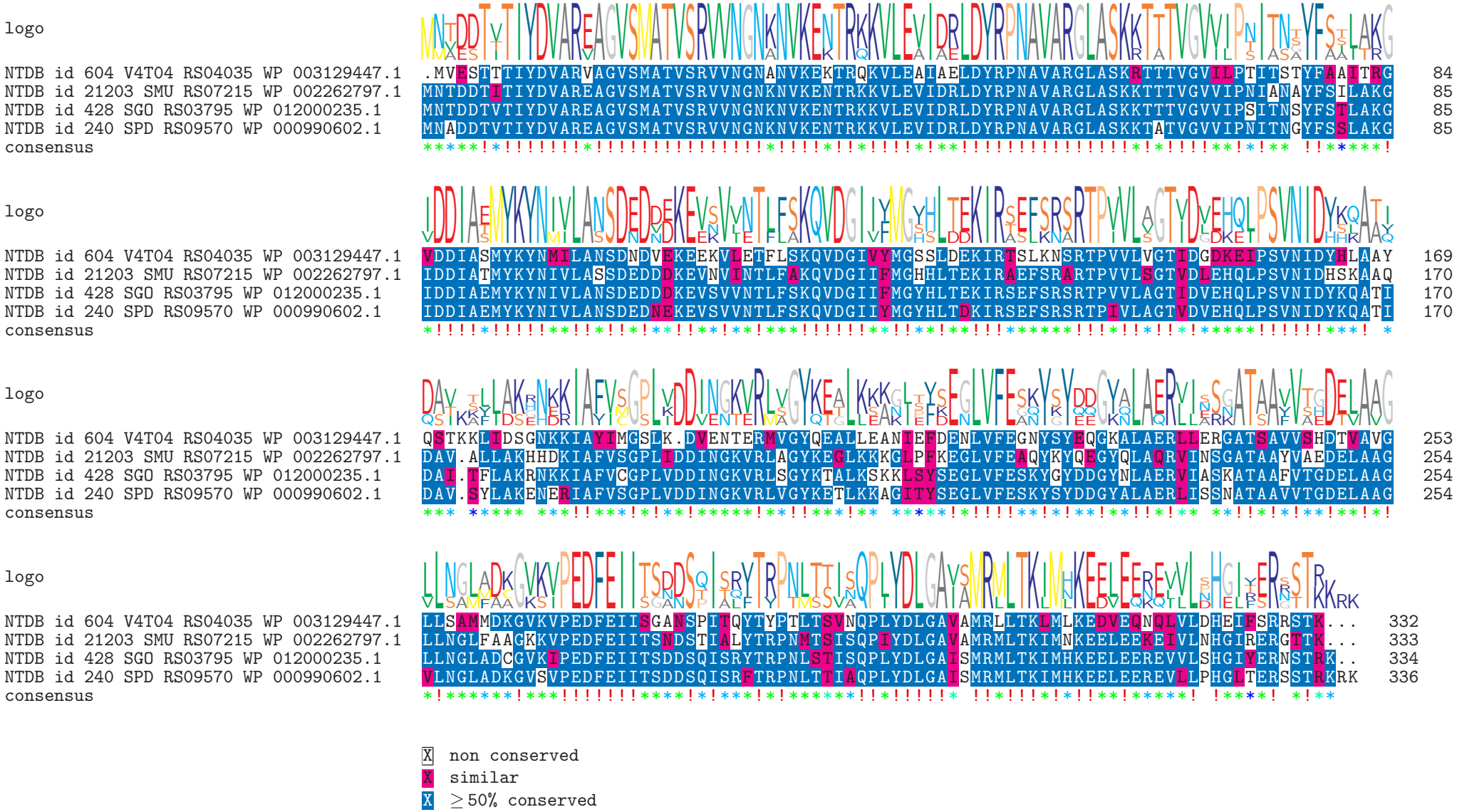

logo
NTDB id 604 V4T04 RS04035 WP 003129447.1 .MVESTTTIYDVARVAGVSMATVSRVVNGNANVKEKTRQKVLEAIAELDYRPNAVARGLASKRTTTVGVILPTITSTYFAAITRG 84
NTDB id 21203 SMU RS07215 WP 002262797.1 MNTDDTITIYDVAREAGVSMATVSRVVNGNKNVKENTRKKVLEVIDRLDYRPNAVARGLASKKTTTVGVVIPNIANAYFSILAKG 85
NTDB id 428 SGO RS03795 WP 012000235.1 MNTDDTVTIYDVAREAGVSMATVSRVVNGNKNVKENTRKKVLEVIDRLDYRPNAVARGLASKKTTTVGVVIPSITNSYFSTLAKG 85
NTDB id 240 SPD RS09570 WP 000990602.1 MNADDTVTIYDVAREAGVSMATVSRVVNGNKNVKENTRKKVLEVIDRLDYRPNAVARGLASKKTATVGVVIPNITNGYFSSLAKG 85
consensus
logo
NTDB id 604 V4T04 RS04035 WP 003129447.1 VDDIASMYKYNMILANSDNDVEKEEKVLETFLSKQVDGIVYMGSSLDEKIRTSLKNSRTPVVLVGTIDGDKEIPSVNIDYHLAAY 169
NTDB id 21203 SMU RS07215 WP 002262797.1 IDDIATMYKYNIVLASSDEDDDKEVNVINTLFAKQVDGIIFMGHHLTEKIRAEFSRARTPVVLSGTVDLEHQLPSVNIDHSKAAQ 170
NTDB id 428 SGO RS03795 WP 012000235.1 IDDIAEMYKYNIVLANSDEDDDKEVSVVNTLFSKQVDGIIFMGYHLTEKIRSEFSRSRTPVVLAGTIDVEHQLPSVNIDYKQATI 170
NTDB id 240 SPD RS09570 WP 000990602.1 IDDIAEMYKYNIVLANSDEDNEKEVSVVNTLFSKQVDGIIYMGYHLTDKIRSEFSRSRTPIVLAGTVDVEHQLPSVNIDYKQATI 170
consensus
