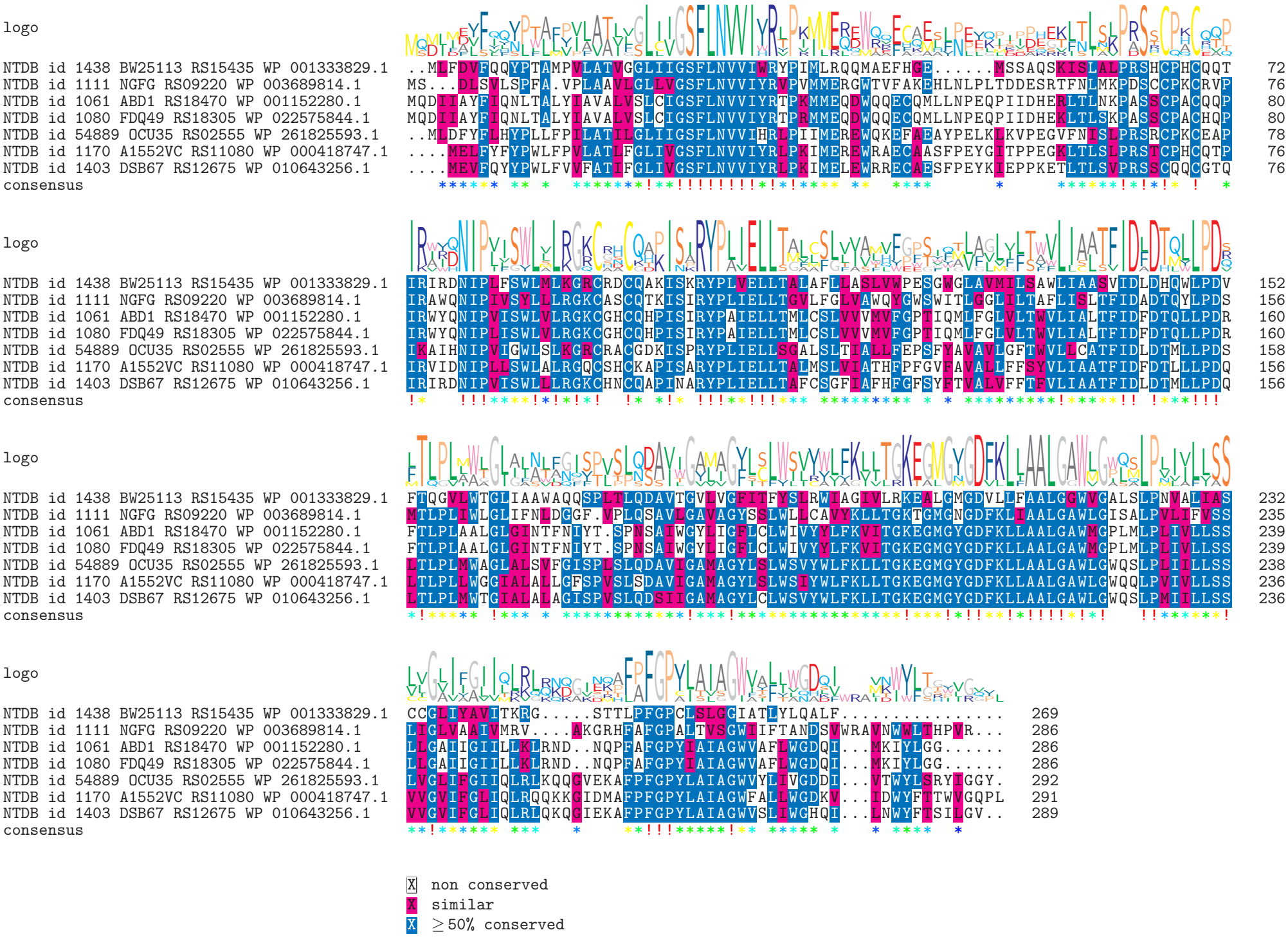
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NTDB id 1111 NGFG RS09220 WP 003689814.1 MS...DLSVLSPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCRVP 76
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
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NTDB id 1170 A1552VC RS11080 WP 000418747.1 ....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMERRWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
NTDB id 1403 DSB67 RS12675 WP 010643256.1 ....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
consensus
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NTDB id 1061 ABD1 RS18470 WP 001152280.1 IRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 IRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
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NTDB id 1111 NGFG RS09220 WP 003689814.1 LIGLVAAIVMRV....AKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
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consensus
X non conserved
X similar
X ≥ 50% conserved