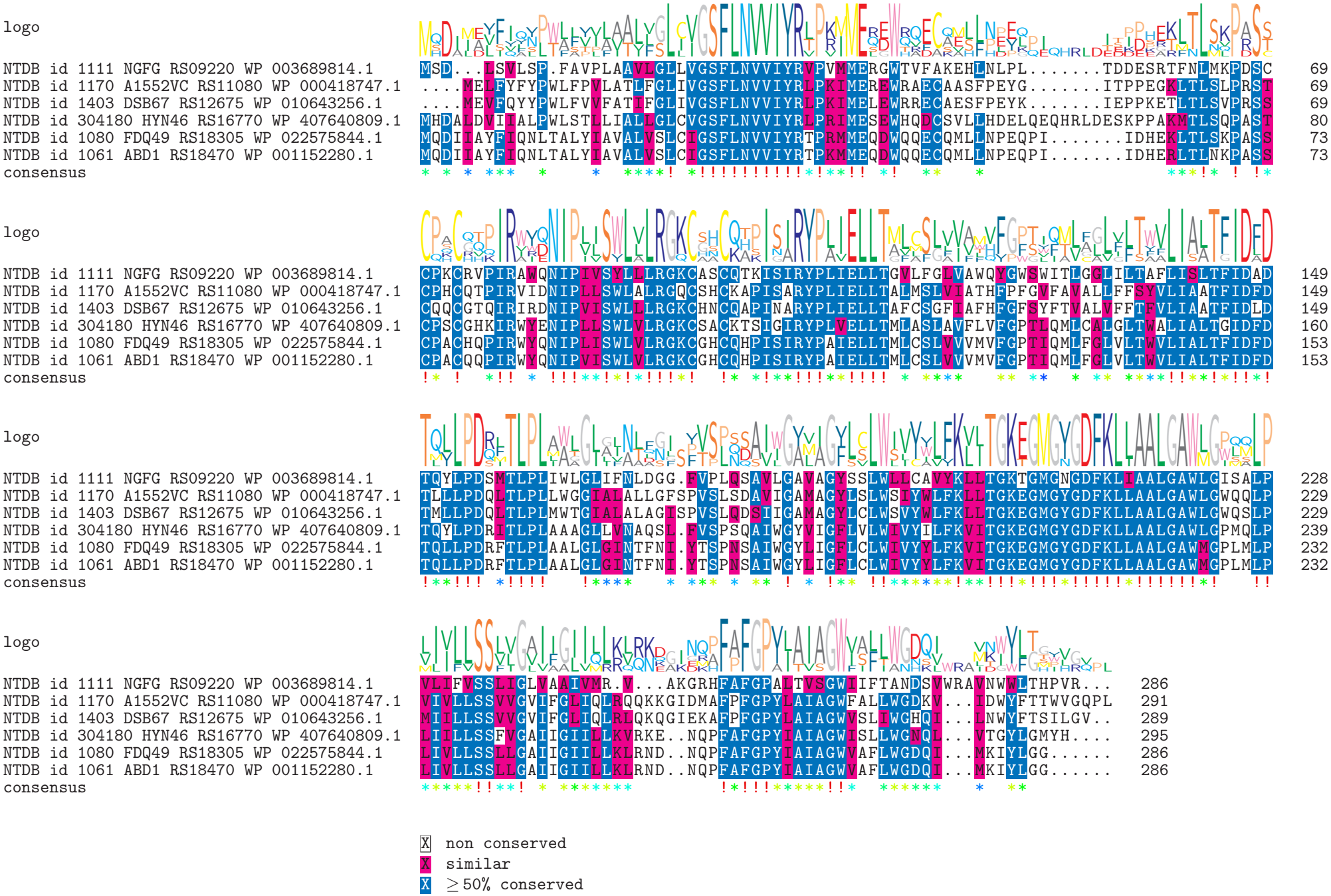
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NTDB id 1170 A1552VC RS11080 WP 000418747.1 ....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMERFWRAECAASFPEYG.....ITPPEGKLTLSLPRST 69
NTDB id 1403 DSB67 RS12675 WP 010643256.1 ....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYK.....IEPPKETLTLSVPRSS 69
NTDB id 304180 HYN46 RS16770 WP 407640809.1 MHDALDVIIALPWLSTLLIALLGLCVGSFLNVVIYRLPRIMESEWHQDCSVLLHDELQEQHRLDESKPPAKMTLSQPAST 80
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPI.....IDHEKLTLSKPASS 73
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPI.....IDHERLTLNKPASS 73
consensus
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NTDB id 1111 NGFG RS09220 WP 003689814.1 CPKCRVPIRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDAD 149
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NTDB id 304180 HYN46 RS16770 WP 407640809.1 TQYLPDRITLPLAAAGLLVNAQSL.FVSPSQAIWGYVIGFLVLWIVYILFKVITGKEGMGYGDFKLLAALGAWLGPMQLP 239
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consensus
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NTDB id 1403 DSB67 RS12675 WP 010643256.1 MIILLSSVVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
NTDB id 304180 HYN46 RS16770 WP 407640809.1 LIILLSSFVGAIIGIILLKVRKE..NQPFAFGPYIAIAGWISLLWGNQL...VTGYLGMYH.... 295
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LIVLLSSLLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
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consensus
X non conserved
X similar
X ≥ 50% conserved