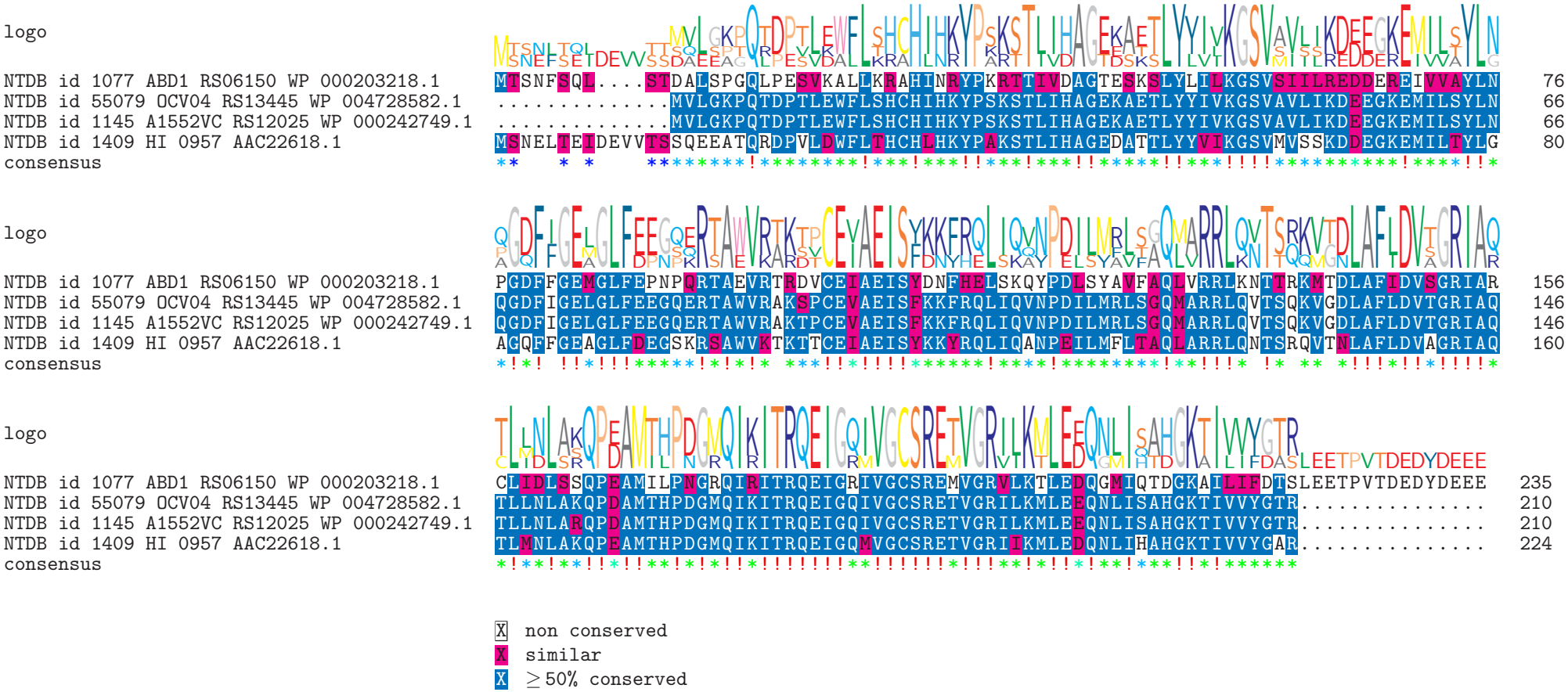
logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1
NTDB id 55079 OCV04 RS13445 WP 004728582.1
NTDB id 1145 A1552VC RS12025 WP 000242749.1
NTDB id 1409 HI 0957 AAC22618.1
consensus
MTSNFSQL....STDALSPGQLPESVKALLKRAHINRYPKRTTIVDAGTESKSLYLILKGSVSIILREDDEREIVVAYLN 76
.............MVLGKPQTDPTLEWFLSHCHIHKYPSKSTLIHAGEKAETLYYIVKGSVAVLIKDEEGKEMILSYLN 66
.............MVLGKPQTDPTLEWFLSHCHIHKYPSKSTLIHAGEKAETLYYIVKGSVAVLIKDEEGKEMILSYLN 66
MSNELTEIDEVVTSSQEEATQRDPVLDWFLTHCHLHKYPAKSTLIHAGEDATTLYYVIKGSVMVSSKDDEGKEMILTYLG 80
NTDB id 1077 ABD1 RS06150 WP 000203218.1
NTDB id 55079 OCV04 RS13445 WP 004728582.1
NTDB id 1145 A1552VC RS12025 WP 000242749.1
NTDB id 1409 HI 0957 AAC22618.1
consensus
PGDFFGEMGLFEPNPQRTAEVRTRDVCEIAEISYDNFHELSKQYPDLSYAVFAQLVRRLKNTTRKMTDLAFIDVSGRIAR 156
QGDFIGELGLFEEGQERTAWVRAKSPCEVAEISFKKFRQLIQVNPDILMRLSGQMARRLQVTSQKVGDLAFLDVTGRIAQ 146
QGDFIGELGLFEEGQERTAWVRAKTPCEVAEISFKKFRQLIQVNPDILMRLSGQMARRLQVTSQKVGDLAFLDVTGRIAQ 146
AGQFFGEAGLFDEGSKRSAWVKTKTTCEIAEISYKKYRQLIQANPEILMFLTAQLARRLQNTSRQVTNLAFLDVAGRIAQ 160
NTDB id 1077 ABD1 RS06150 WP 000203218.1
NTDB id 55079 OCV04 RS13445 WP 004728582.1
NTDB id 1145 A1552VC RS12025 WP 000242749.1
NTDB id 1409 HI 0957 AAC22618.1
consensus
CLIDLSSQPEAMILPNGRQIRITRQEIGRIVGCSREMVGRVLKTLEDQGMIQTDGKAILIFDTSLEETPVTDEDYDEEE 235
TLLNLAKQPDAMTHPDGMQIKITRQEIGQIVGCSRETVGRILKMLEEQNLISAHGKTIVVYGTR.............. 210
TLLNLARQPDAMTHPDGMQIKITRQEIGQIVGCSRETVGRILKMLEEQNLISAHGKTIVVYGTR.............. 210
TLMNLAKQPEAMTHPDGMQIKITRQEIGQMVGCSRETVGRIIKMLEDQNLIHAHGKTIVVYGAR.............. 224
NTDB id 1077 ABD1 RS06150 WP 000203218.1
NTDB id 55079 OCV04 RS13445 WP 004728582.1
NTDB id 1145 A1552VC RS12025 WP 000242749.1
NTDB id 1409 HI 0957 AAC22618.1
consensus
X non conserved
X similar
X ≥ 50% conserved