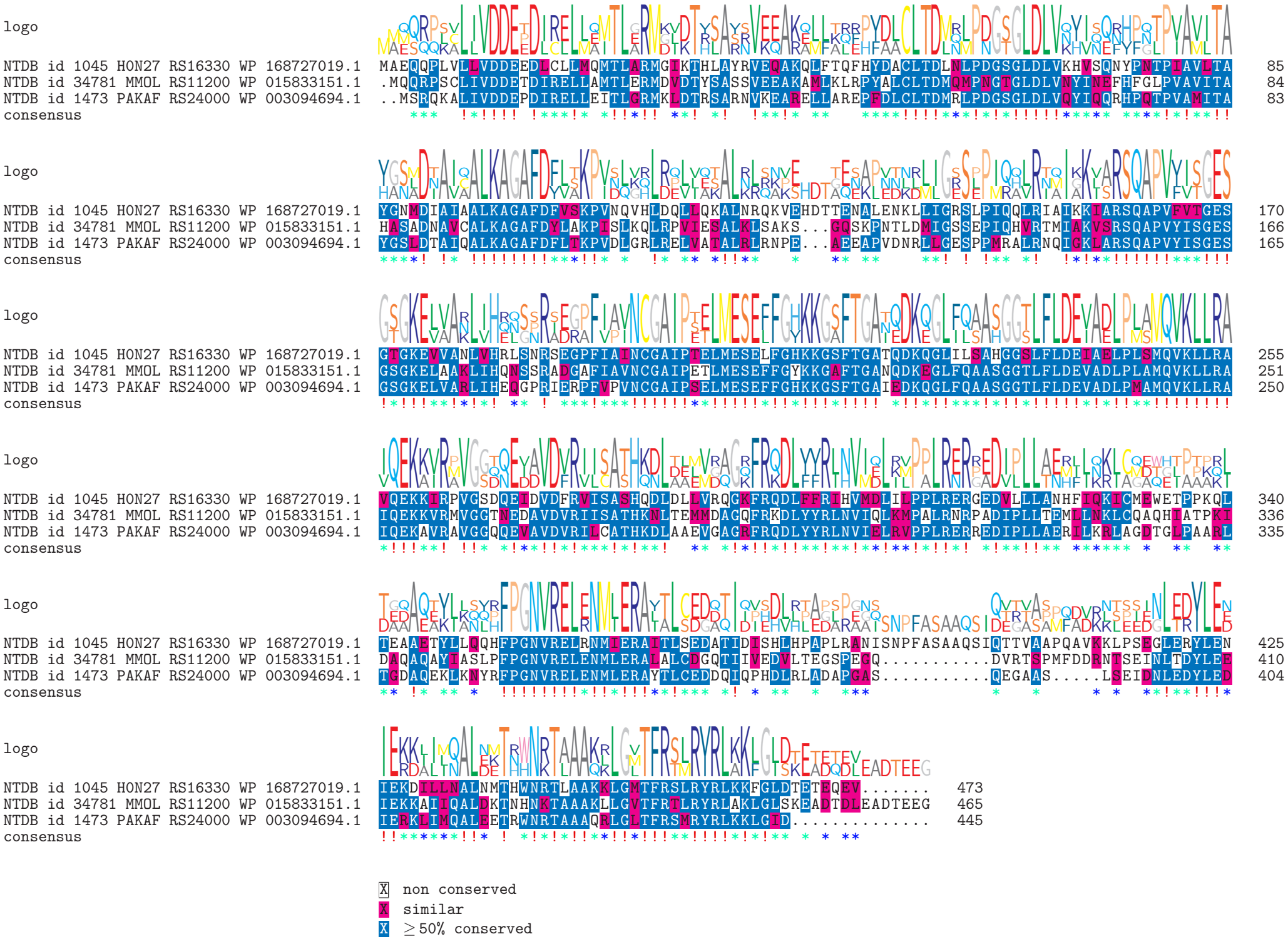
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NTDB id 1045 HON27 RS16330 WP 168727019.1 MAEQQPLVLLVDDEEDLCLLMQMTLARMGIKTHLAYRVEQAKQLFTQFHYDACLTDLNLPDGSGLDLVKHVSQNYPNTPIAVLTA 85
NTDB id 34781 MMOL RS11200 WP 015833151.1 .MQQRPSCLIVDDETDIRELLAMTLERMDVDTYSASSVEEAKAMLKLRPYALCLTDMQMPNGTGLDLVNYINEFHFGLPVAVITA 84
NTDB id 1473 PAKAF RS24000 WP 003094694.1 ..MSRQKALIVDDEPDIRELLEITLGRMKLDTRSARNVKEARELLAREPFDLCLTDMRLPDGSGLDLVQYIQQRHPQTPVAMITA 83
consensus *** !*!!!! !***!* *!!*!! **! *! !**!* ** ****!!!!*** !*!*!!!!!***** ****!*!**!!
