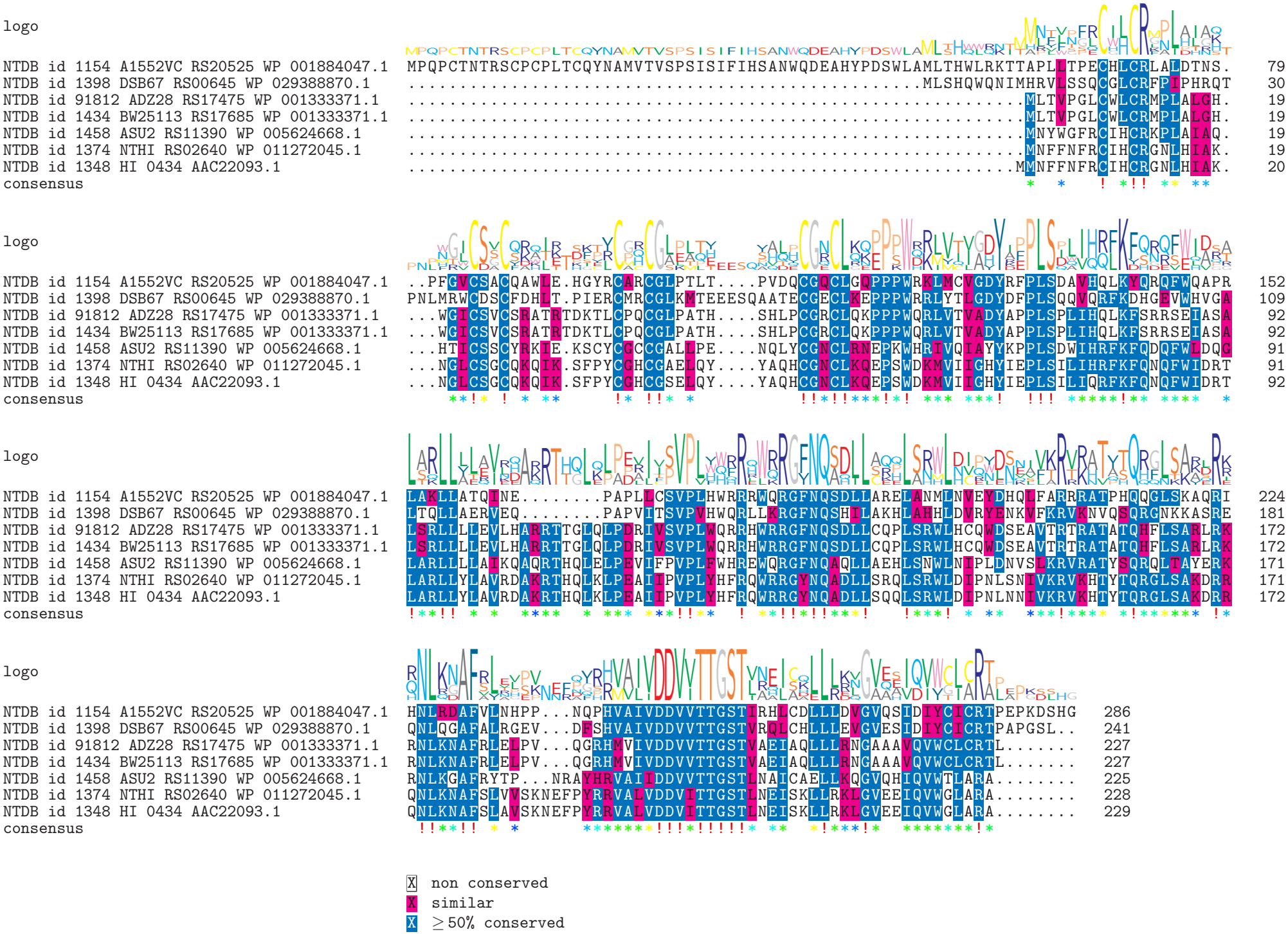
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1 MPQPCTNTRSCPCPLTCQYNAMVTVSPSISIFIHSANWQDEAHYPDSWLAMLTHWLRKTTAPLLTPECHLCRLALDTNS. 79
NTDB id 1398 DSB67 RS00645 WP 029388870.1 .....MLSHQWQNIMHRVLSSQCGLCRFPIPHRQT 30
NTDB id 91812 ADZ28 RS17475 WP 001333371.1 .....MLTVPGLCWLCRMPLALGH. 19
NTDB id 1434 BW25113 RS17685 WP 001333371.1 .....MLTVPGLCWLCRMPLALGH. 19
NTDB id 1458 ASU2 RS11390 WP 005624668.1 .....MNYWGFRCIHCRKPLAIAQ. 19
NTDB id 1374 NTHI RS02640 WP 011272045.1 .....MNFFNFRCIHCRGNLHIAK. 19
NTDB id 1348 HI 0434 AAC22093.1 .....MMNFFNFRCIHCRGNLHIAK. 20
consensus
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1 ..PFGVCSACQAWLE.HGYRCARCGLPTLT....PVDQCGQCLGQPPPWRKLMCVGDYRFPLSDAVHQLKYQRQFWQAPR 152
NTDB id 1398 DSB67 RS00645 WP 029388870.1 PNLMRWCDSCFDHLT.PIERCMRCGLKMTEEESQAATECGECLKEPPPWRRLYTLGDYDFPLSQQVQRFKDHGEVWHVGA 109
NTDB id 91812 ADZ28 RS17475 WP 001333371.1 ...WGICSVCSRATRTDKTLCPQCGLPATH....SHLPCGRCLQKPPPWQRLVTVADYAPPLSPLIHQLKFSRRSEIASA 92
NTDB id 1434 BW25113 RS17685 WP 001333371.1 ...WGICSVCSRATRTDKTLCPQCGLPATH....SHLPCGRCLQKPPPWQRLVTVADYAPPLSPLIHQLKFSRRSEIASA 92
NTDB id 1458 ASU2 RS11390 WP 005624668.1 ...HTICSSCYRKIE.KSCYCGCCGALLPE....NQLYCGNCLRNEPKWHRIVQIAYYKPPLSDWIHRFKFQDQFWLDQG 91
NTDB id 1374 NTHI RS02640 WP 011272045.1 ...NGLCSGCQKQIK.SFPYCGHCGAELQY....YAQHCGNCLKQEPSWDKMVIIGHYIEPLSILIHRFKFQNQFWIDRT 91
NTDB id 1348 HI 0434 AAC22093.1 ...NGLCSGCQKQIK.SFPYCGHCGSELQY....YAQHCGNCLKQEPSWDKMVIIGHYIEPLSILIQRFKFQNQFWIDRT 92
consensus
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1 LAKLLATQINE.....PAPLLCSVPLHWRRRWQRGFNQSDLLARELANMLNVEYDHQLFARRRATPHQQGLSKAQRI 224
