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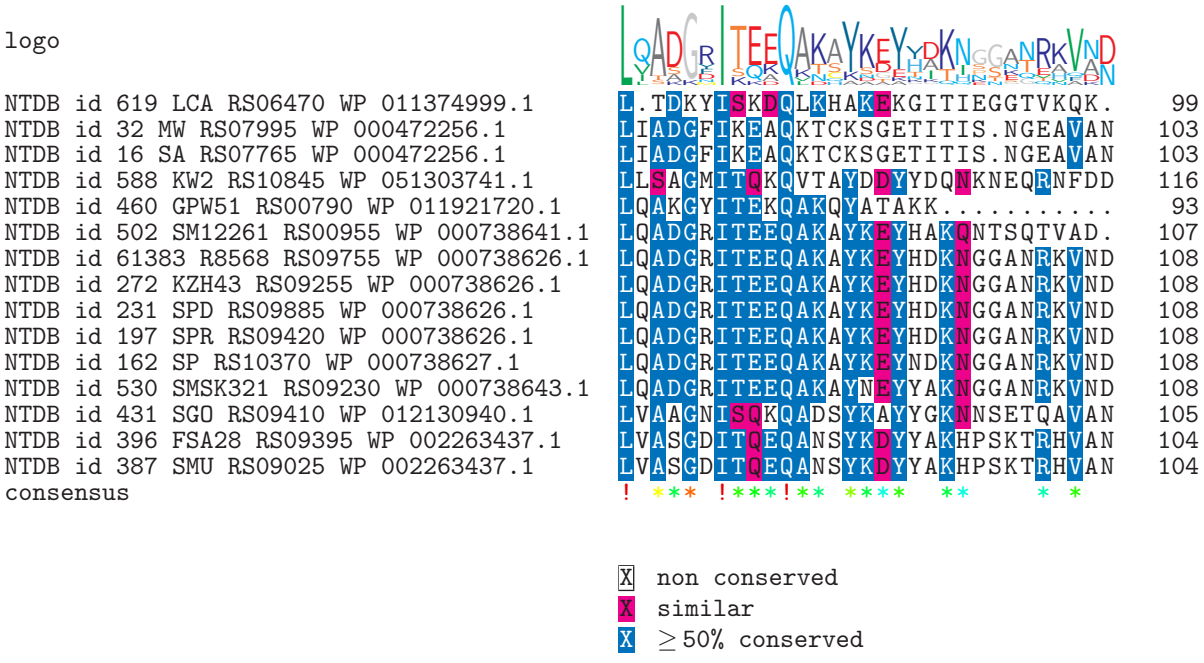
NTDB id 387 SMU RS09025 WP 002263437.1 .....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK 73
consensus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NTDB id 16 SA RS07765 WP 000472256.1 LIADGFIKEAQKTCKSGETITIS.NGEAVAN 103
NTDB id 588 KW2 RS10845 WP 051303741.1 LLSAGMITQKQVTAYDDYYDQNKNEQRNFDD 116
NTDB id 460 GPW51 RS00790 WP 011921720.1 LQAKGYITEKQAKQYATAKK..... 93
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NTDB id 61383 R8568 RS09755 WP 000738626.1 LQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 272 KZH43 RS09255 WP 000738626.1 LQADGRITEEQAKAYKEYHDKNGGANRKVND 108
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NTDB id 396 FSA28 RS09395 WP 002263437.1 LVASGDITQEQANSYKDYYAKHPSKTRHVAN 104
NTDB id 387 SMU RS09025 WP 002263437.1 LVASGDITQEQANSYKDYYAKHPSKTRHVAN 104
consensus
X non conserved
X similar
X ≥ 50% conserved