
logo
NTDB id 604 V4T04 RS04035 WP 003129447.1 .MVESTTTIYDVARVAGVSMATVSRVVNGNANVKEKTRQKVLEAIAELDYRPNAVARGLASKRTTTVGVILPTITSTYFA 79
NTDB id 86686 CUGBS98 RS03550 WP 001090621.1 MNTDDTITIYDVAREAGVSMATVSRVVNGNKNVKENTRKKVLEVIDRLDYRPNAVARGLASKKTTTVGVVIPNIANSYFS 80
NTDB id 428 SGO RS03795 WP 012000235.1 MNTDDTVTIYDVAREAGVSMATVSRVVNGNKNVKENTRKKVLEVIDRLDYRPNAVARGLASKKTTTVGVVIPSITNSYFS 80
NTDB id 240 SPD RS09570 WP 000990602.1 MNADDTVTIYDVAREAGVSMATVSRVVNGNKNVKENTRKKVLEVIDRLDYRPNAVARGLASKKTATVGVVIPNITNGYFS 80
consensus
logo
NTDB id 604 V4T04 RS04035 WP 003129447.1 AITRGVDDIASMYKYNMILANSDNDVEKEEKVLETFLSKQVDGIVYMGSSLDEKIRTSLKNSRTPVVLVGTIDGDKEIPS 159
NTDB id 86686 CUGBS98 RS03550 WP 001090621.1 ILARGIDDIAAMYKYNIVLASSDEDDDKEVNVVNTLFAKQVDGIIFMGHHLTEKIRAEFSRSRTPIVLAGTVDLEHQLPS 160
NTDB id 428 SGO RS03795 WP 012000235.1 TLAKGIDDIAEMYKYNIVLANSDEDDDKEVSVVNTLFSKQVDGIIFMGYHLTEKIRSEFSRSRTPVVLAGTIDVEHQLPS 160
NTDB id 240 SPD RS09570 WP 000990602.1 SLAKGIDDIAEMYKYNIVLANSDEDNEKEVSVVNTLFSKQVDGIIYMGYHLTDKIRSEFSRSRTPIVLAGTVDVEHQLPS 160
consensus
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NTDB id 604 V4T04 RS04035 WP 003129447.1 VNIDYHLAAYQSTKKLIDSGNKKIAYIMGSLK.DVENTERMVGYQEALLEANIEFDENLVFEGNYSYEQGKALAERLLER 238
NTDB id 86686 CUGBS98 RS03550 WP 001090621.1 VNIDYKAAAVDVIDILAGNHKDIAFVSGPLIDDINGKVRLAGYKEGLKKNGLNFKEGLVFEANYRYAEGFALAQRVINA 239
NTDB id 428 SGO RS03795 WP 012000235.1 VNIDYKQATIDAITFLAKRNKKIAFVCGPLVDDINGKVRLSGYKTALKSKKLSYSEGLVFESKYGYDDGYNLAERVIAS 239
NTDB id 240 SPD RS09570 WP 000990602.1 VNIDYKQATIDAVSYLAKENERIAFVSGPLVDDINGKVRLVGYKETLKKAGITYSEGLVFESKYSYDDGYALAERLISS 239
consensus
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NTDB id 604 V4T04 RS04035 WP 003129447.1 GATSAVVSHDTVAVGLLSAMMDKGVKVPEDFEIISGANSPITQYTYPTLTSVNQPLYDLGAVAMRLLTKLMLKEDVEQNQ 318
NTDB id 86686 CUGBS98 RS03550 WP 001090621.1 GATAAYVAEDELAAGLLNGLFEAGKRVPEDFEIITSNDSPIAQYTRPNLTSISQPVYDLGAVSMRMLTKIMHKEELEEKE 319
NTDB id 428 SGO RS03795 WP 012000235.1 KATAAFVTGDELAAGLLNGLADCGVKIPEDFEIITSDDSQISRYTRPNLSTISQPLYDLGAISMRMLTKIMHKEELEERE 319
NTDB id 240 SPD RS09570 WP 000990602.1 NATAAVVTGDELAAGVLNGLADKGVSVPEDFEIITSDDSQISRFTRPNLTTIAQPLYDLGAISMRMLTKIMHKEELEERE 319
consensus
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NTDB id 604 V4T04 RS04035 WP 003129447.1 LVLDHEIFSRRSTK... 332
NTDB id 86686 CUGBS98 RS03550 WP 001090621.1 IVLNHGIVKRGTTNN.. 334
NTDB id 428 SGO RS03795 WP 012000235.1 VVLSHGIYERNSTRK.. 334
NTDB id 240 SPD RS09570 WP 000990602.1 VLLPHGLTERSSTRKRK 336
consensus
X non conserved
X similar
X ≥ 50% conserved