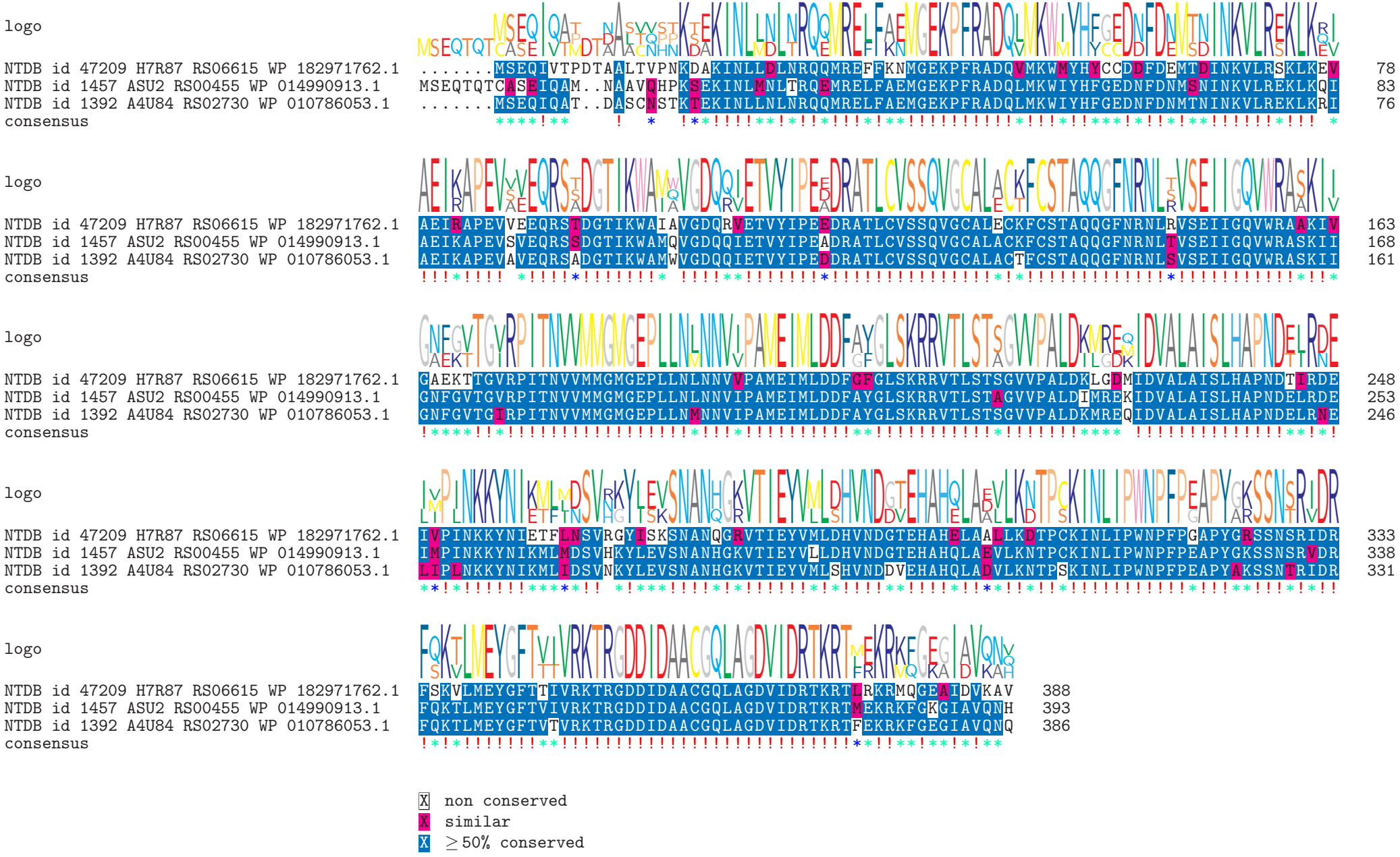
logo
NTDB id 47209 H7R87 RS06615 WP 182971762.1 .......MSEQIVTPDTAALTVPNKDAKINLLDLNRQQMREFFKNMGEKPFRADQVMKWMYHYCCDDFDEMTDINKVLRSKLKEV 78
NTDB id 1457 ASU2 RS00455 WP 014990913.1 MSEQTQTCASEIQAM..NAAVQHPKSEKINLMNLTRQEMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMSNINKVLREKLKQI 83
NTDB id 1392 A4U84 RS02730 WP 010786053.1 .......MSEQIQAT..DASCNSTKTEKINLLNLNRQQMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMTNINKVLREKLKRI 76
consensus
logo
NTDB id 47209 H7R87 RS06615 WP 182971762.1 AEIRAPEVVEEQRSTDGTIKWAIAVGDQRVETVYIPEEDRATLCVSSQVGCALECKFCSTAQQGFNRNLRVSEIIGQVWRAAKIV 163
NTDB id 1457 ASU2 RS00455 WP 014990913.1 AEIKAPEVSVEQRSSDGTIKWAMQVGDQQIETVYIPEADRATLCVSSQVGCALACKFCSTAQQGFNRNLTVSEIIGQVWRASKII 168
NTDB id 1392 A4U84 RS02730 WP 010786053.1 AEIKAPEVAVEQRSADGTIKWAMWVGDQQIETVYIPEDDRATLCVSSQVGCALACTFCSTAQQGFNRNLSVSEIIGQVWRASKII 161
consensus
logo
NTDB id 47209 H7R87 RS06615 WP 182971762.1 GAEKTGVRPITNVVMMGMGEPLLNLNNVVPAMEIMLDDFGFGLSKRRVTLSTSGVVPALDKLGDMIDVALAISLHAPNDTIRDE 248
