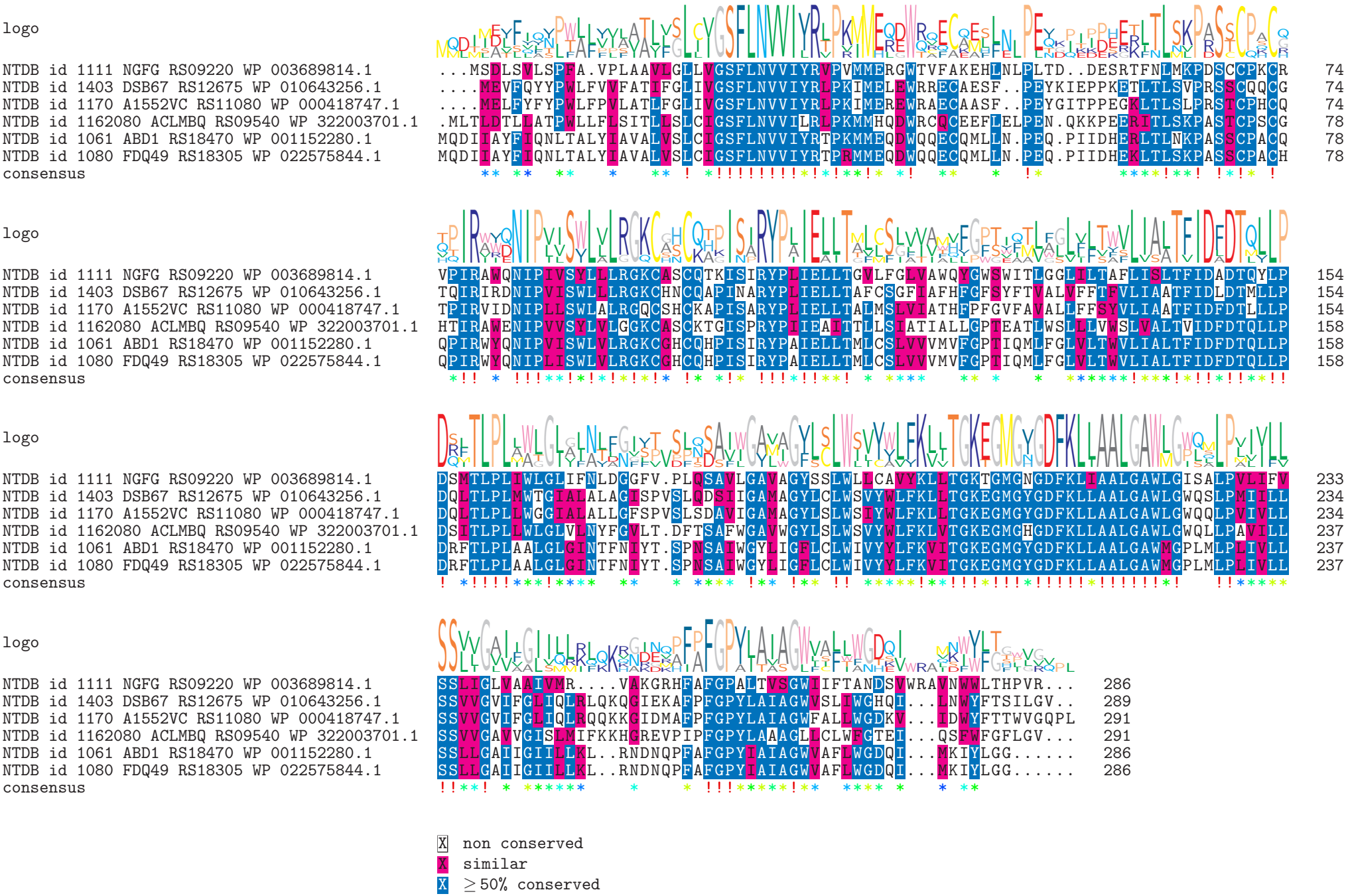
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NTDB id 1162080 ACLMBQ RS09540 WP 322003701.1 .MLTLDTLLATPWLLFLSITLLSLCIGSFLNVVILRLPKMMHQDWRCQCEEFLELPEN.QKKPEERITLSKPASTCPSCG 78
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLN.PEQ.PIIDHERLTLNKPASSCPACQ 78
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLN.PEQ.PIIDHEKLTLSKPASSCPACH 78
consensus
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NTDB id 1111 NGFG RS09220 WP 003689814.1 VPIRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLP 154
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NTDB id 1162080 ACLMBQ RS09540 WP 322003701.1 HTIRAWENIPVVSYLVLGGKCASCKTGISPRYPIIEAITTLLSIATIALLGPTEATLWSLLLVWSLVALTVIDFDTQLLP 158
NTDB id 1061 ABD1 RS18470 WP 001152280.1 QPIRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLP 158
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consensus
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NTDB id 1162080 ACLMBQ RS09540 WP 322003701.1 DSITLPLLWLGLVLNYFGVLT.DFTSAFWGAVWGYLSLWSVYWLFKLVTGKEGMGHGDFKLLAALGAWLGWQLLPAVILL 237
NTDB id 1061 ABD1 RS18470 WP 001152280.1 DRFTLPLAALGLGINTFNIYT.SPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLL 237
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 DRFTLPLAALGLGINTFNIYT.SPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLL 237
consensus
logo
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NTDB id 1170 A1552VC RS11080 WP 000418747.1 SSVVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
NTDB id 1162080 ACLMBQ RS09540 WP 322003701.1 SSVVGAVVGISLMIFKKHGREVPIPFGPYLAAAGLLCLWFGTEI...QSFWFGFLGV... 291
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consensus
X non conserved
X similar
X ≥ 50% conserved