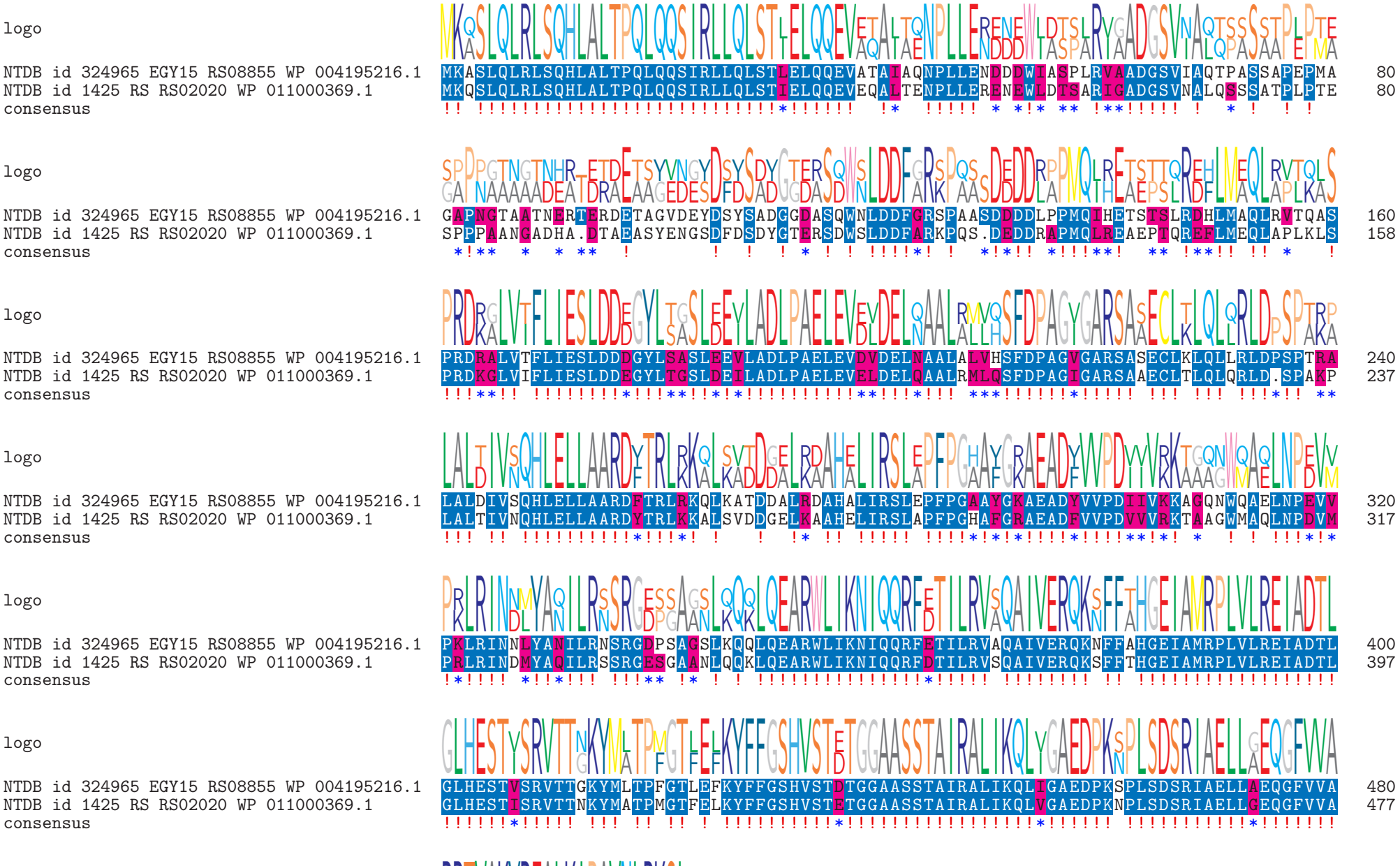

logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 MKASLQLRLSQHLALTPQLQQSIRLLQLSTLELQQEVATAIAQNPLLENDDDWIASPLRVAADGSVIAQTPASSAPEPMA 80
NTDB id 1425 RS RS02020 WP 011000369.1 MKQSLQLRLSQHLALTPQLQQSIRLLQLSTIELQQEVEQALTENPLLERENEWLDTSARIGADGSVNALQSSSATPLPTE 80
consensus
logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 GAPNGTAATNERTERDETAGVDEYDSYSADGGDASQWNLDDFGRSPAASDDDDLPPMQIHETSTSLRDHLMAQLRVTQAS 160
NTDB id 1425 RS RS02020 WP 011000369.1 SPPPAANGADHA.DTAEASYENGSDFDSDYGTERSDWSLDDFARKPQS.DEDDRAPMQLREAEPTQREFLMEQLAPLKLS 158
consensus
logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 PRDRALVTFLIESLDDDGYLSASLEEVLADLPAELEVDVDELNAALALVHSFDPAGVGARSASECLKLQLLRLDPSPTRA 240
NTDB id 1425 RS RS02020 WP 011000369.1 PRDKGLVIFLIESLDDEGYLTGSLDEILADLPAELEVELDELQAALRMLQSFDPAGIGARSAAECLTLQLQRLD.SPAKP 237
consensus
logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 LALDIVSQHLELLAARDFTRLRKQLKATDDALRDAHALIRSLEPFPGAAYGKAEADYVVPDIIVKKAGQNWQAELNPEVV 320
NTDB id 1425 RS RS02020 WP 011000369.1 LALTIVNQHLELLAARDYTRLKKALSVDDGELKAAHELIRSLAPFPGHAFGRAEADFVVPDVVVRKTAAGWMAQLNPDVM 317
consensus
logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 PKLRINNLYANILRNSRGDPSAGSLKQQLQEARWLIKNIQQRFETILRVAQAIVERQKNFFAHGEIAMRPLVLREIADTL 400
NTDB id 1425 RS RS02020 WP 011000369.1 PRLRINDMYAQILRSSRGESGAANLQQKLQEARWLIKNIQQRFDTILRVSQAIVERQKSFFTHGEIAMRPLVLREIADTL 397
consensus
logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 GLHESTVSRVTTGKYMLTPFGTLEFKYFFGSHVSTDTGGAASSTAIRALIKQLIGAEDPKSPLSDSRIAELLAEQGFVVA 480
NTDB id 1425 RS RS02020 WP 011000369.1 GLHESTISRVTTNKYMATPMGTFELKYFFGSHVSTETGGAASSTAIRALIKQLVGAEDPKNPLSDSRIAELLGEQGFVVA 477
consensus
logo
NTDB id 324965 EGY15 RS08855 WP 004195216.1 RRTVAKYREALKIPAVNLRKSL 502
NTDB id 1425 RS RS02020 WP 011000369.1 RRTVAKYREALKIPAVNLRKSL 499
consensus

X non conserved

X similar

X $\geq$ 50% conserved