NTDB id 1457 ASU2 RS00455 WP 014990913.1 GNFGVTGVRPITNVVMMGMGEPLLNLNNVIPAMEIMLDDFAYGLSKRRVTLSTAGVVPALDIMREKIDVALAISLHAPNDELRDE 253
NTDB id 1392 A4U84 RS02730 WP 010786053.1 GNFGVTGIRPITNVVMMGMGEPLLNMNNVIPAMEIMLDDFAYGLSKRRVTLSTSGVVPALDKMREQIDVALAISLHAPNDELRNE 246
consensus
logo
NTDB id 47209 H7R87 RS06615 WP 182971762.1 IVPINKKYNIETFLNSVRGYISKSNANQGRVTIEYVMLDHVNDGTEHAHELAALLKDTPCKINLIPWNPFPGAPYGRSSNSRIDR 333
NTDB id 1457 ASU2 RS00455 WP 014990913.1 IMPINKKYNIKMLMDSVHKYLEVSNANHGKVTIEYVLLDHVNDGTEHAHQLAEVLKNTPCKINLIPWNPFPEAPYGKSSNSRVDR 338
NTDB id 1392 A4U84 RS02730 WP 010786053.1 LIPLNKKYNIKMLIDSVNKYLEVSNANHGKVTIEYVMLSHVNDDVEHAHQLADVLKNTPSKINLIPWNPFPEAPYAKSSNTRIDR 331
consensus
logo
NTDB id 47209 H7R87 RS06615 WP 182971762.1 FSKVLMEYGFTTIVRKTRGDDIDAACGQLAGDVIDRTKRTLRKRMQGEAIDVKAV 388
NTDB id 1457 ASU2 RS00455 WP 014990913.1 FQKTLMEYGFTVIVRKTRGDDIDAACGQLAGDVIDRTKRTMEKRKFGKGIAVQNH 393
NTDB id 1392 A4U84 RS02730 WP 010786053.1 FQKTLMEYGFTVTVRKTRGDDIDAACGQLAGDVIDRTKRTFEKRKFGEGIAVQNQ 386
consensus
X non conserved
X similar
X ≥ 50% conserved