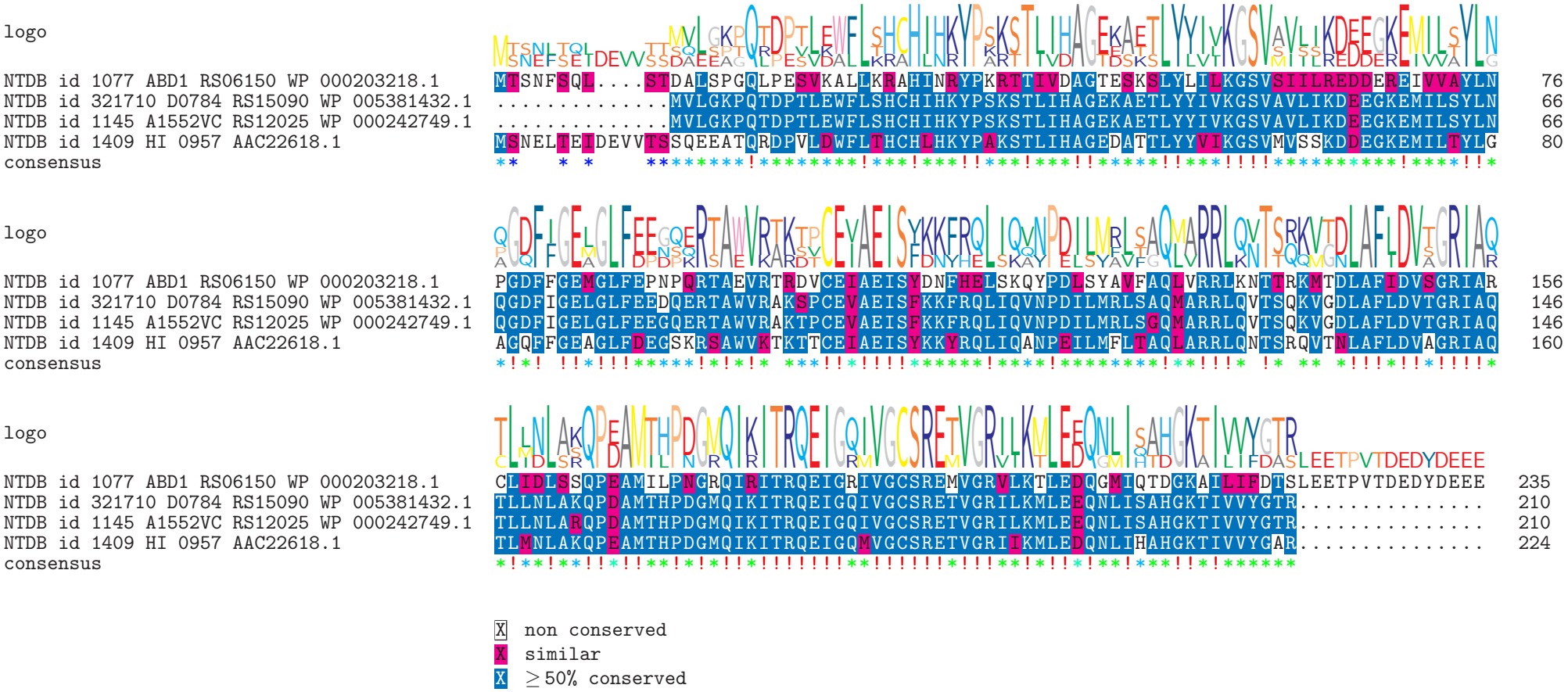
logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1 MTSNFSQL....STDALSPGQLPESVKALLKRAHINRYPKRTTIVDAGTESKSLYLILKGSVSIILREDDEREIVVAYLN 76
NTDB id 321710 D0784 RS15090 WP 005381432.1 .............MVLGKPQTDPTLEWFLSHCHIHKYPSKSTLIHAGEKAETLYYIVKGSVAVLIKDEEGKEMILSYLN 66
NTDB id 1145 A1552VC RS12025 WP 000242749.1 .............MVLGKPQTDPTLEWFLSHCHIHKYPSKSTLIHAGEKAETLYYIVKGSVAVLIKDEEGKEMILSYLN 66
NTDB id 1409 HI 0957 AAC22618.1 MSNELTEIDEVVTSSQEEATQRDPVLDWFLTHCHLHKYPAKSTLIHAGEDATTLYYVIKGSVMVSSKDDEGKEMILTYLG 80
consensus
logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1 PGDFFGEMGLFEPNPQRTAEVRTRDVCEIAEISYDNFHELSKQYPDLSYAVFAQLVRRLKNTTRKMTDLAFIDVSGRIAR 156
NTDB id 321710 D0784 RS15090 WP 005381432.1 QGDFIGELGLFEEDQERTAWVRAKSPCEVAEISFKKFRQLIQVNPDILMRLSAQMARRLQVTSQKVGDLAFLDVTGRIAQ 146
NTDB id 1145 A1552VC RS12025 WP 000242749.1 QGDFIGELGLFEEGQERTAWVRAKTPCEVAEISFKKFRQLIQVNPDILMRLSGQMARRLQVTSQKVGDLAFLDVTGRIAQ 146
NTDB id 1409 HI 0957 AAC22618.1 AGQFFGEAGLFDEGSKRSAWVKTKTTCEIAEISYKKYRQLIQANPEILMFLTAQLARRLQNTSRQVTNLAFLDVAGRIAQ 160
consensus
logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1 CLIDLSSQPEAMILPNGRQIRITRQEIGRIVGCSREMVGRVLKTLEDQGMIQTDGKAILIFDTSLEETPVTDEDYDEEE 235
NTDB id 321710 D0784 RS15090 WP 005381432.1 TLLNLAKQPDAMTHPDGMQIKITRQEIGQIVGCSRETVGRILKMLEEQNLISAHGKTIVVYGTR.............. 210
NTDB id 1145 A1552VC RS12025 WP 000242749.1 TLLNLARQPDAMTHPDGMQIKITRQEIGQIVGCSRETVGRILKMLEEQNLISAHGKTIVVYGTR.............. 210
NTDB id 1409 HI 0957 AAC22618.1 TLMNLAKQPEAMTHPDGMQIKITRQEIGQMVGCSRETVGRIIKMLEDQNLIHAHGKTIVVYGAR.............. 224
consensus
X non conserved
X similar
X ≥ 50% conserved