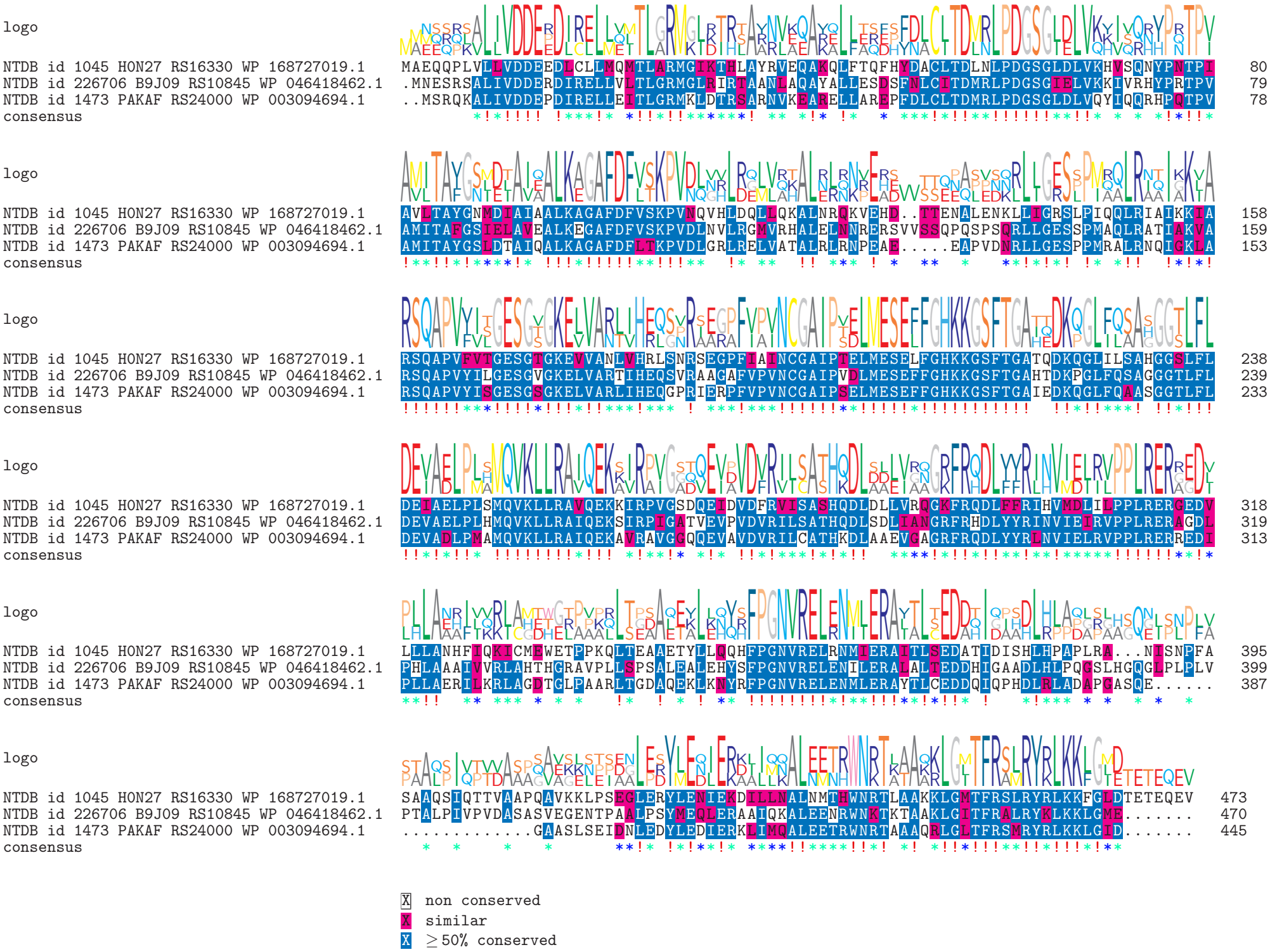
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 MAEQQPLVLLLVDDEEDLCLLMQMTLARMGIKTHLAYRVEQAKQLFTQFHYDACLTDLNLPDGSGLDLVKHVSQNYPNTPI 80
NTDB id 226706 B9J09 RS10845 WP 046418462.1 .MNESRSALIVDDERDIRELLVLTLGRMGLRIRTAANLAQAYALLESDSFNLCITDMRLPDGSGIELVKKIVRHYPRTPV 79
NTDB id 1473 PAKAF RS24000 WP 003094694.1 ..MSRQKALIVDDEPDIRELLEITLGRMKLDTRSARNVKEARELLAREPFDLCLTDMRLPDGSGLDLVQYIQQRHPQTPV 78
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 AVLTAYGNMDIAIAALKAGAFDFVSKPVNQVHLDQLLQKALNRQKVEHD..TTENALENKLLIGRSLPIQQLRIAIKKIA 158
NTDB id 226706 B9J09 RS10845 WP 046418462.1 AMITAFGSIELAVEALKEGAFDFVSKPVDLNVLRGMVRHALELNNRERSVVSSQPQSPSQRLLGESSPMAQLRATIAKVA 159
NTDB id 1473 PAKAF RS24000 WP 003094694.1 AMITAYGSLDTAIQALKAGAFDFLTKPVDLGRLRELVATALRLRNPEAE.....EAPVDNRLLGESPPMRALRNQIGKLA 153
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 RSQAPVFVTGESGTGKEVVANLVHRLSNRSEGPFIAINCGAIPTELMESELFGHKKGSFTGATQDKQGLILSAHGGSLFL 238
NTDB id 226706 B9J09 RS10845 WP 046418462.1 RSQAPVYILGESGVGKELVARTIHEQSVRAAGAFVPVNCGAIPVDLMESEFFGHKKGSFTGAHTDKPGLFQSAGGGTLFL 239
NTDB id 1473 PAKAF RS24000 WP 003094694.1 RSQAPVYISGESGSGKELVARLIHEQGPRIERPFVPVNCGAIPSELMESEFFGHKKGSFTGAIEDKQGLFQAASGGTLFL 233
