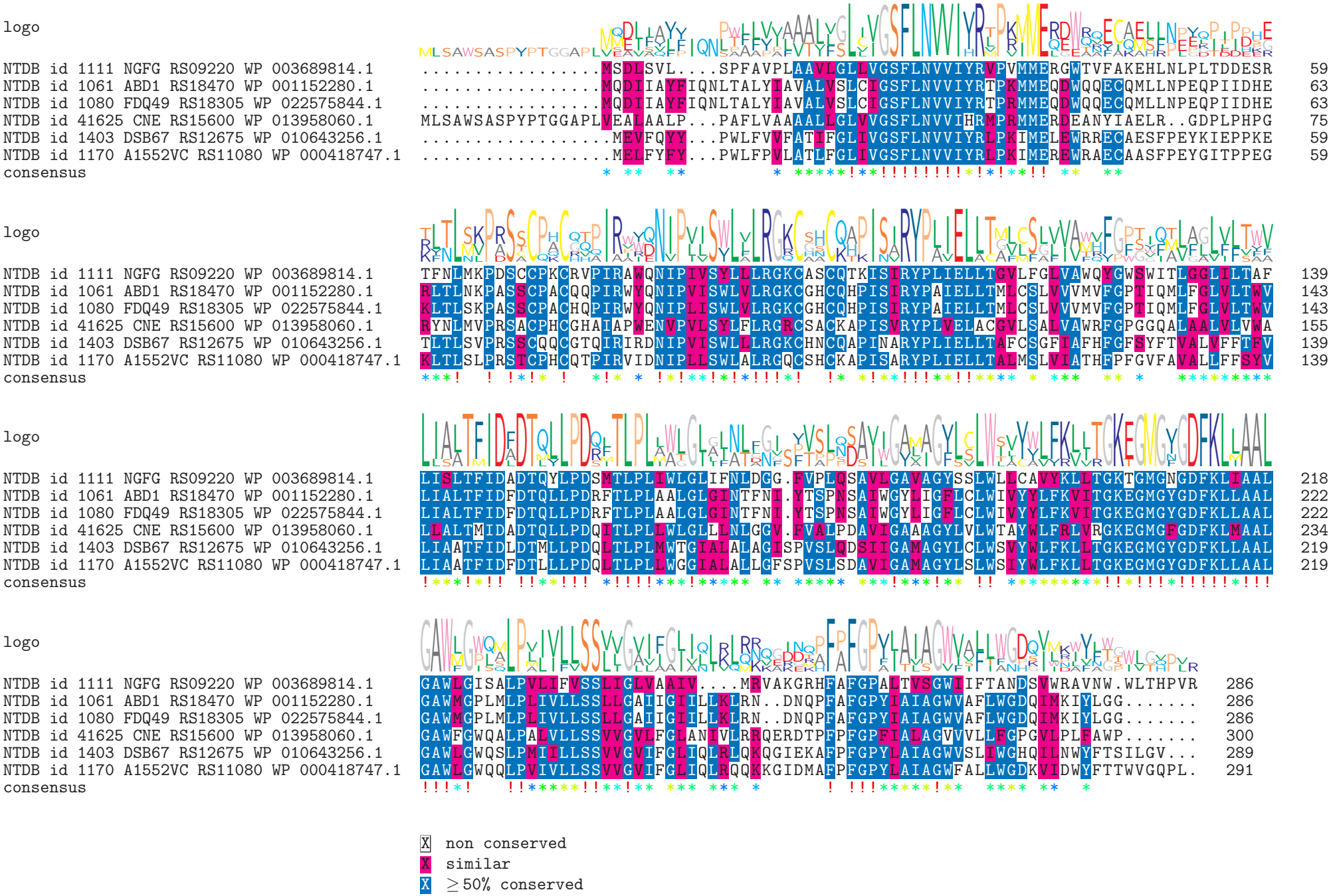
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 .....MSDLSVL....SPFAVPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESR 59
NTDB id 1061 ABD1 RS18470 WP 001152280.1 .....MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHE 63
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 .....MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHE 63
NTDB id 41625 CNE RS15600 WP 013958060.1 MLSAWSASPYPTGGAPLVEALAALP...PAFLVAAAALLGLVVGSFLNVVIHRMPRMMERDEANYIAELR..GDPLPHPG 75
NTDB id 1403 DSB67 RS12675 WP 010643256.1 .....MEVFQYY...PWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKE 59
NTDB id 1170 A1552VC RS11080 WP 000418747.1 .....MELFYFY...PWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEG 59
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 TFNLMKPDSCCPKCRVPIRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAF 139
NTDB id 1061 ABD1 RS18470 WP 001152280.1 RLTLNKPASSCPACQQPIRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWV 143
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 KLTLSKPASSCPACHQPIRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWV 143
NTDB id 41625 CNE RS15600 WP 013958060.1 RYNLMVPRSACPHCGHAIAPWENVPVLSYLFLRGRCSACKAPISVRYPLVELACGVLSALVAWRFGPGGQALAALVLVWA 155
NTDB id 1403 DSB67 RS12675 WP 010643256.1 TLTLSVPRSSCQQCGTQIRIRDNIPVISWLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFV 139
NTDB id 1170 A1552VC RS11080 WP 000418747.1 KLTLSLPRSTCPHCQTPIRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYV 139
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 LISLTFIDADTQYLPDSMTLPLIWLGLIFNLDGG.FVPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAAL 218
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LIALTFIDFDTQLLPDRFTLPLAALGLGINTFNI.YTSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAAL 222
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LIALTFIDFDTQLLPDRFTLPLAALGLGINTFNI.YTSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAAL 222
NTDB id 41625 CNE RS15600 WP 013958060.1 LLALTMIDADTQLLPDQLTLPLLWLGLLLNLGGV.FVALPDAVIGAAAGYLVLWTAYWLFRLVRGKEGMGRGDFKLMAAL 234
NTDB id 1403 DSB67 RS12675 WP 010643256.1 LIAATFIDLDTMLLPDQLTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAAL 219
NTDB id 1170 A1552VC RS11080 WP 000418747.1 LIAATFIDFDTLLLPDQLTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAAL 219
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 GAWLGISALPVLIFVSSLIGLVAAIV....MRVAKGRHFAFGPALTVSGWIIFTANDSVWRAVNW.WLTHPVR 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 GAWMGPLMLPLIVLLSSLLGAIIGIILLKLRN..DNQPFAFGPYIAIAGWVAFLWGDQIMKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 GAWMGPLMLPLIVLLSSLLGAIIGIILLKLRN..DNQPFAFGPYIAIAGWVAFLWGDQIMKIYLGG..... 286
NTDB id 41625 CNE RS15600 WP 013958060.1 GAWFGWQALPALVLLSSVVGVLFGLANIVLRRQERDTPFPFGPFIALAGVVVLLFGPGVLPLFAWP..... 300
NTDB id 1403 DSB67 RS12675 WP 010643256.1 GAWLGWQSLPMIILLSSVVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQILNWYFTSILGV... 289
NTDB id 1170 A1552VC RS11080 WP 000418747.1 GAWLGWQQLPVIVLLSSVVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKVIDWYFTTWVGQPL. 291
consensus
X non conserved
X similar
X ≥ 50% conserved