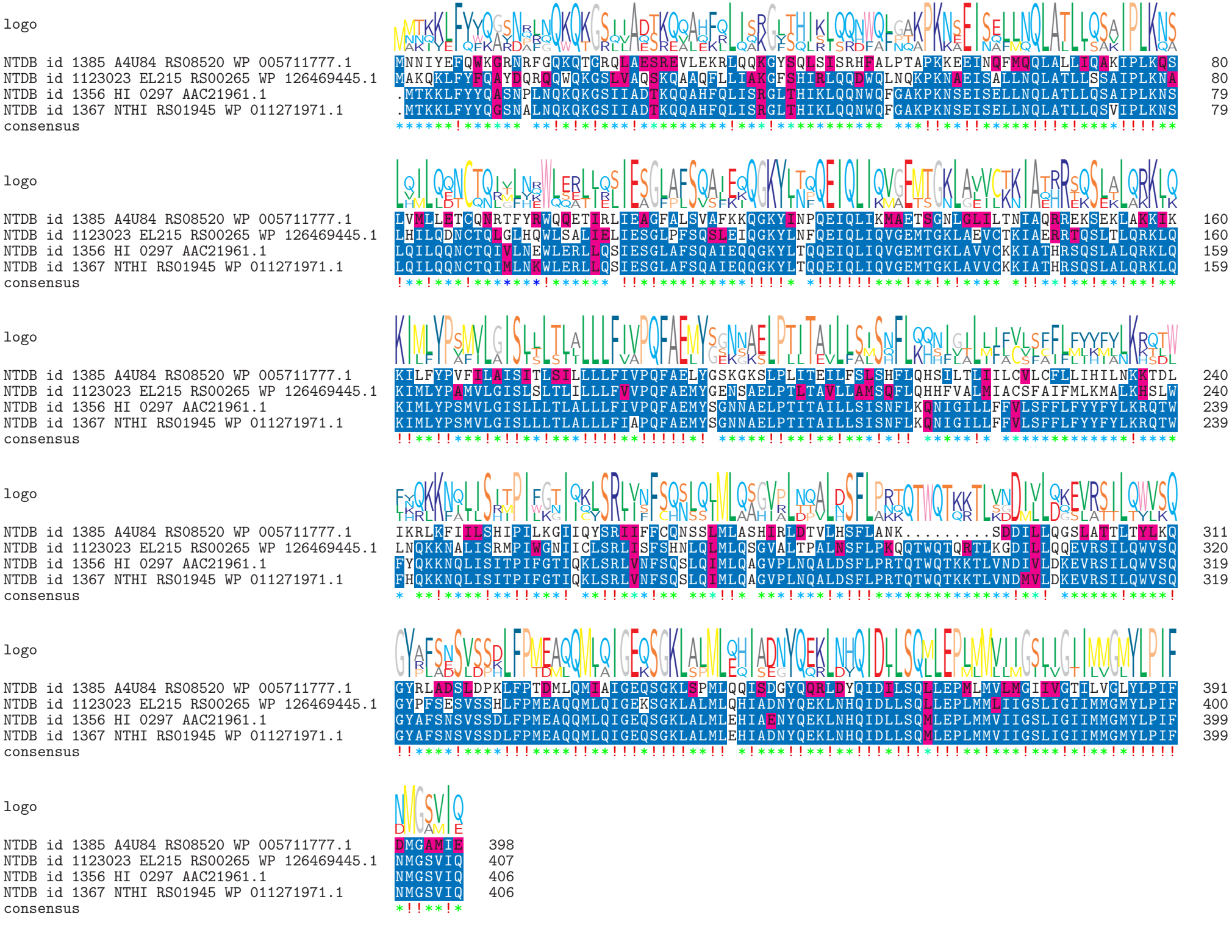
logo
NTDB id 1385 A4U84 RS08520 WP 005711777.1
NTDB id 1123023 EL215 RS00265 WP 126469445.1
NTDB id 1356 HI 0297 AAC21961.1
NTDB id 1367 NTHI RS01945 WP 011271971.1
consensus
MNNIYEFQWKGRNRFGQKQTGRQLAESREVLEKRLQQKGYSQLSISRHFALPTAPKKEEINQFMQQLALLIQAKIPLKQS 80
MAKQKLFYFQAYDQRQQWQKGSLVAQSKQAAQFLLIAKGFSHIRLQQDWQLNQKPKNAEISALLNQLATLLSSAIPLKNA 80
.MTKKLFYYQASNPLNQKQKGSIIADTKQQAHFQLISRGLTHIKLQQNWQFGAKPKNSEISELLNQLATLLQSAIPLKNS 79
.MTKKLFYYQGSNALNQKQKGSIIADTKQQAHFQLISRGLTHIKLQQNWQFGAKPKNSEISELLNQLATLLQSVIPLKNS 79
LQILQQNCTQIYLNRWLERLLQSIESGLAFSQAIEQQGKYLTQQEIQLLQVGEMTGKLAVVCTKIATRRSQSLALQRKLQ 160
LHILQDNCTQLGLHQWLSALIELIESGLPFSQSLEIQGKYLNFQEIQLIQVGEMTGKLAEVCTKIAERRTQSLTLQRKLQ 160
LQILQQNCTQIVLNEWLERLLQSIESGLAFSQAIEQQGKYLTQQEIQLIQVGEMTGKLAVVCKKIATHRSQSLALQRKLQ 159
LQILQQNCTQIMLNKWLERLLQSIESGLAFSQAIEQQGKYLTQQEIQLIQVGEMTGKLAVVCKKIATHRSQSLALQRKLQ 159
KIMLYPSMVLGISLLLTLALLLFIVPQFAEMYSGNNAELPTITAILLSISNFLKQNIGILLFFVLSFFLFYYFYLKRQTW 240
