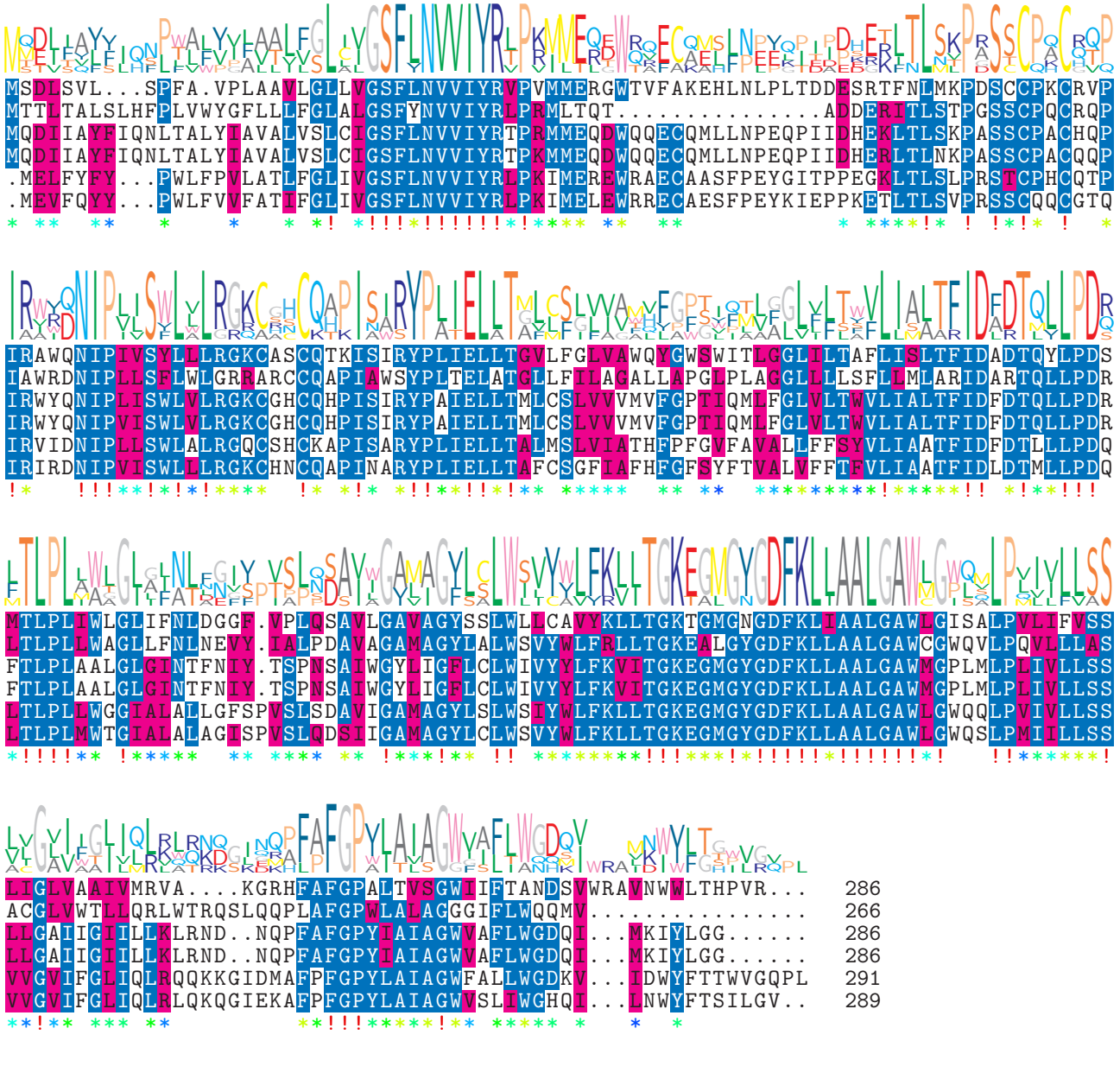

logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 MSDLSVL...SPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCRVP 76
NTDB id 203204 BBD64 RS04645 WP 008807384.1 MTTLTALSLHFPLVWYGFLLLFGLALGSFYNVVIYRLPRMLTQT..............ADDERITLSTPGSSCPQCRQP 65
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
NTDB id 1170 A1552VC RS11080 WP 000418747.1 .MELFYFY...PWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
NTDB id 1403 DSB67 RS12675 WP 010643256.1 .MEVFQYY...PWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 IRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPDS 156
NTDB id 203204 BBD64 RS04645 WP 008807384.1 IAWRDNIPLLSFLWLGRRARCCQAPIAWSYPLTELATGLLFILAGALLAPGLPLAGGLLLLSFLLMLARIDARTQLLPDR 145
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 IRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1061 ABD1 RS18470 WP 001152280.1 IRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1170 A1552VC RS11080 WP 000418747.1 IRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPDQ 156
NTDB id 1403 DSB67 RS12675 WP 010643256.1 IRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLPDQ 156
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 MTLPLIWLGLIFNLDGGF.VPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFVSS 235
NTDB id 203204 BBD64 RS04645 WP 008807384.1 LTLPLLWAGLLFNLNEVY.IALPDAVAGAMAGYLALWSVYWLFRLLTGKEALGYGDFKLLAALGAWCGWQVLPQVLLLAS 224
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1061 ABD1 RS18470 WP 001152280.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1170 A1552VC RS11080 WP 000418747.1 LTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLSS 236
NTDB id 1403 DSB67 RS12675 WP 010643256.1 LTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLSS 236
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 LIGLVAAIVMRVA....KGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 203204 BBD64 RS04645 WP 008807384.1 ACGLVWTLLQRLWTRQSLQQPLAFGPWLALAGGGIFLWQQMV.............. 266
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1170 A1552VC RS11080 WP 000418747.1 VVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
NTDB id 1403 DSB67 RS12675 WP 010643256.1 VVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
consensus
X non conserved
X similar
X ≥ 50% conserved