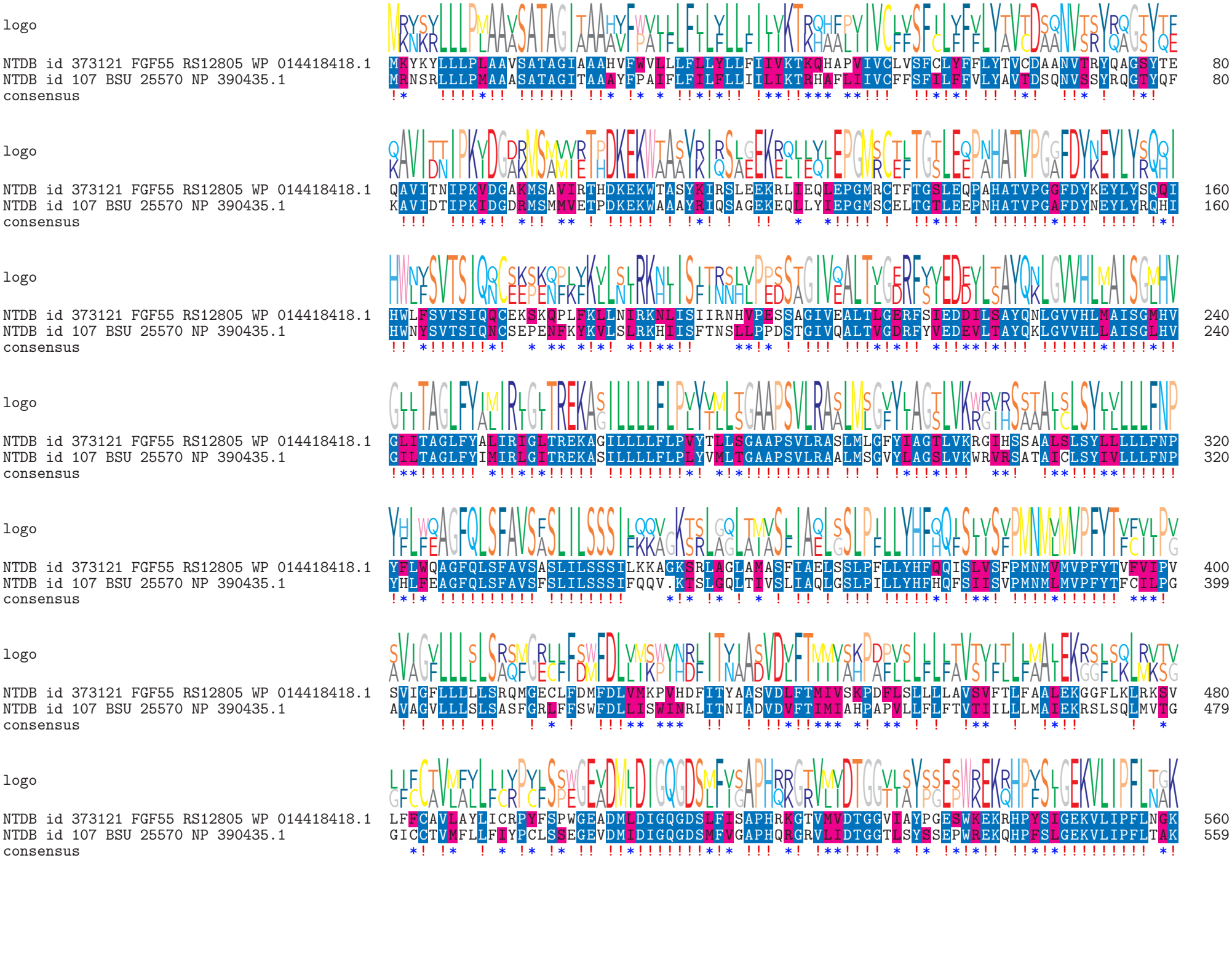
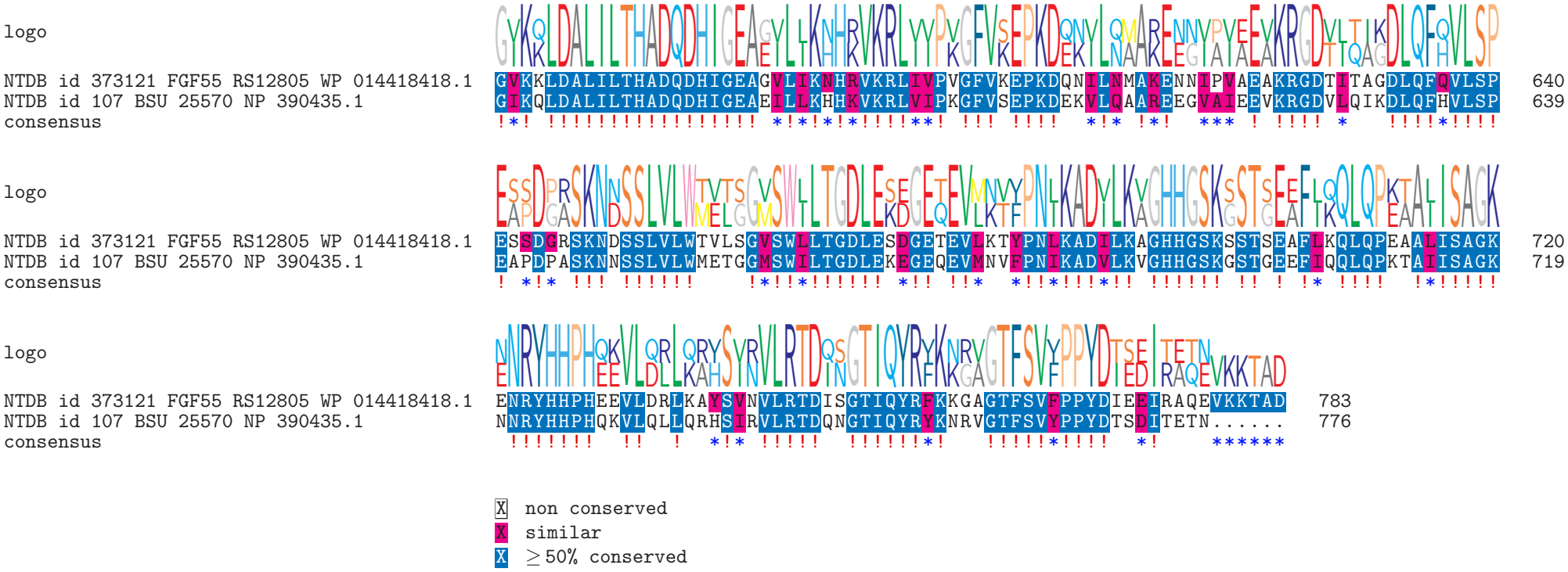

logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 MKYKYLLLPLAAVSATAGIAAAHVFWVLLLFLLYLLFIIVKTKQHAPVIVCLVSFCLYFFLYTVCDAANVTRYQAGSYTE 80
NTDB id 107 BSU 25570 NP 390435.1 MRNSRLLLPMAAASATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF 80
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 QAVITNIPKVDGAKMSAVIRTHDKEKWTASYKIRSLEEKRLIEQLEPGMRCTFTGSLEQPAHATVPGGFDYKEYLYSQQI 160
NTDB id 107 BSU 25570 NP 390435.1 KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHI 160
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 HWLFSVTSIQQCEKSKQPLFKLLNIRKNLISIIRNHVPESSAGIVEALTLGERFSIEDDILSAYQNLGVVHLMAISGMHV 240
NTDB id 107 BSU 25570 NP 390435.1 HWNYSVTSIQNCSEPENFKYKVLSLRKHIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGVVHLLAISGLHV 240
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 GLITAGLFYALIRIGLTREKAGILLLLFLPVYTLLSGAAPSVLRASLMLGFYIAGTLVKRGIHSSAALSLSYLLLLLFNP 320
NTDB id 107 BSU 25570 NP 390435.1 GILTAGLFYIMIRLGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVRSATAICLSYIVLLLFNP 320
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 YFLWQAGFQLSFAVSASLILSSSILKKAGKSRLAGLAMASFIAELSSLPFLLYHFQQISLVSFPMNMVMVPFYTVFVIPV 400
NTDB id 107 BSU 25570 NP 390435.1 YHLFEAGFQLSFAVSFSLILSSSIFQQV.KTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPG 399
