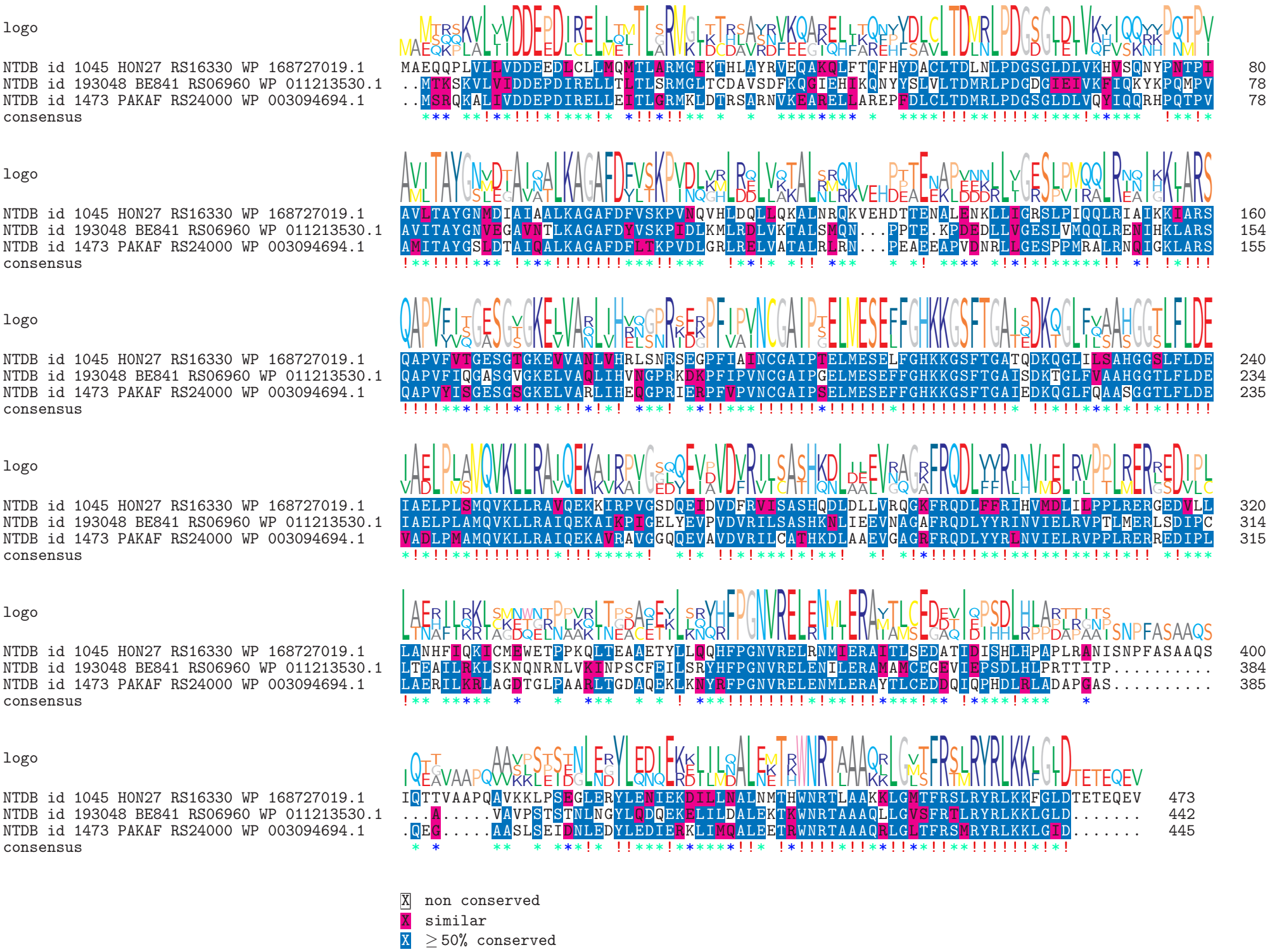

logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 MAEQQPLVLLVDDEEDLCLLMQMTLARMGIKTHLAYRVEQAKQLFTQFHYDACLTDLNLPDGSGLDLVKHVSQNYPNTPI 80
NTDB id 193048 BE841 RS06960 WP 011213530.1 ..MTKSKVLVIDDEPDIRELLTLTLSRMGLTCDAVSDFKQGIEHIKQNYYSLVLTDMRLPDGDGIEIVKFIQKYKPQMPV 78
NTDB id 1473 PAKAF RS24000 WP 003094694.1 ..MSRQKALIVDDEPDIRELLEITLGRMKLDTRSARNVKEARELLAREPFDLCLTDMRLPDGSGLDLVQYIQQRHPQTPV 78
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 AVLTAYGNMDIAIAALKAGAFDFVSKPVNQVHLDQLLQKALNRQKVEHDTTENALENKLLIGRSLPIQQLRIAIKKIARS 160
NTDB id 193048 BE841 RS06960 WP 011213530.1 AVITAYGNVEGAVNTLKAGAFDYVSKPIDLKMLRDLVKTALSMQN...PPTE.KPDEDLLVGESLVMQQLRENIHKLARS 154
NTDB id 1473 PAKAF RS24000 WP 003094694.1 AMITAYGSLDTAIQALKAGAFDFLTKPVDLGRLRELVATALRLRN...PEAEEAPVDNRLLGESPPMRALRNQIGKLARS 155
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 QAPVFVTGESGTGKELVANLVHRLSNRSEGPFIAINCGAIPTELMESELFGHKKGSFTGATQDKQGLILSAHGGSLFLDE 240
NTDB id 193048 BE841 RS06960 WP 011213530.1 QAPVFIQGASGVGKELVAQLIHVNGPRKDKPFIPVNCGAIPGELMESEFFGHKKGSFTGAISDKTGLFVAAHGGTLFLDE 234
NTDB id 1473 PAKAF RS24000 WP 003094694.1 QAPVYISGESGSGKELVARLIHEQGPRIERPFVPVNCGAIPSELMESEFFGHKKGSFTGAIEDKQGLFQAASGGTLFLDE 235
