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NTDB id 1045 HON27 RS16330 WP 168727019.1 MAEQQPLVLLVDDEEDLCLLMQMTLARMGIKTHLAYRVEQAKQLFTQFHYDACLTDLNLPDGSGLDLVKHVSQNYPNTPIAVLTA 85
NTDB id 67004 TAO RS02955 WP 096526546.1 .MRTSPLALIVDDEPDILALLELTLAPMGIRCQKAANLEQAQQCLQQQGFNLCLTDMRLPDGDGITLVKHINQAYPQLPVAVITA 84
NTDB id 1473 PAKAF RS24000 WP 003094694.1 ..MSRQKALIVDDEPDIRELLEITLGRMKLDTRSARNVKEARELLAREPFDLCLTDMRLPDGSGLDLVQYIQQRHPQTPVAMITA 83
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 YGNMDIAIAALKAGAFDFVSKPVNQVHLDQLLQKALNRQKVEHDTTENALENKLLIGRSLPIQQLRIAIKKIARSQAPVFVTGES 170
NTDB id 67004 TAO RS02955 WP 096526546.1 HGNVEAAVTALKSGAFDFISKPVDLGTLRNLVNTALKLPSTYP..QKERRTRHQLLGESPPMRVVRAQIAKLARSQAPVYISGES 167
NTDB id 1473 PAKAF RS24000 WP 003094694.1 YGSLDTAIQALKAGAFDFLTKPVDLGRLRELVATALRLRNPEA...EEAPVDNRLLGESPPMRALRNQIGKLARSQAPVYISGES 165
consensus
logo
NTDB id 1045 HON27 RS16330 WP 168727019.1 GTGKEVVANLVHRLSNRSEGPFIAINCGAIPTELMESELFGHKKGSFTGATQDKQGLILSAHGGSLFLDEIAELPLSMQVKLLRA 255
NTDB id 67004 TAO RS02955 WP 096526546.1 GTGKELVARLIHAQSPRSAGPFVPVNCGAIPDSLLESEFFGYEKGSFTGAHRDKAGLFQSSEGGTLFLDEVGDLSLAMQVKLLRA 252
NTDB id 1473 PAKAF RS24000 WP 003094694.1 GSGKELVARLIHEQGPRIERPFVPVNCGAIPSELMESEFFGHKKGSFTGAIEDKQGLFQAASGGTLFLDEVADLPMAMQVKLLRA 250
consensus
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NTDB id 1045 HON27 RS16330 WP 168727019.1 VQEKKIRPVGSDQEIDVDFRVISASHQDLDLLVRQGKFRQDLFFRIHVMDLILPPLRERGEDVLLLANHFIQKICMEWETPPKQL 340
NTDB id 67004 TAO RS02955 WP 096526546.1 IQERAVRPLGSASEISVDVRILSATHQDLALLVKRGEFRQDLFYRLNIIELPLPPLRARSTDIPALADHILTRLAATNQMICPQL 337
NTDB id 1473 PAKAF RS24000 WP 003094694.1 IQEKAVRAVGGQQEVAVDVRILCATHKDLAAEVGAGRFRQDLYYRLNVIELRVPPLRERREDIPLLAERILKRLAGDTGLPAARL 335
consensus
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NTDB id 1045 HON27 RS16330 WP 168727019.1 TEAAETYLLQQHFPGNVRELRNMIERAITLSEDATIDISHLHPAPLRANISNPFASAAQSIQTTVAAPQAVKKLPSEGLERYLEN 425
NTDB id 67004 TAO RS02955 WP 096526546.1 TPEALAKLQAYHFPGNVRELENILERALILREEELIQVEDLQLPVT.....SHGLSSTLTVDSLENRLES 402
NTDB id 1473 PAKAF RS24000 WP 003094694.1 TGDAQEKLKNYRFPGNVRELENMLERAYTLCEDDQIQPHDLRLADAPGAS.....QEGAASLSEIDNLEDYLED 404
consensus
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NTDB id 1045 HON27 RS16330 WP 168727019.1 IEKDTLLNALNMTHWNRTLAAKKLGMTFRSLRYRLKKFGLDTETEQEV 473
NTDB id 67004 TAO RS02955 WP 096526546.1 VEKQAILHALAQTGYNKTAAAKLLGITFRALRYRLKKLDLE..... 443
NTDB id 1473 PAKAF RS24000 WP 003094694.1 IERKLIMQALEETRWNRTAAAQRLGLTFRSMRYRLKKLGID..... 445
consensus
X non conserved
X similar
X ≥ 50% conserved