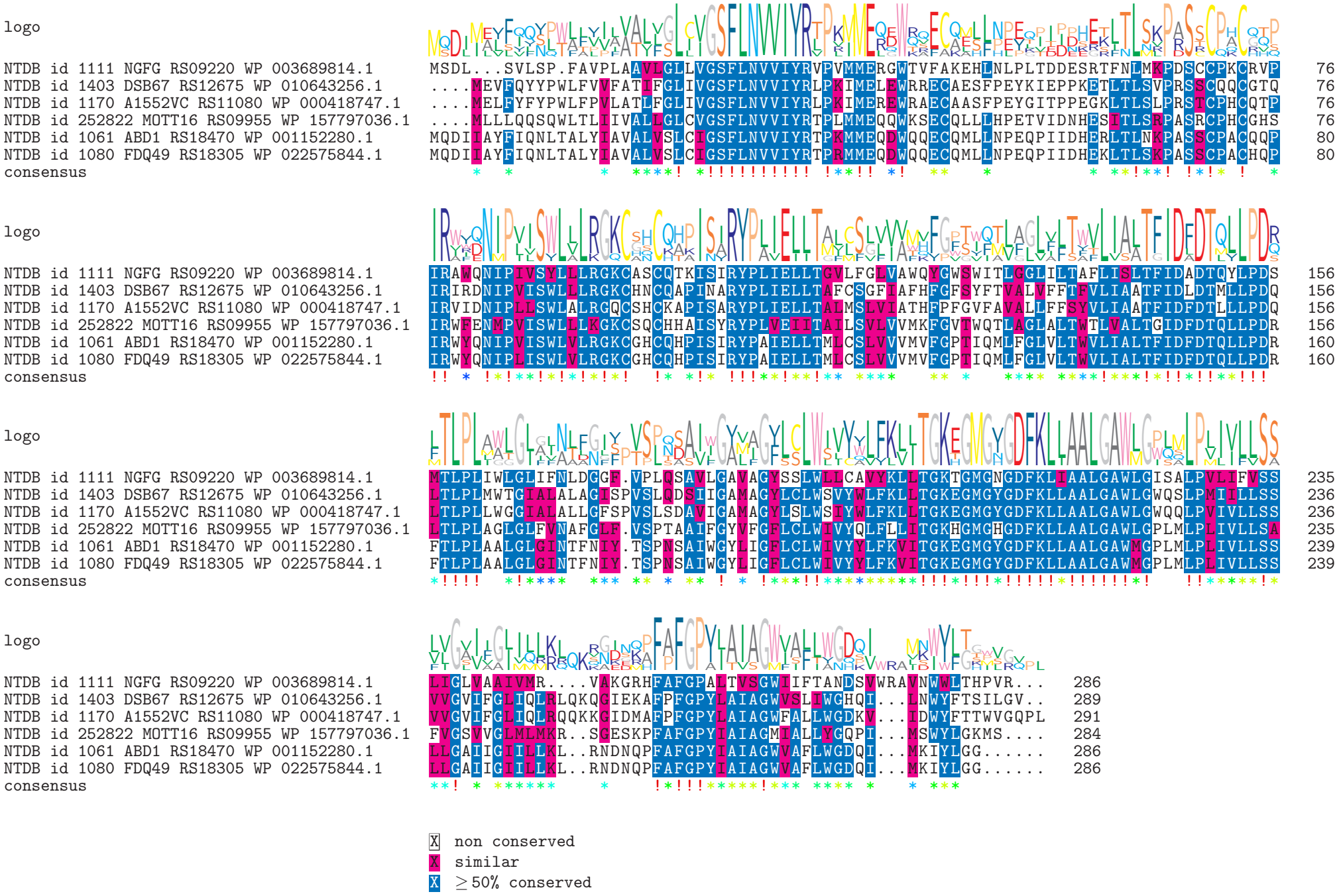
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NTDB id 1403 DSB67 RS12675 WP 010643256.1 ....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
NTDB id 1170 A1552VC RS11080 WP 000418747.1 ....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
NTDB id 252822 MOTT16 RS09955 WP 157797036.1 ....MLLLQQSQWLTLIIVALLGLCVGSFLNVVIYRTPLMMEQQWKSECQLLLHPETVIDNHESITLSRPASRCPHCGHS 76
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
consensus
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NTDB id 1111 NGFG RS09220 WP 003689814.1 IRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPDS 156
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NTDB id 252822 MOTT16 RS09955 WP 157797036.1 IRWFENMPVISWLLLKGKCSQCHHAISYRYPLVEIITAILSVLVVMKFGVTWQTLAGLALTWTLVALTGIDFDTQLLPDR 156
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consensus
logo
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NTDB id 1061 ABD1 RS18470 WP 001152280.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
consensus
logo
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NTDB id 1170 A1552VC RS11080 WP 000418747.1 VVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
NTDB id 252822 MOTT16 RS09955 WP 157797036.1 FVGSVVGLMLMKR..SGESKPFAFGPYIAIAGMIALLYGQPI...MSWYLGKMS.... 284
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LLGAIIGIILLKL..RNDNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LLGAIIGIILLKL..RNDNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
consensus
X non conserved
X similar
X ≥ 50% conserved