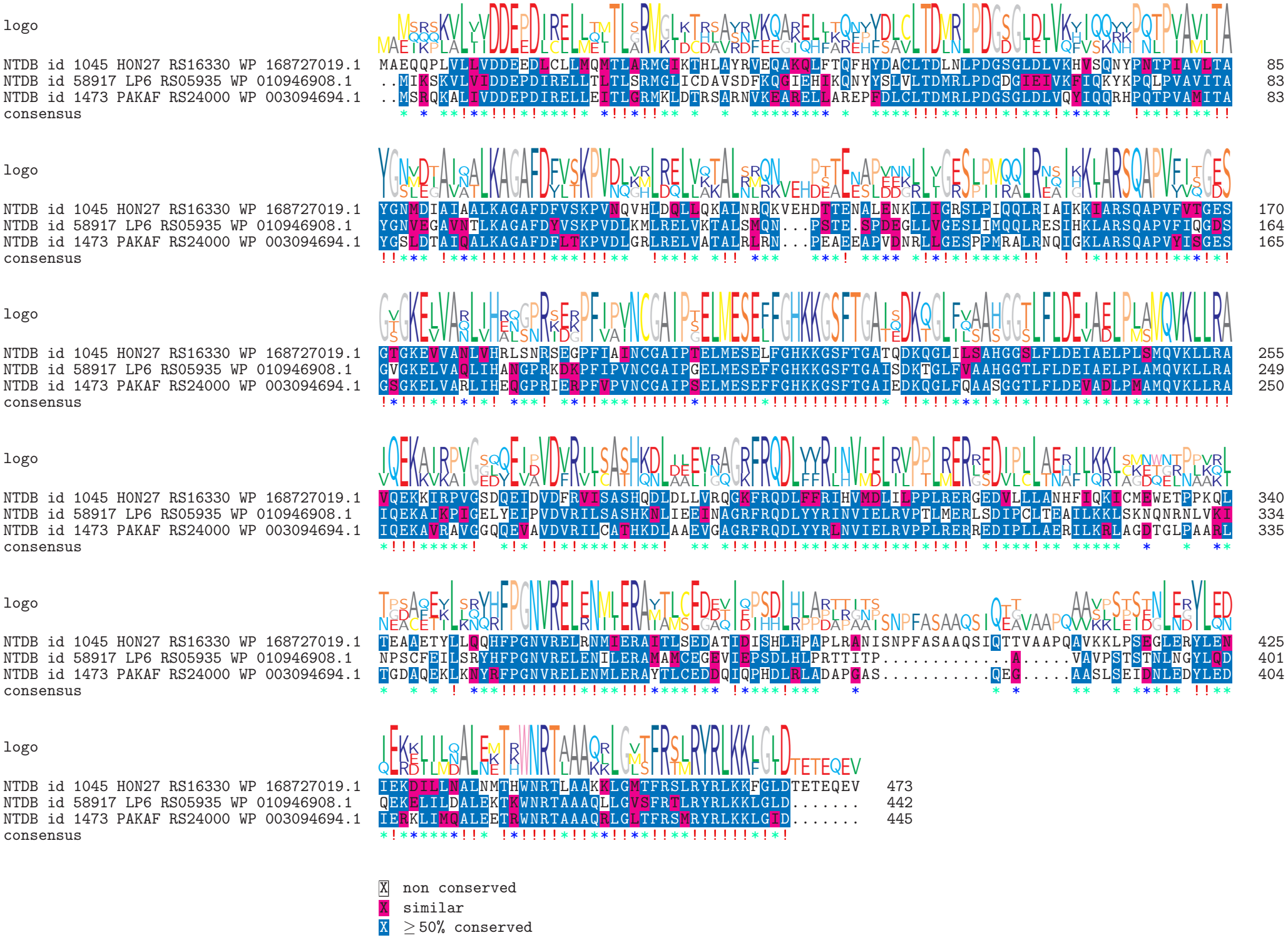
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 MAEQQPLVLLVDDEEDLCLLMQMTLARMGIKTHLAYRVEQAKQLFTQFHYDACLTDLNLPDGSGLDLVKHVSQNYPNTPIAVLTA 85
NTDB id 58917 LP6 RS05935 WP 010946908.1 ..MIKSKVLVIDDEPDIRELLTLTLSRMGLICDAVSDFKQGIEHIKQNYYSLVLTDMRLPDGDGIEIVKFIQKYKPQLPVAVITA 83
NTDB id 1473 PAKAF RS24000 WP 003094694.1 ..MSRQKALIVDDEPDIRELLEITLGRMKLDTRSARNVKEARELLAREPFDLCLTDMRLPDGSGLDLVQYIQQRHPQTPVAMITA 83
consensus
NTDB id 1045 H0N27 RS16330 WP 168727019.1 YGNMDIAIAALKAGAFDFVSKPVNQVHLDQLLQKALNRQKVEHDTTENALENKLLIGRSLPIQQLRIAIKKIARSQAPVFVTGES 170
NTDB id 58917 LP6 RS05935 WP 010946908.1 YGNVEGAVNTLKAGAFDYVSKPVDLKMLRELVKTALSMQN...PSTE.SPDEGLLVGESLIMQQLRESIHKLARSQAPVFIQGDS 164
NTDB id 1473 PAKAF RS24000 WP 003094694.1 YGSLDTAIQALKAGAFDFLTKPVDLGRLRELVATALRLRN...PEAEEAPVDNRLLGESPPMRALRNQIGKLARSQAPVYISGES 165
consensus
NTDB id 1045 H0N27 RS16330 WP 168727019.1 GTGKEVVANLVHRLSNRSEGPFIAINCGAIPTELMESELFGHKKGSFTGATQDKQGLILSAHGGSLFLDEIAELPLSMQVKLLRA 255
NTDB id 58917 LP6 RS05935 WP 010946908.1 GVGKELVAQLIHANGPRKDKPFIPVNCGAIPGELMESEFFGHKKGSFTGAISDKTGLFVAAHGGTLFLDEIAELPLAMQVKLLRA 249
NTDB id 1473 PAKAF RS24000 WP 003094694.1 GSGKELVARLIHEQGPRIERPFVPVNCGAIPSELMESEFFGHKKGSFTGAIEDKQGLFQAASGGTLFLDEVADLPMAMQVKLLRA 250
