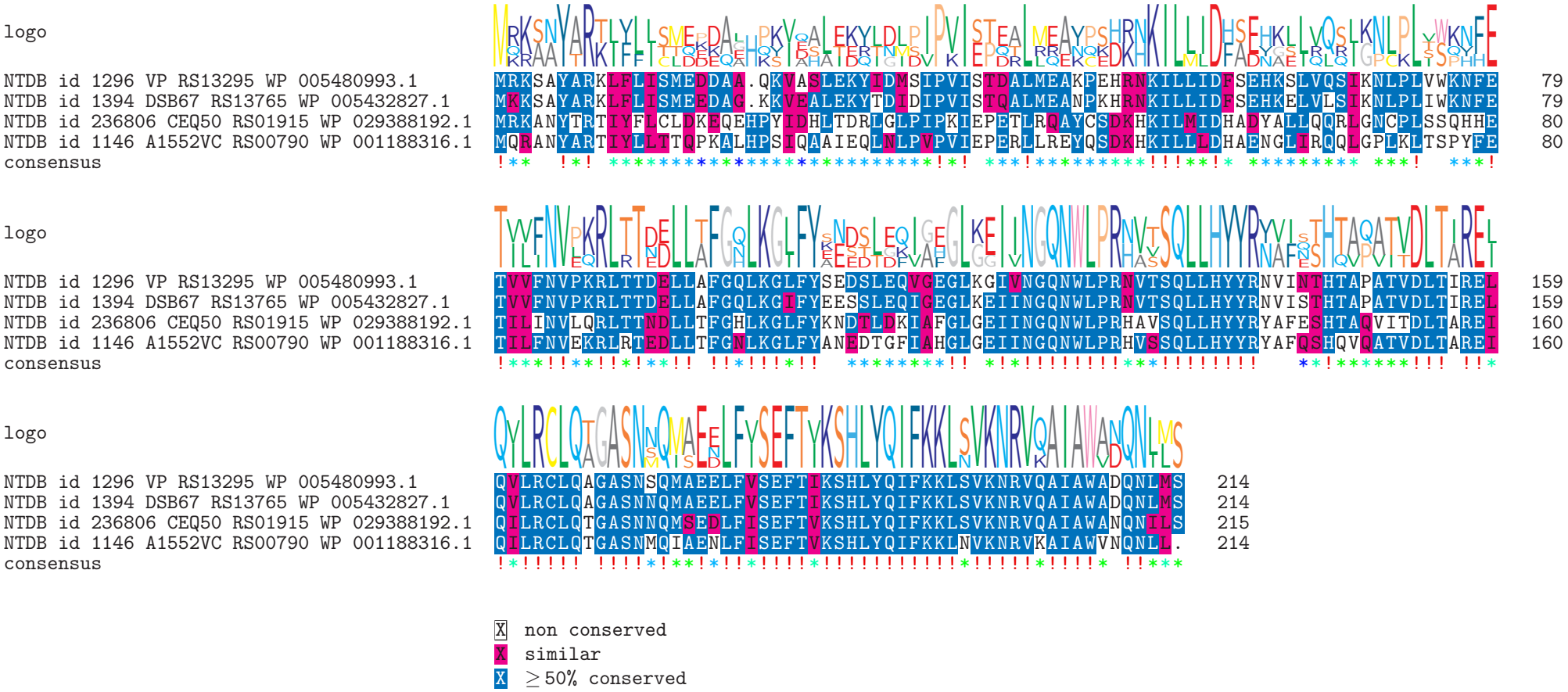
logo
NTDB id 1296 VP RS13295 WP 005480993.1 MRKSAYARKLFLISMEDDAA.QKVASLEKYIDMSIPVISTDALMEAKPEHRNKILLIDFSEHKSLVQSIKNLPLVWKNFE 79
NTDB id 1394 DSB67 RS13765 WP 005432827.1 MKKSAYARKLFLISMEEDAG.KKVEALEKYTDIDIPVISTQALMEANPKHRNKILLIDFSEHKELVLSIKNLPLIWKNFE 79
NTDB id 236806 CEQ50 RS01915 WP 029388192.1 MRKANYTRTIYFLCLDKEQEHPYIDHLTDRLGLPIPKIEPETLRQAYCSDKHKILMIDHADYALLQQRLGNCPLSSQHHE 80
NTDB id 1146 A1552VC RS00790 WP 001188316.1 MQRANYARTIYLLTTQPKALHPSIQAAIEQLNLPVPVIEPERLLREYQSDKHKILLLDHAENGLIRQQLGPLKLTSPYFE 80
consensus
logo
NTDB id 1296 VP RS13295 WP 005480993.1 TVVFNVPKRLTTDELLAFGQLKGLFYSEDSLEQVGEGLKGIVNGQNWLPRNVTSQLLHYYRNVINTHTAPATVDLTIREL 159
NTDB id 1394 DSB67 RS13765 WP 005432827.1 TVVFNVPKRLTTDELLAFGQLKGIFYEESSLEQIGEGLKEIINGQNWLPRNVTSQLLHYYRNVISTHTAPATVDLTIREL 159
NTDB id 236806 CEQ50 RS01915 WP 029388192.1 TILINVLQRLTTNDLLTFGHLKGLFYKNDTLDKIAFGLGEIINGQNWLPRHAVSQLLHYYRYAFESHTAQVITDLTAREI 160
NTDB id 1146 A1552VC RS00790 WP 001188316.1 TILFNVEKRLRTEDLLTFGNLKGLFYANEDTGFIAHGLGEIINGQNWLPRHVSSQLLHYYRYAFQSHQVQATVDLTAREI 160
consensus
logo
NTDB id 1296 VP RS13295 WP 005480993.1 QVLRCLQAGASNSQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS 214
NTDB id 1394 DSB67 RS13765 WP 005432827.1 QVLRCLQAGASNNQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS 214
NTDB id 236806 CEQ50 RS01915 WP 029388192.1 QILRCLQTGASNNQMSEDLFISEFTVKSHLYQIFKKLSVKNRVQAIAWANQNILS 215
NTDB id 1146 A1552VC RS00790 WP 001188316.1 QILRCLQTGASNMQIAENLFISEFTVKSHLYQIFKKLNVKNRVKAIAWVNQNLL. 214
consensus
X non conserved
X similar
X ≥ 50% conserved