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 DEIAELPLSMQVKLLRAVQEKKIRPVGSDQEIDVDFRVISASHQDLDLLVRQGKFRQDLFFRIHVMDLILPPLRERGEDV 318
NTDB id 226706 B9J09 RS10845 WP 046418462.1 DEVAELPLHMQVKLLRAIQEKSIRPIGATVEVPVDVRILSATHQDLSDLIANGRFRHDLYYRINVIEIRVPPLRERAGDL 319
NTDB id 1473 PAKAF RS24000 WP 003094694.1 DEVADLPMAMQVKLLRAIQEKAVRAVGGQQEVAVDVRILCATHKDLAAEVGAGRFRQDLYYRLNVIELRVPPLRERREDI 313
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 LLLANHFIQKICMEWETPPKQLTEAAETYLLQQHFPGNVRELRNMIERAITLSEDATIDISHLHPAPLRA...NISNPFA 395
NTDB id 226706 B9J09 RS10845 WP 046418462.1 PHLAAAIVVRLAHTHGRAVPLLSPSALEALEHYSFPGNVRELENILERALALTEDDHIGAADLHLPQGSLHGQGLPLPLV 399
NTDB id 1473 PAKAF RS24000 WP 003094694.1 PLLAERILKRLAGDTGLPAARLTGDAQEKLKNYRFPGNVRELENMLERAYTLCEDDQIQPHDLRLADAPGASQE..... 387
consensus
logo
NTDB id 1045 H0N27 RS16330 WP 168727019.1 SAAQSIQTTVAAPQAVKKLPSEGLERYLENIEKDILLNALNMTHWNRTLAAKKLGMTFRSLRYRLKKFGLDTETEQEV 473
NTDB id 226706 B9J09 RS10845 WP 046418462.1 PTALPIVPVDASASVEGENTPAALPSYMEQLERAAIQKALEENRWNKTKTAAKLGITFRALRYKLKKLGME..... 470
NTDB id 1473 PAKAF RS24000 WP 003094694.1 .....GAASLSEIDNLEDYLEDIERKLIMQALEETRWNRTAAAQRLGLTFRSMRYRLKKLGID..... 445
consensus
X non conserved
X similar
X ≥ 50% conserved