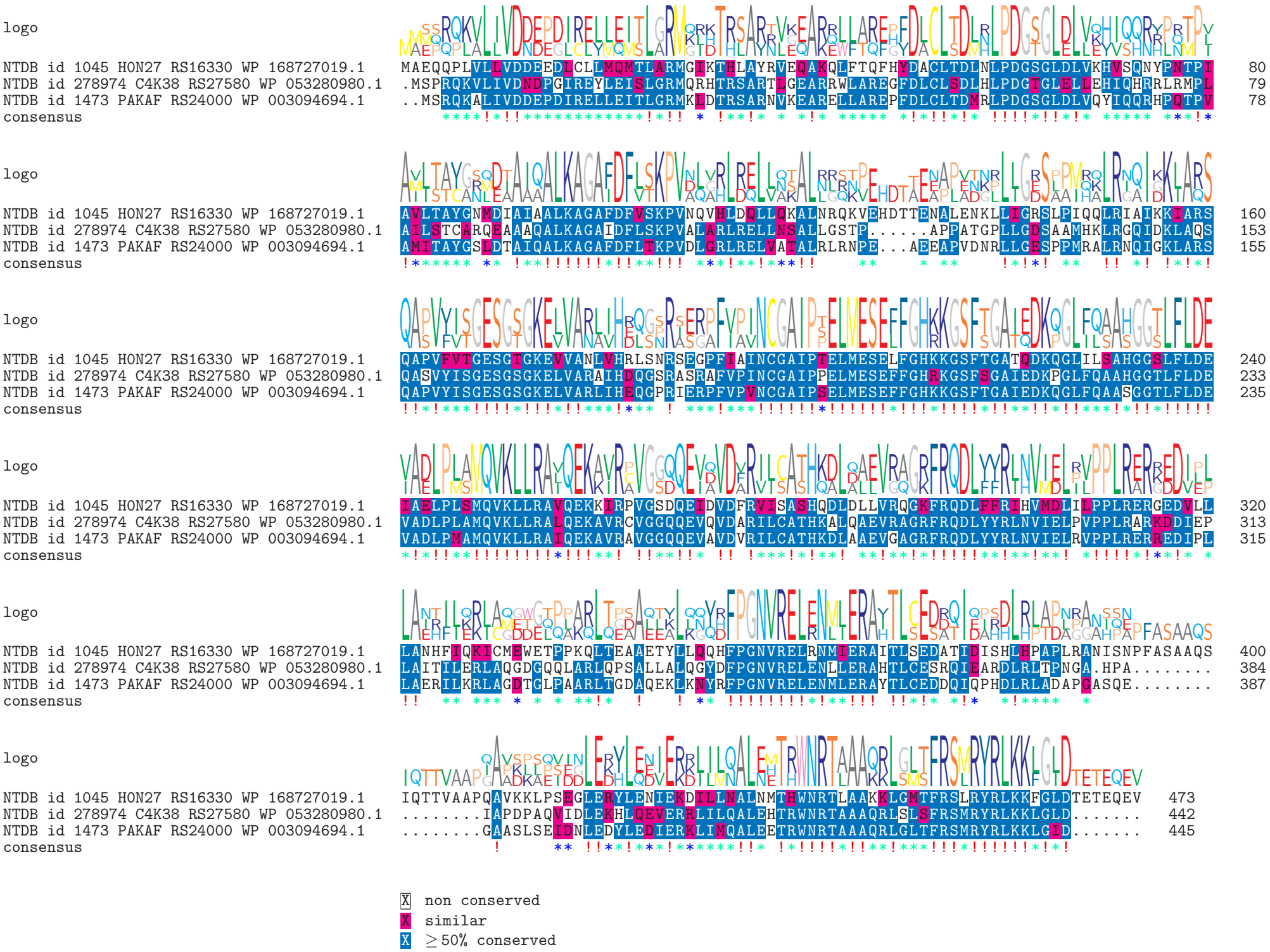
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 MAEQQPLVLLVDDEEDLCLLMQMTLARMGIKTHLAYRVEQAKQLFTQFHYDACLTDLNLPDGSGLDLVKHVSQNYPNTPI 80
NTDB id 278974 C4K38 RS27580 WP 053280980.1 .MSPRQKVLIVDNDPGIREYLEISLGRMQRHTRSARTLGEARRWLAREGFDLCLSDLHLPDGTGLELLEHIQHRRLRMPL 79
NTDB id 1473 PAKAF RS24000 WP 003094694.1 ..MSRQKALIVDDEPDIRELLEITLGRMKLDTRSARNVKEARELLAREPFDLCLTDMRLPDGSGLDLVQYIQQRHPQTPV 78
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 AVLTAYGNMDIAIAALKAGAFDFVSKPVNQVHLDQLLQKALNRQKVEHDTTENALENKLLIGRSLPIQQLRIAIKKIARS 160
NTDB id 278974 C4K38 RS27580 WP 053280980.1 AILSTCARQEAAAQALKAGAIDFLSKPVALARLRELLNSALLGSTP.....APPATGPLLGDSAAMHKLRGQIDKLAQS 153
NTDB id 1473 PAKAF RS24000 WP 003094694.1 AMITAYGSLDTAIQALKAGAFDFLTKPVDLGRLRELVATALRLRNPE...AEEAPVDNRLLGESPPMRALRNQIGKLARS 155
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 QAPVFVTGESGTGKELVANLVHRLSNRSEGPFIAINCGAIPTELMESELFGHKKGSFTGATQDKQGLILSAHGGSLFLDE 240
NTDB id 278974 C4K38 RS27580 WP 053280980.1 QASVYISGESGSGKELVARAIHDQGSRASRAFVPINCGAIPPELMESEFFGHRKGSFSGAIEDKPGLFQAAHGGTLFLDE 233
NTDB id 1473 PAKAF RS24000 WP 003094694.1 QAPVYISGESGSGKELVARLIHRQGPRIERPFVPVNCGAIPSELMESEFFGHKKGSFTGAIEDKQGLFQAASGGTLFLDE 235