KIMLYPAMVLGISLSLTLILLLFVVPQFAEMYGENSAELPTLTAVLLAMSQFLQHHFVALMIACSFAIFMLKMALKHSLW 240
KIMLYPSMVLGISLLLTLALLLFIVPQFAEMYSGNNAELPTITAILLSISNFLKQNIGILLFFVLSFFLFYYFYLKRQTW 239
KIMLYPSMVLGISLLLTLALLLFIAPQFAEMYSGNNAELPTITAILLSISNFLKQNIGILLFFVLSFFLFYYFYLKRQTW 239
IKRLKFIILSHIPILKGIIQYSRIIFFCQNSSLMLASHIRLDTVLHSFLANK.....SDDILLQGSLATTLTYLKQ 311
LNQKKNALISRMPIWGNIICLSRLISFSHNLQLMLQSGVALTPALNSFLPKQQTWQTQRTLKGDILLQQEVRSILQWVSQ 320
FYQKKNQLISITPIFGTIQKLSRLVNFSQSLQIMLQAGVPLNQALDSFLPRTQTWQTKKTLVNDIVLDKEVRSILQWVSQ 319
FHQKKNQLISITPIFGTIQKLSRLVNFSQSLQIMLQAGVPLNQALDSFLPRTQTWQTKKTLVNDMVLDKEVRSILQWVSQ 319
GYRLADSLDPKLFPTDMLQMIAIGEQSGKLSPMLQQISDGYQQRLDYQIDILSQLLEPMLMVLMGIIVGTILVGLYLPIF 391
GYPFSESVSSHLFPMEAQQMLQIGEKSGKLALMLQHIADNYQEKLNHQIDLLSQLLEPLMMLIIGSLIGIIMMGMYLPIF 400
GYAFSNSVSSDLFPMEAQQMLQIGEQSGKLALMLEHIAENYQEKLNHQIDLLSQMLEPLMMVIIGSLIGIIMMGMYLPIF 399
GYAFSNSVSSDLFPMEAQQMLQIGEQSGKLALMLEHIADNYQEKLNHQIDLLSQMLEPLMMVIIGSLIGIIMMGMYLPIF 399
DMGAMIE 398
NMGSVIQ 407
NMGSVIQ 406
NMGSVIQ 406

X non conserved
X similar
X ≥ 50% conserved