NTDB id 1398 DSB67 RS00645 WP 029388870.1 LTQLLAERVEQ.....PAPVITSVPVHWQRLLKRGFNQSHILAKHLAHHLDVRYENKVFKRVKNVQSQRGNKKASRE 181
NTDB id 91812 ADZ28 RS17475 WP 001333371.1 LSRLLLLEVLHARRTTGLQLPDRIVSVPLWQRRHWRRGFNQSDLLCQPLSRWLHCQWDSEAVTRTRATATQHFLSARLRK 172
NTDB id 1434 BW25113 RS17685 WP 001333371.1 LSRLLLLEVLHARRTTGLQLPDRIVSVPLWQRRHWRRGFNQSDLLCQPLSRWLHCQWDSEAVTRTRATATQHFLSARLRK 172
NTDB id 1458 ASU2 RS11390 WP 005624668.1 LARLLLLAIKQAQRTHQLELPEVIFPVPLFWHREWQRGFNQAQLLAEHLSNWLNTPLDNVSLKRVRATYSQRQLTAYERK 171
NTDB id 1374 NTHI RS02640 WP 011272045.1 LARLLYLAVRDAKRTHQLKLPEAIIPVPLYHFRQWRRGYNQADLLSRQLSRWLDIPNLSNIVKRVKHTYTQRGLSAKDRR 171
NTDB id 1348 HI 0434 AAC22093.1 LARLLYLAVRDAKRTHQLKLPEAIIPVPLYHFRQWRRGYNQADLLSQQLSRWLDIPNLNNIVKRVKHTYTQRGLSAKDRR 172
consensus
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1 HNLRDAFVLNHPP...NQPHVAIVDDVVTTGSTIRHLCDLLLDVGVQSIDIYCICRTPEPKDSHG 286
NTDB id 1398 DSB67 RS00645 WP 029388870.1 QNLQGAFALRGEV...DFSHVAIVDDVVTTGSTVRQLCHLLLEVGVESIDIYCICRTPAPGSL.. 241
NTDB id 91812 ADZ28 RS17475 WP 001333371.1 RNLKNAFRLELPV...QGRHMVIVDDVVTTGSTVAEIAQLLLRNGAAAVQVWCLCRTL..... 227
NTDB id 1434 BW25113 RS17685 WP 001333371.1 RNLKNAFRLELPV...QGRHMVIVDDVVTTGSTVAEIAQLLLRNGAAAVQVWCLCRTL..... 227
NTDB id 1458 ASU2 RS11390 WP 005624668.1 RNLKGAFRYTP...NRAYHRVAIIDDVVTTGSTLNAICAELLKQGVQHIQVWTLARA..... 225
NTDB id 1374 NTHI RS02640 WP 011272045.1 QNLKNAFSLVVSKNEFPYRRVALVDDVITTGSTLNEISKLLRKLGVEEIQVWGLARA..... 228
NTDB id 1348 HI 0434 AAC22093.1 QNLKNAFSLAVSKNEFPYRRVALVDDVITTGSTLNEISKLLRKLGVEEIQVWGLARA..... 229
consensus
X non conserved
X similar
X ≥ 50% conserved