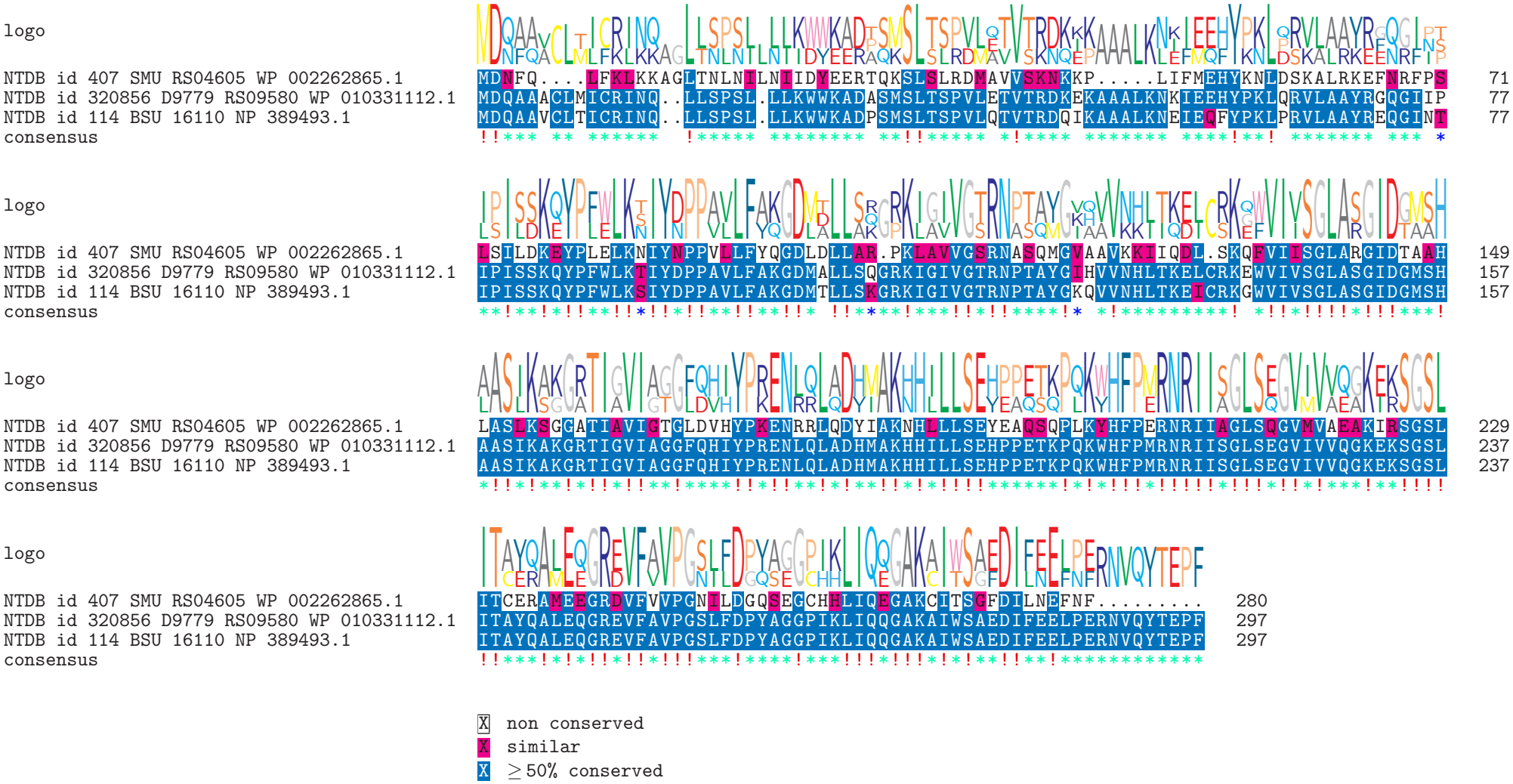
logo
NTDB id 407 SMU RS04605 WP 002262865.1 MDNFQ....LFKLKKAGLTNLNILNIIDYEERTQKSLSLRDMAVVSKNKKP.....LIFMEHYKNLDSKALRKEFNRFPS 71
NTDB id 320856 D9779 RS09580 WP 010331112.1 MDQAAACLMICRINQ..LLSPSL.LLKWWKADASMSLTSPVLETVTRDKEKAAALKNKIEEHYPKLQRVLAAYRGQGIIP 77
NTDB id 114 BSU 16110 NP 389493.1 MDQAAVCLTICRINQ..LLSPSL.LLKWWKADPSMSLTSPVLQTVTRDQIKAAALKNEIEQFYPKLPRVLAAYREQGINT 77
consensus
logo
NTDB id 407 SMU RS04605 WP 002262865.1 LSILDKEYPLELKNIYNPPVLLFYQGDLDLLAR.PKLAVVGSRNASQMGVAAVKKIIQDL.SKQFVIISGLARGIDTAAH 149
NTDB id 320856 D9779 RS09580 WP 010331112.1 IPISSKQYPFWLKTIYDPPAVLFAKGDMALLSQGRKIGIVGTRNPTAYGIHVVNHLTKELCRKEWVIVSGLASGIDGMSH 157
NTDB id 114 BSU 16110 NP 389493.1 IPISSKQYPFWLKSIYDPPAVLFAKGDMTLLSKGRKIGIVGTRNPTAYGKQVVNHLTKEICRKGWVIVSGLASGIDGMSH 157
consensus
logo
NTDB id 407 SMU RS04605 WP 002262865.1 LASLKSGGATIAVIGTGLDVHYPKENRRLQDYIAKNHLLLSEYEAQSQPLKYHFPERNRIIAGLSQGVMVAEAKIRSGSL 229
NTDB id 320856 D9779 RS09580 WP 010331112.1 AASIKAKGRTIGVIAGGFQHIYPRENLQLADHMAKHHILLSEHPPETKPQKWHFPMRNRIISGLSEGVIVVQGKEKSGSL 237
NTDB id 114 BSU 16110 NP 389493.1 AASIKAKGRTIGVIAGGFQHIYPRENLQLADHMAKHHILLSEHPPETKPQKWHFPMRNRIISGLSEGVIVVQGKEKSGSL 237
consensus
logo
NTDB id 407 SMU RS04605 WP 002262865.1 ITCERAMEEGRDVFVVPGNILDGQSEGCHHLIQEGAKCITSGFDILNEFNF..... 280
NTDB id 320856 D9779 RS09580 WP 010331112.1 ITAYQALEQGREVFAVPGSLFDPYAGGPIKLIQQGAKAIWSAEDIFEELPERNVQYTEPF 297
NTDB id 114 BSU 16110 NP 389493.1 ITAYQALEQGREVFAVPGSLFDPYAGGPIKLIQQGAKAIWSAEDIFEELPERNVQYTEPF 297
consensus
X non conserved
X similar
X ≥ 50% conserved