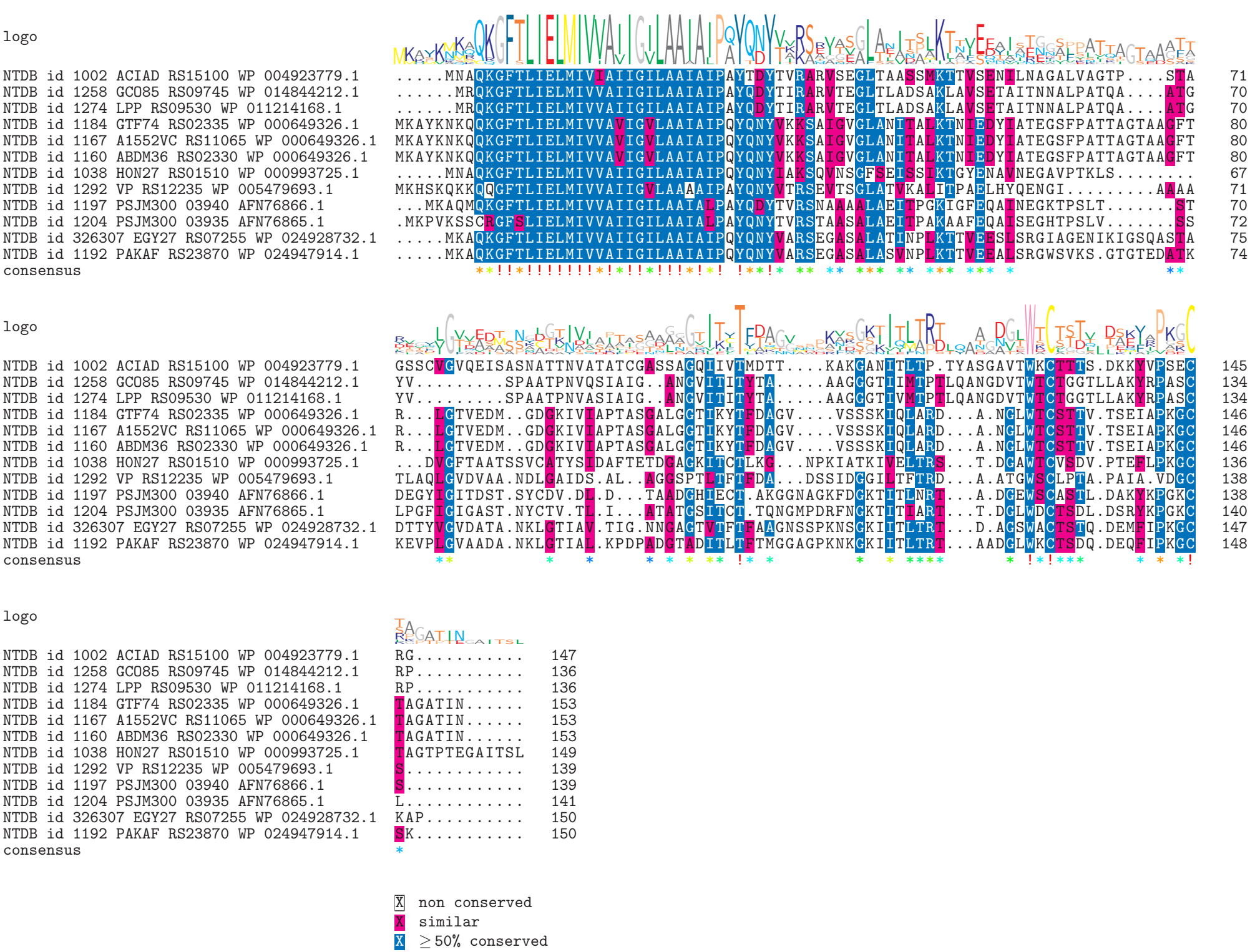
logo
NTDB id 1002 ACIAD RS15100 WP 004923779.1 .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTP....STA 71
NTDB id 1258 GCO85 RS09745 WP 014844212.1 ......MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQA....ATG 70
NTDB id 1274 LPP RS09530 WP 011214168.1 ......MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQA....ATG 70
NTDB id 1184 GTF74 RS02335 WP 000649326.1 MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT 80
NTDB id 1167 A1552VC RS11065 WP 000649326.1 MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT 80
NTDB id 1160 ABDM36 RS02330 WP 000649326.1 MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT 80
NTDB id 1038 HON27 RS01510 WP 000993725.1 .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAVPTKLS........ 67
NTDB id 1292 VP RS12235 WP 005479693.1 MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.........AAAA 71
NTDB id 1197 PSJM300 03940 AFN76866.1 ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLT.......ST 70
NTDB id 1204 PSJM300 03935 AFN76865.1 .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLV.......SS 72
NTDB id 326307 EGY27 RS07255 WP 024928732.1 .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALATINPLKTTVEESLSRGIAGENIKIGSQASTA 75
NTDB id 1192 PAKAF RS23870 WP 024947914.1 .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKS.GTGTEDATK 74
consensus
logo
NTDB id 1002 ACIAD RS15100 WP 004923779.1 GSSCVGVQEISASNATTNVATATCGASSAGQQIIVTMDTT....KAKGANITLTP.TYASGAVTWKCTTTS.DKKYVPSEC 145
NTDB id 1258 GCO85 RS09745 WP 014844212.1 YV.........SPAATPNVQSIAIG..ANGVITITYTA.....AAGGGTIIMTPTLQANGDVTWTCTGGTLLAKYRPASC 134
NTDB id 1274 LPP RS09530 WP 011214168.1 YV.........SPAATPNVASIAIG..ANGVITITYTA.....AAGGGTIVMTPTLQANGDVTWTCTGGTLLAKYRPASC 134
NTDB id 1184 GTF74 RS02335 WP 000649326.1 R...LGTVEDM..GDGKIVIAPTASGALGGTIKYTFDAGV....VSSSKIQLARD...A.NGLWTCSTTV.TSEIAPKGC 146
