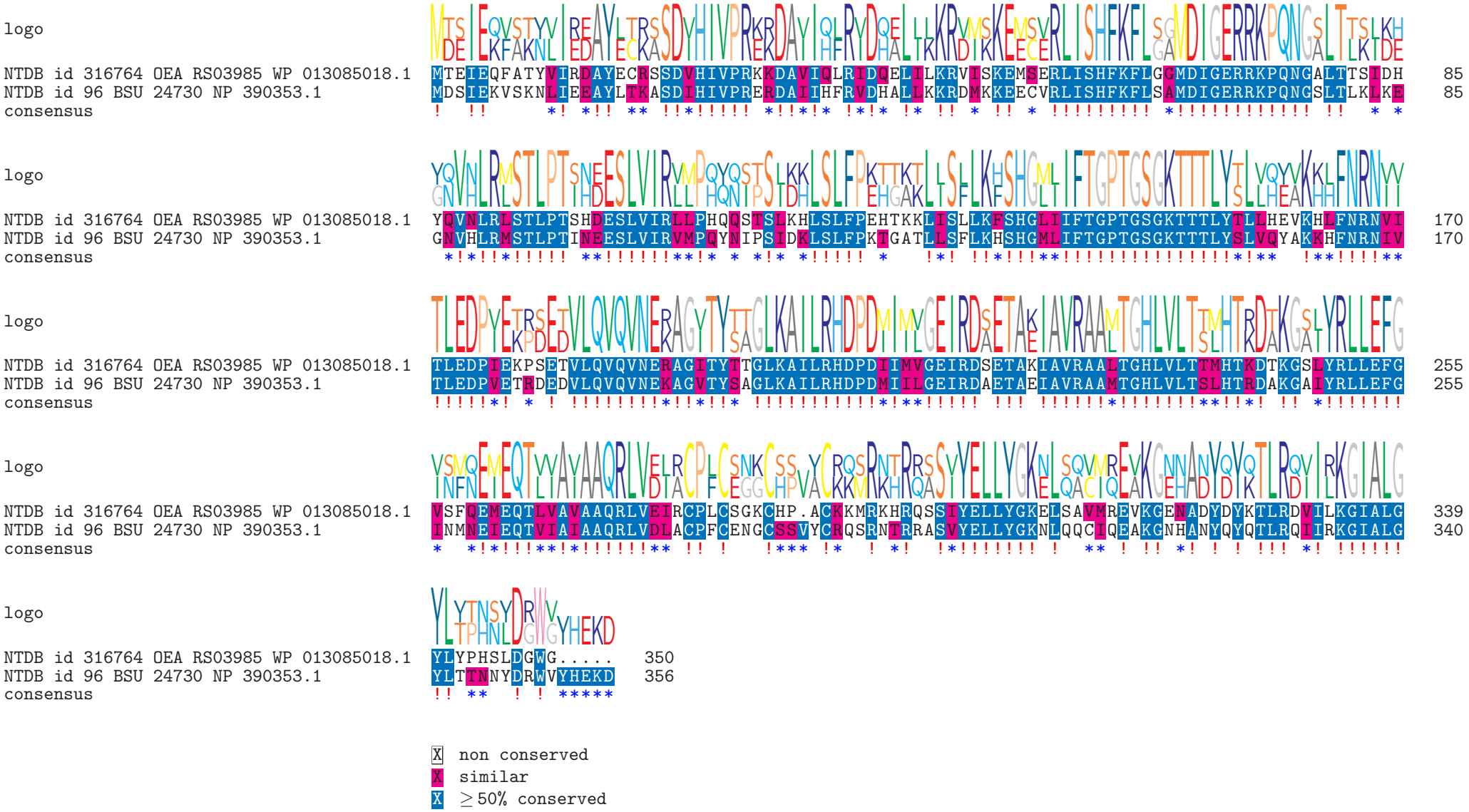

logo
NTDB id 316764 OEA RS03985 WP 013085018.1 MTEIEQFATYVIRDAYECRSSDVHIVPRKKDAVIQLRIDQELILKRVISKEMSERLISHFKFLGGMDIGERRKPQNGALTTSIDH 85
NTDB id 96 BSU 24730 NP 390353.1 MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLTLKLKE 85
consensus
logo
NTDB id 316764 OEA RS03985 WP 013085018.1 YQVNLRLSTLPTSHDESLVIRLLPHQQSTSLKHLSLFPEHTKKLISLLKFSHGLIIFTGPTGSGKTTTLYTLLHEVKHLFNRNVI 170
NTDB id 96 BSU 24730 NP 390353.1 GNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQYAKKHFNRNIV 170
consensus
logo
NTDB id 316764 OEA RS03985 WP 013085018.1 TLEDPIEKPSETVLQVQVNERAGITYTTGLKAILRHDPDIIMVGEIRDSETAKIAVRAALTGHLVLTTMHTKDTKGSLYRLLEFG 255
NTDB id 96 BSU 24730 NP 390353.1 TLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLHTRDAKGAIYRLLEFG 255
consensus
logo
NTDB id 316764 OEA RS03985 WP 013085018.1 VSFQEMEQTLVAVAAQRLVEIRCPLCSGKCHP.ACKKMRKHRQSSIYELLYGKELSAVMREVKGENADYDYKTLRDVILKGIALG 339
NTDB id 96 BSU 24730 NP 390353.1 INMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGNHANYQYQTLRQIIRKGIALG 340
consensus
logo
NTDB id 316764 OEA RS03985 WP 013085018.1 YLYPHSLDGWG..... 350
NTDB id 96 BSU 24730 NP 390353.1 YLTTNNYDRWVYHEKD 356
consensus
X non conserved
X similar
X ≥ 50% conserved