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 SVIGFLLLLLSRQMGECLFDMFDLVMKPVHDFITYAASVDLFTMIVSKPDFLSLLLLAVSVFTLFAALEKGGFLKLRKSV 480
NTDB id 107 BSU 25570 NP 390435.1 AVAGVLLLSLSASFGRLFFSWFDLLISWINRLITNIADVDVFTIMIAHPAPVLLFLFTVTIILLLMAIEKRSLSQLMVTG 479
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 LFFCAVLAYLICRPYFSPWGEADMLDIGQGDSLFISAPHRKGTVMVDTGGVIAYPGESWKEKRHPYSIGEKVLIPFLNGK 560
NTDB id 107 BSU 25570 NP 390435.1 GICCTVMFLLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLIDTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAK 559
consensus


logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 GVKKLDALILTHADQDHIGEAGVLIKNHRVKRLIVPVGFVKEPKDQNILNMAKENNIPVAEAKRGDTITAGDLQFQVLSP 640
NTDB id 107 BSU 25570 NP 390435.1 GIKQLDALILTHADQDHIGEAEILLKHHKVKRLVIPKGFVSEPKDEKVLQAAREEGVAIEEVKRGDVLQIKDLQFHVLSP 639
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 ESSDGRSKNDSSLVLWTVLSGVSWLLTGDLESDGETEVLKTYPNLKADILKAGHHGSKSSTSEAFLKQLQPEAALISAGK 720
NTDB id 107 BSU 25570 NP 390435.1 EAPDPASKNNSSLVLWMETGGMSWILTGDLEKEGEQEVMNVFPNIKADVLKVGHHGSKGSTGEEFIQQLQPKTAIISAGK 719
consensus
logo
NTDB id 373121 FGF55 RS12805 WP 014418418.1 ENRYHHPHEEVLDRLKAYSVNVLRTDISGTIQYRFKKGAGTFSVFPPYDIEEIRAQEVKKTAD 783
NTDB id 107 BSU 25570 NP 390435.1 NNRYHHPHQKVLQLLQRHSIRVLRTDQNGTIQYRYKNRVGTFSVYPPYDTSDITETN..... 776
consensus
X non conserved
X similar
X ≥ 50% conserved