NTDB id 1167 A1552VC RS11065 WP 000649326.1 R...LGTVEDM..GDGKIVIAPTASGALGGTIKYTFDAGV....VSSSKIQLARD...A.NGLWTCSTTV.TSEIAPKGC 146
NTDB id 1160 ABDM36 RS02330 WP 000649326.1 R...LGTVEDM..GDGKIVIAPTASGALGGTIKYTFDAGV....VSSSKIQLARD...A.NGLWTCSTTV.TSEIAPKGC 146
NTDB id 1038 HON27 RS01510 WP 000993725.1 ...DVGFTAATSSVCATYSIDAFTETDGAGKITCTLKG...NPKIATKIVELTRS...T.DGAWTCVSDV.PTEFLPKGC 136
NTDB id 1292 VP RS12235 WP 005479693.1 TLAQLGVDVAA.NDLGAIDS.AL..AGGSPTLTFTFDA...DSSIDGGILTFTRD...A.ATGWSCLPTA.PAIA.VDGC 138
NTDB id 1197 PSJM300 03940 AFN76866.1 DEGYIGITDST.SYCDV.DL.D...TAADGHIECT.AKGGNAGKFDGKTITLNRT...A.DGEWSCASTL.DAKYKPGKC 138
NTDB id 1204 PSJM300 03935 AFN76865.1 LPGFIGIGAST.NYCTV.TL.I...ATATGSITCT.TQNGMPDRFNGKTITTART...T.DGLWDCTSDL.DSRYKPGKC 140
NTDB id 1038 HON27 RS01510 WP 000993725.1 DTTYVGVDATA.NKLGTIAV.TIG.NNGAGTVTFTFAAGNSSPKNSGKIITLTRT...D.AGSWACTSTQ.DEMFIPKGC 147
NTDB id 1192 PAKAF RS23870 WP 024947914.1 KEVPLGVAADA.NKLGTIAL.KPDPADGTADITLTFTMGGAGPKNKGKIITLTRT...AADGLWKCTSDQ.DEQFIPKGC 148
consensus
logo
NTDB id 1002 ACIAD RS15100 WP 004923779.1 RG.......... 147
NTDB id 1258 GCO85 RS09745 WP 014844212.1 RP.......... 136
NTDB id 1274 LPP RS09530 WP 011214168.1 RP.......... 136
NTDB id 1184 GTF74 RS02335 WP 000649326.1 TAGATIN..... 153
NTDB id 1167 A1552VC RS11065 WP 000649326.1 TAGATIN..... 153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1 TAGATIN..... 153
NTDB id 1038 HON27 RS01510 WP 000993725.1 TAGTPTEGAITSL 149
NTDB id 1292 VP RS12235 WP 005479693.1 S.......... 139
NTDB id 1197 PSJM300 03940 AFN76866.1 S.......... 139
NTDB id 1204 PSJM300 03935 AFN76865.1 L.......... 141
NTDB id 326307 EGY27 RS07255 WP 024928732.1 KAP......... 150
NTDB id 1192 PAKAF RS23870 WP 024947914.1 SK.......... 150
consensus
non conserved
similar
≥ 50% conserved