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 IAELPLSMQVKLLRAVQEKKIRPVGSDQEIDVDFRVISASHQDLDLLVRQGKFRQDLFFRIHVMDLILPPLRERGEDVLL 320
NTDB id 278974 C4K38 RS27580 WP 053280980.1 VADLPLAMQVKLLRALQEKAVRCVGGQQEVQVDARILCATHKALQAEVRAGRFRQDLYYRLNVIELPVPPLRARKDDIEP 313
NTDB id 1473 PAKAF RS24000 WP 003094694.1 VADLPMAMQVKLLRAIQEKAVRAVGGQQEVAVDVRILCATHKDLAAEVGAGRFRQDLYYRLNVIELRVPPLRERREDIPL 315
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 LANHFIQKICMEWETPPKQLTEAAETYLLQQHFPGNVRELRNMIERAITLSEDATIDISHLHPAPLRANISNPFASAAQS 400
NTDB id 278974 C4K38 RS27580 WP 053280980.1 LAITILERLAQGDGQQLARLQPSALLALQGYDFPGNVRELENLLERAHTLCESRQIEARDLRLTPNGA.HPA..... 384
NTDB id 1473 PAKAF RS24000 WP 003094694.1 LAERILKRLAGDTGLPAARLTGDAQEKLKNYRFPGNVRELENMLERAYTLCEDDQIQPHDLRLADAPGASQE..... 387
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 IQTTVAAPQAVKKLPSEGLERYLENIEKDILLNALNMTHWNRTLAAKKLGMTFRSLRYRLKKFGLDTETEQEV 473
NTDB id 278974 C4K38 RS27580 WP 053280980.1 .....IAPDPAQVIDLEKHLQEVERRLILQALEHTRWNRTAAAQRLSLSFRSMRYRLKKLGLD..... 442
NTDB id 1473 PAKAF RS24000 WP 003094694.1 .....GAASLSEIDNLEDYLEDIERKLIMQALEETRWNRTAAAQRLGLTFRSMRYRLKKLGID..... 445
consensus
X non conserved
X similar
X ≥ 50% conserved