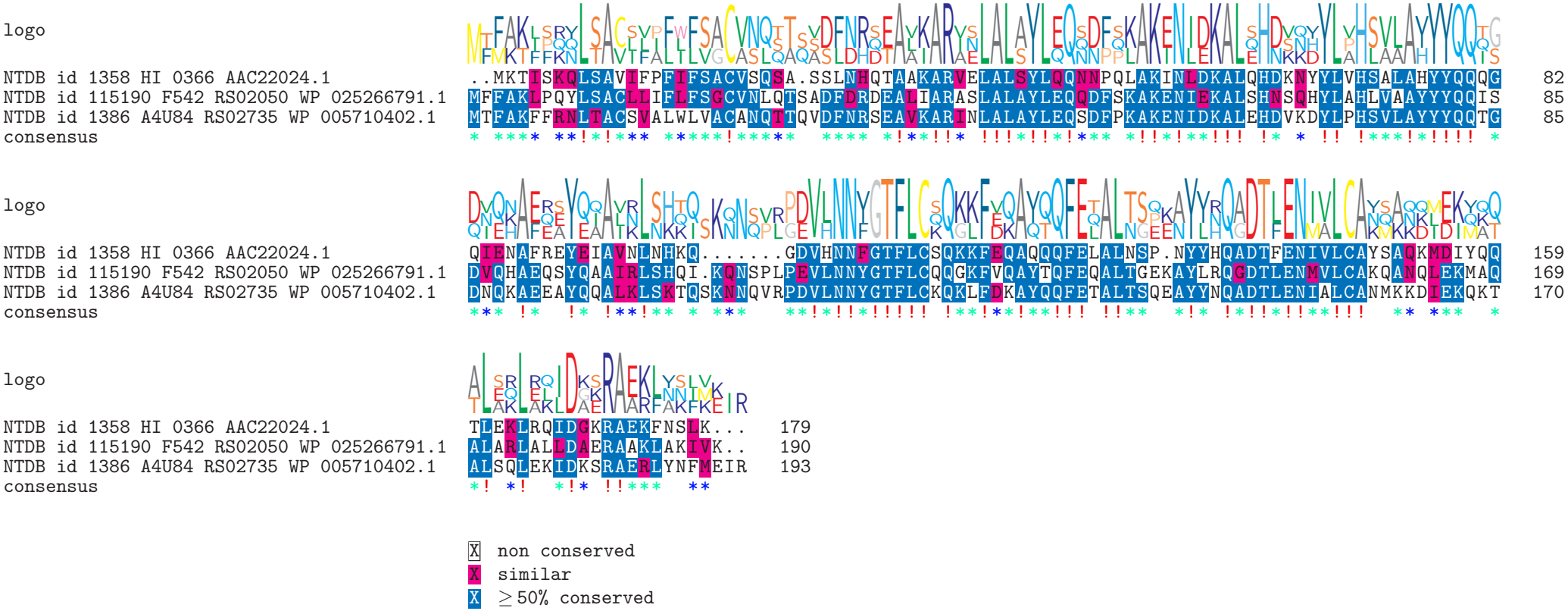
logo
NTDB id 1358 HI 0366 AAC22024.1 ..MKTISKQLSAVIFPFIFSACVSQSA.SSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG 82
NTDB id 115190 F542 RS02050 WP 025266791.1 MFFAKLPQYLSACLLIFLFSGCVNLQTSADFDRDEALIARASLALAYLEQQDFSKAKENIEKALSHNSQHYLAHLVAAYYYQQIS 85
NTDB id 1386 A4U84 RS02735 WP 005710402.1 MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG 85
consensus
logo
NTDB id 1358 HI 0366 AAC22024.1 QIENAFREYEIAVNLNHKQ.......GDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVLCAYSAQKMDIYQQ 159
NTDB id 115190 F542 RS02050 WP 025266791.1 DVQHAEQSYQAAIRLSHQI.KQNSPLPEVLNNYGTFLCQQGKFVQAYTQFEQALTGEKAYLRQGDTLENMVLCAKQANQLEKMAQ 169
NTDB id 1386 A4U84 RS02735 WP 005710402.1 DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT 170
consensus
logo
NTDB id 1358 HI 0366 AAC22024.1 TLEKLRQIDGKRAEKFNSLK... 179
NTDB id 115190 F542 RS02050 WP 025266791.1 ALARLALLDAERAAKLAKLVK.. 190
NTDB id 1386 A4U84 RS02735 WP 005710402.1 ALSQLEKIDKSRAERLYNFMEIR 193
consensus
X non conserved
X similar
X ≥ 50% conserved