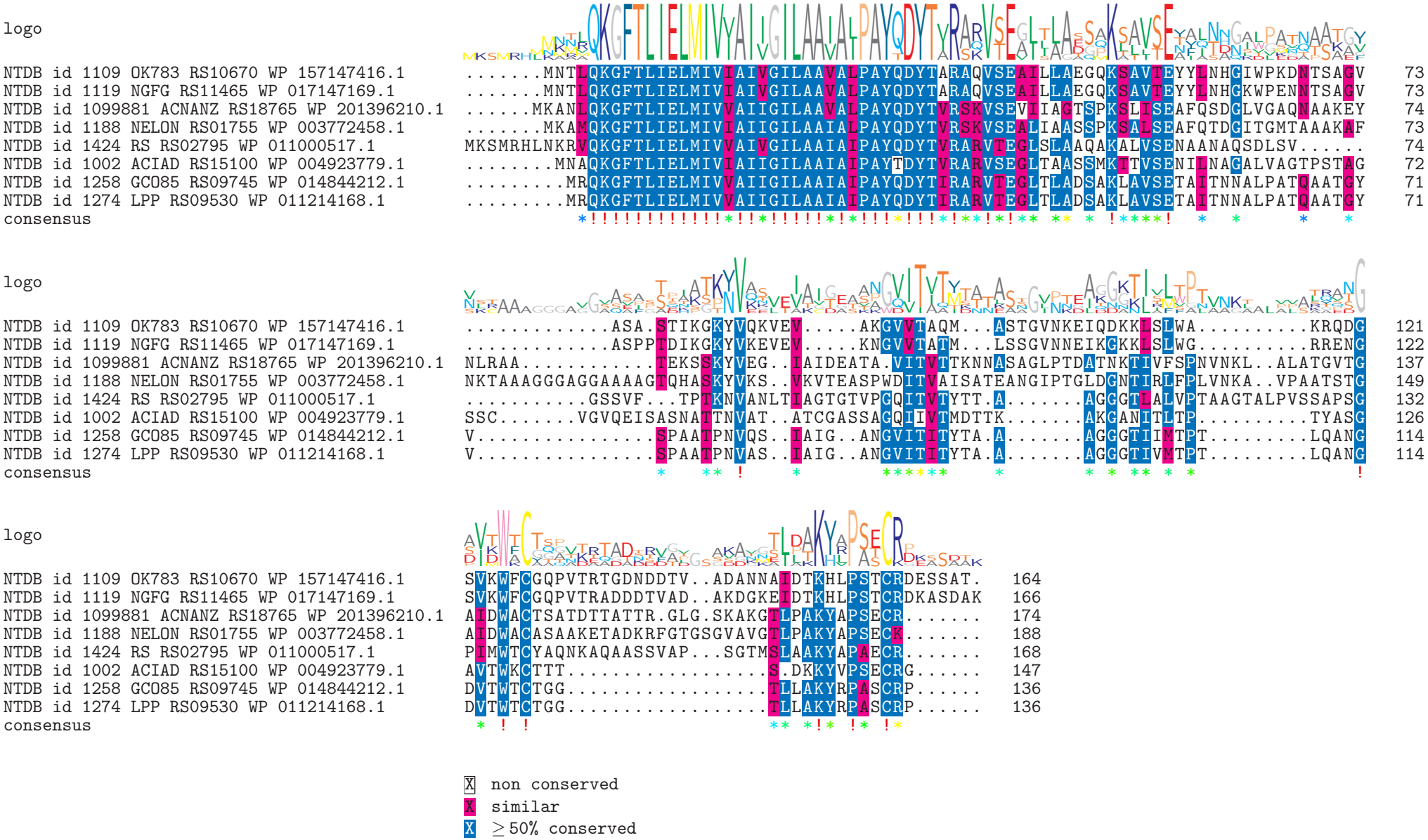
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NTDB id 1109 OK783 RS10670 WP 157147416.1 .....MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGIWPKDNTSAGV 73
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NTDB id 1099881 ACNANZ RS18765 WP 201396210.1 .....MKANLQKGFTLIELMIVVAIIGILAAVALPAYQDYTVRSKVSEVIIAGTSPKSLISEAFQSDGLVGAQNAAKEY 74
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NTDB id 1424 RS RS02795 WP 011000517.1 MKSMRHLNKRVQKGFTLIELMIVVAIVGILAAIAIPAYQDYTVRARVTECLSLAAQAKALVSENAANAQSDLSV..... 74
NTDB id 1002 ACIAD RS15100 WP 004923779.1 .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSECLTAASSMKTTVSENILNAGALVAGTPSTAG 72
NTDB id 1258 GCO85 RS09745 WP 014844212.1 .....MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGY 71
NTDB id 1274 LPP RS09530 WP 011214168.1 .....MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGY 71
consensus
NTDB id 1109 OK783 RS10670 WP 157147416.1 .....ASA.STIKGKYVQKVEV.....AKGVVTAQM...ASTGVNKEIQDKKLSLWA.....KRQDG 121
NTDB id 1119 NGFG RS11465 WP 017147169.1 .....ASPPTDIKGKYVKEVEV.....KNGVVTATM...LSSGVNNEIKGKKLSLWG.....RRENG 122
NTDB id 1099881 ACNANZ RS18765 WP 201396210.1 NLRAA.....TEKSSKYVEG..IAIDEATA.VITVTTKNNASAGLPTDATNKTIVFSPNVNKL..ALATGVTG 137
NTDB id 1188 NELON RS01755 WP 003772458.1 NKTAAAGGGAGGAAAAGTQHASKYVKS..VKVTEASPWDITVAISATEANGIPTGLDGNTIRLFPLVNKA..VPAATSTG 149
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NTDB id 1274 LPP RS09530 WP 011214168.1 V.....SPAATPNVAS..IAIG..ANGVITITYTA.A.....AGGGTIVMTPT.....LQANG 114
consensus
logo
NTDB id 1109 OK783 RS10670 WP 157147416.1 SVKWFCGQPVTRTGDNDDTV..ADANNAIDTKHLPSTCRDESSAT. 164
NTDB id 1119 NGFG RS11465 WP 017147169.1 SVKWFCGQPVTRADDDTVAD..AKDGKEIDTKHLPSTCRDKASDAK 166
NTDB id 1099881 ACNANZ RS18765 WP 201396210.1 AIDWACTSATDTTATTR.GLG.SKAKGTLPAKYAPSECR..... 174
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NTDB id 1258 GCO85 RS09745 WP 014844212.1 DVTWTCTGG.....TLLAKYRPASCRP..... 136
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consensus
X non conserved
X similar
X ≥50% conserved