consensus
NTDB id 1045 H0N27 RS16330 WP 168727019.1 VQEKKIRPVGSDQEIDVDFRVISASHQDLDLLVRQGKFRQDLFFRIHVMDLILPPLRERGEDVLLLANHFIQQICMEWETPPKQL 340
NTDB id 58917 LP6 RS05935 WP 010946908.1 IQEKAIKPIGELYEIPVDVRILSASHKNLIEEINAGRFRQDLYYRINVIELRVPTLMERLSDIPCLTEAILKKLSKNQNRNLVKI 334
NTDB id 1473 PAKAF RS24000 WP 003094694.1 IQEKAVRAVGGQQEVAVDVRILCATHKDLAAEVGAGRFRQDLYYRLNVIELRVPPLRERREDIPLLAERILKRLAGDTGLPAARL 335
consensus
NTDB id 1045 H0N27 RS16330 WP 168727019.1 TEAAETYLLQQHFPGNVRELRNMIERAITLSEDATIDISHLHPAPLRANISNPFASAAQSIQTTVAAPQAVKKLPSEGLERYLEN 425
NTDB id 58917 LP6 RS05935 WP 010946908.1 NPSCFEILSRYHFPGNVRELENILERAMAMCEGEVIEPSDLHLPRTTITP.............A.....VAVPSTSTNLNGYLQD 401
NTDB id 1473 PAKAF RS24000 WP 003094694.1 TGDAQEKLKNYRFPGNVRELENMLERAYTLCEDDQIQPHDLRLADAPGAS...........QEG.....AASLSEIDNLEDYLED 404
consensus
NTDB id 1045 H0N27 RS16330 WP 168727019.1 IEKDILLNALNMTHWNRTLAAKKLGMTFRSLRYRLKKFGLDTETEQEV 473
NTDB id 58917 LP6 RS05935 WP 010946908.1 QEKELILDALEKTKWNRTAAAQLLGVSFRTLRYRLKKLGLD..... 442
NTDB id 1473 PAKAF RS24000 WP 003094694.1 IERKLIMQALEETRWNRTAAAQRLGLTFRSMRYRLKKLGID..... 445
consensus
X non conserved
X similar
X ≥ 50% conserved