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 IAELPLSMQVKLLRAVQEKKIRPVGSDQEIDVDFRVISASHQDLDLLVRQGKFRQDLFFRIHVMDLILPPLRERGEDVLL 320
NTDB id 193048 BE841 RS06960 WP 011213530.1 IAELPLAMQVKLLRAIQEKAIKPIGELYEVPVDVRILSASHKNLIEEVNAGAFRQDLYYRINVIELRVPTLMERLSDIPC 314
NTDB id 1473 PAKAF RS24000 WP 003094694.1 VADLPMAMQVKLLRAIQEKAVRAVGGQQEVAVDVRILCATHKDLAAEVGAGRFRQDLYYRLNVIELRVPPLRERREDIPL 315
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 LANHFIQKICMEWETPPKQLTEAAETYLLQQHFPGNVRELRNMIERAITLSEDATIDISHLHPAPLRANISNPFASAAQS 400
NTDB id 193048 BE841 RS06960 WP 011213530.1 LTEAILRKLSKNQNRNLVKINPSCFEILSRYHFPGNVRELENILERAMAMCEGEVIEPSDLHLPRTTITP..... 384
NTDB id 1473 PAKAF RS24000 WP 003094694.1 LAERILKRLAGDTGLPAARLTGDAQEKLKNYRFPGNVRELENMLERAYTLCEDDQIQPHDLRLADAPGAS..... 385
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 IQTTVAAPQAVKKLPSEGLERYLENIEKDILLNALNMTHWNRTLAAKKLGMTFRSLRYRLKKFGLDTETEQEV 473
NTDB id 193048 BE841 RS06960 WP 011213530.1 ...A.....VAVPSTSTNLNGYLQDQEKELILDALEKTKWNRTAAAQLLGVSFRTLRYRLKKLGLD..... 442
NTDB id 1473 PAKAF RS24000 WP 003094694.1 .QEG.....AASLSEIDNLEDYLEDIERKLIMQALEETRWNRTAAAQRLGLTFRSMRYRLKKLGID..... 445
consensus
X non conserved
X similar
X ≥ 50% conserved