logo
NTDB id 604 V4T04 RS04035 WP 003129447.1 QSTKKLIDSGNKKIAYIMGSLK.DVENTERMVGYQEALLEANIEFDENLVFEGNYSYEQGKALAERLLERGATSAVVSHDTVAVG 253
NTDB id 21203 SMU RS07215 WP 002262797.1 DAV.ALLAKHHDKIAFVSGPLIDDINGKVRLAGYKEGLKKKGLPFKEGLVFEAQYKYQEGYQLAQRVINSGATAAYVAEDELAAG 254
NTDB id 428 SGO RS03795 WP 012000235.1 DAI.TFLAKRNKKIAFVCGPLVDDINGKVRLSGYKTALKSKKLSYSEGLVFESKYGYDDGYNLAERVIASKATAAFVTGDELAAG 254
NTDB id 240 SPD RS09570 WP 000990602.1 DAV.SYLAKENERIAFVSGPLVDDINGKVRLVGYKETLKKAGITYSEGLVFESKYSYDDGYALAERLISSNATAAVVTGDELAAG 254
consensus
logo
NTDB id 604 V4T04 RS04035 WP 003129447.1 LLSAMMDKGVKVPEDFEIISGANSPITQYTYPTLTSVNQPLYDLGAVAMRLLTKLMLKEDVEQNQLVLDHEIFSRRSTK... 332
NTDB id 21203 SMU RS07215 WP 002262797.1 LLNGLFAAGKKVPEDFEIITSNDSTIALYTRPNMTSISQPIYDLGAVAMRMLTKIMNKEELEEKEIVLNHGIRERGTTK... 333
NTDB id 428 SGO RS03795 WP 012000235.1 LLNGLADCGVKIPEDFEIITSDDSQISRYTRPNLSTISQPLYDLGAISMRMLTKIMHKEELEEREVVLSHGIYERNSTRK... 334
NTDB id 240 SPD RS09570 WP 000990602.1 VLNGLADKGVSVPEDFEIITSDDSQISRFTRPNLTTIAQPLYDLGAISMRMLTKIMHKEELEEREVLLPHGLTERSSTRKRK 336
consensus
X non conserved
X similar
X ≥ 50% conserved