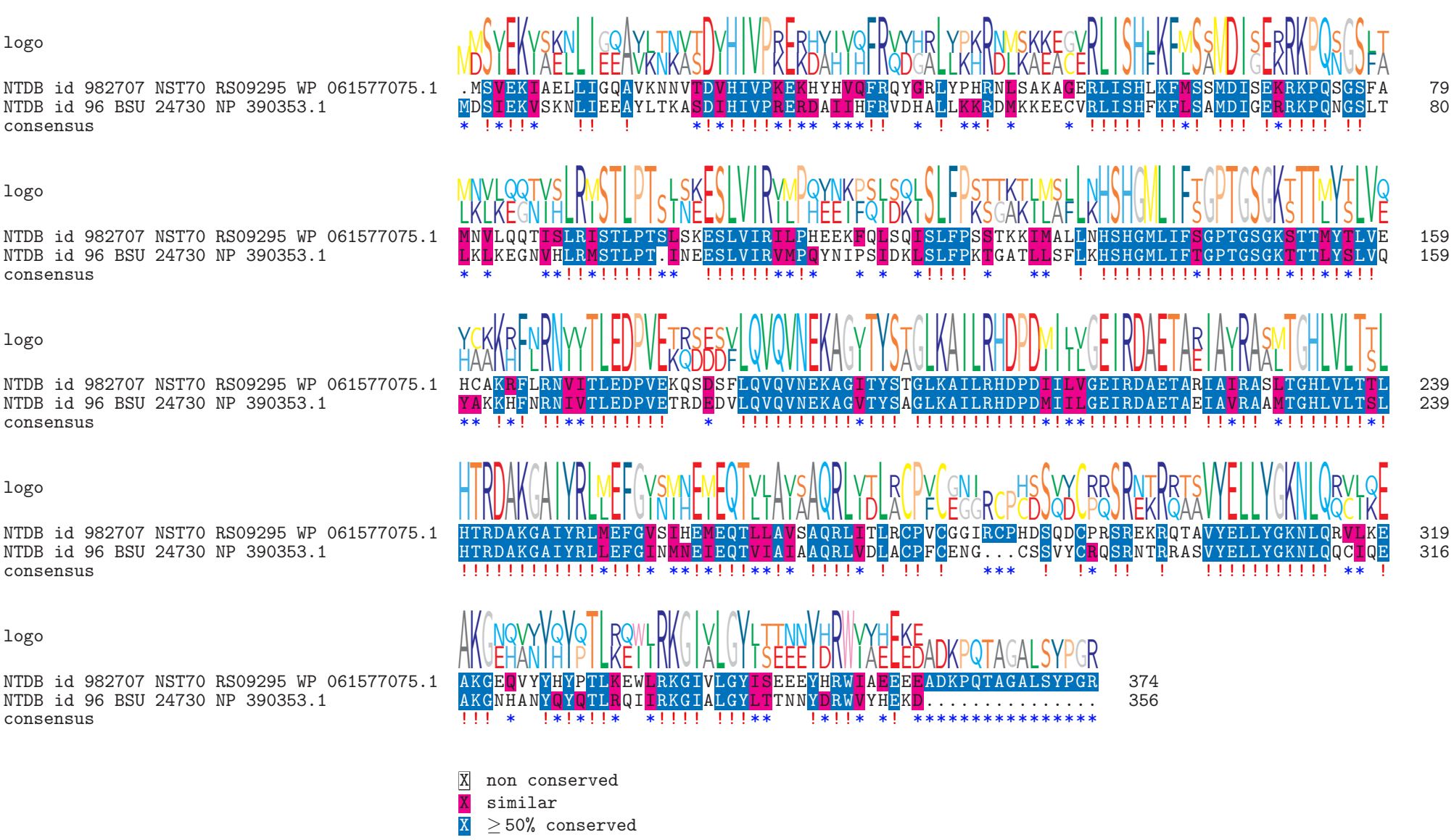
logo
NTDB id 982707 NST70 RS09295 WP 061577075.1 .MSVEKIAELLIGQAVKNNVTDVHIVPKEKHYHVQFRQYGRLYPHRNLSAKAGERLISHLKFMSSMDISEKRKPQSGSFA 79
NTDB id 96 BSU 24730 NP 390353.1 MDSTEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT 80
consensus
logo
NTDB id 982707 NST70 RS09295 WP 061577075.1 MNVLQQTISLRISTLPTSLSKESLVIRILPHEEKFQLSQISLFPSSTKKIMALLNHSHGMLIFSGPTGSGKSTTMYTLVE 159
NTDB id 96 BSU 24730 NP 390353.1 LKLKEGNVHLRMSTLPT.INEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQ 159
consensus
logo
NTDB id 982707 NST70 RS09295 WP 061577075.1 HCAKRFLRNVITLEDPVEKQSDSFLQVQVNEKAGITYSTGLKAILRHDPDIILVGEIRDAETARIAIRASLTGHLVLTTL 239
NTDB id 96 BSU 24730 NP 390353.1 YAKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSL 239
consensus
logo
NTDB id 982707 NST70 RS09295 WP 061577075.1 HTRDAKGAIYRLMEFGVSIHEMEQTLLAVSAQRLITLRCPVCGGIRCPHDSQDCPRSREKRQTAVYELLYGKNLQRVLKE 319
NTDB id 96 BSU 24730 NP 390353.1 HTRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENG...CSSVYCRQSRNTRRASVYELLYGKNLQQCIQE 316
consensus
logo
NTDB id 982707 NST70 RS09295 WP 061577075.1 AKGEQVYYHYPTLKEWLRKGIVLGYISEEEYHRWIAEEEEADKPQTAGALSYPGR 374
NTDB id 96 BSU 24730 NP 390353.1 AKGNHANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD..... 356
consensus
X non conserved
X similar
X ≥ 50% conserved