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 YGNMDIAIAALKAGAFDFVSKPVNQVHLDQLLQKALNRQKVEHDTTENALENKLLICRSLPIQQLRIAIKKIARSQAPVFVTGES 170
NTDB id 34781 MMOL RS11200 WP 015833151.1 HASADNAVCALKAGAFDYLAKPISLKQLRPVIESALKLSAKS...GQSKPNTLDMICSSEPIQHVRTMIAKVSRSQAPVYISGES 166
NTDB id 1473 PAKAF RS24000 WP 003094694.1 YGSLDTAIQALKAGAFDFLTKPVDLGRLRELVATALRLRNPE...AEEAPVDNRLLGESPPMRALRNQIGKLARSQAPVYISGES 165
consensus ***** !* !!!!!!!!***!!* * !* ** *!!** * ** ** **! ! !** *! !*!**!!!!!!***!!!
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 GTGKEVVANLVHRLSNRSEGPFIAINCGAIPTELMESELFGHKKGSFTGATQDKQGLILSAHGGSLFLDEIAELPLSMQVKLLRA 255
NTDB id 34781 MMOL RS11200 WP 015833151.1 GSGKELAAKLIHQNSSRADGAFIAVNCGAIPETLMESEFFGYKKGAFTGANQDKEGLFQAASGGTLFLDEVADLPLAMQVKLLRA 251
NTDB id 1473 PAKAF RS24000 WP 003094694.1 GSGKELVARLIHEQGPRIERPFVPVNCGAIPSELMESEFFGHKKGSFTGAIEDKQGLFQAASGGTLFLDEVADLPMAMQVKLLRA 250
consensus !*!!!**!*!*! ** ! ***!***!!!!!!*!!!!!*!!*!!!*!!!! *!!*!!***!*!!*!!!!!*!*!!**!!!!!!!!
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 VQEKKIRPVGSDQEIDVDFRVISASHQDLDLLVRQGKFRQDLFFRIHVMDLILPPLRERGEDVLLLANHFIQKICMEWETPPKQL 340
NTDB id 34781 MMOL RS11200 WP 015833151.1 IQEKKVRMVGGTNEDAVDVRIISATHKNLTEMMDAGQFRKDLYYRLNVIQLKMPALRNRPADIPLLTEMLLNKLCQAQHIATPKI 336
NTDB id 1473 PAKAF RS24000 WP 003094694.1 IQEKAVRAVGGQQEVAVDVRILCATHKDLAAEVGAGRFRQDLYYRLNVIELRVPPLRERREDIPLLAERILKRLAGDTGLPAARL 335
consensus *!!!**! !!* *!**!!*!***!*!**! ** *!*!!*!!**!**!**!**!*!!*! *!**!!** ****** * ** **
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 TEAAETYLLQQHFPGNVRELRNMIERAITLSEDATIDTSHLHPAPLRANISNPFASAAQSIQTTVAAPQAVKKLPSEGLERYLEN 425
NTDB id 34781 MMOL RS11200 WP 015833151.1 DAQAQAYIASLPFPGNVRELENMLERALALCDGQTIIVEDVLTEGSPEGQ.....DVRTSPMFDDRNTSEINLTDYLEE 410
NTDB id 1473 PAKAF RS24000 WP 003094694.1 TGDAQEKLKNYRFPGNVRELENMLERAYTLCEDDQIQPHDLRLADAPGAS.....QEGAASLSEIDNLEDYLED 404
consensus ** !* ** * !!!!!!!!*!!*!!!**!*** *! * ** * *** * * * * * **!**!!!*
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 IEKDTLLNALNMTHWNRTLAAKKLGMTFRSLRYRLKKFGLDTETEQEV..... 473
NTDB id 34781 MMOL RS11200 WP 015833151.1 IEKKAIIQALDKTNHNKTAAAKLLGVTFRTLRYRLARKLGLSKEADTDLEADTEEG 465
NTDB id 1473 PAKAF RS24000 WP 003094694.1 IERKLIMQALEETRWNRTAAAQRLGLTFRSMRYRLKKLGID..... 445
consensus !!******!!* ! *!*!*!!**!!*!!!**!!!!*!*!** * * **
X non conserved
X similar
X ≥ 50% conserved