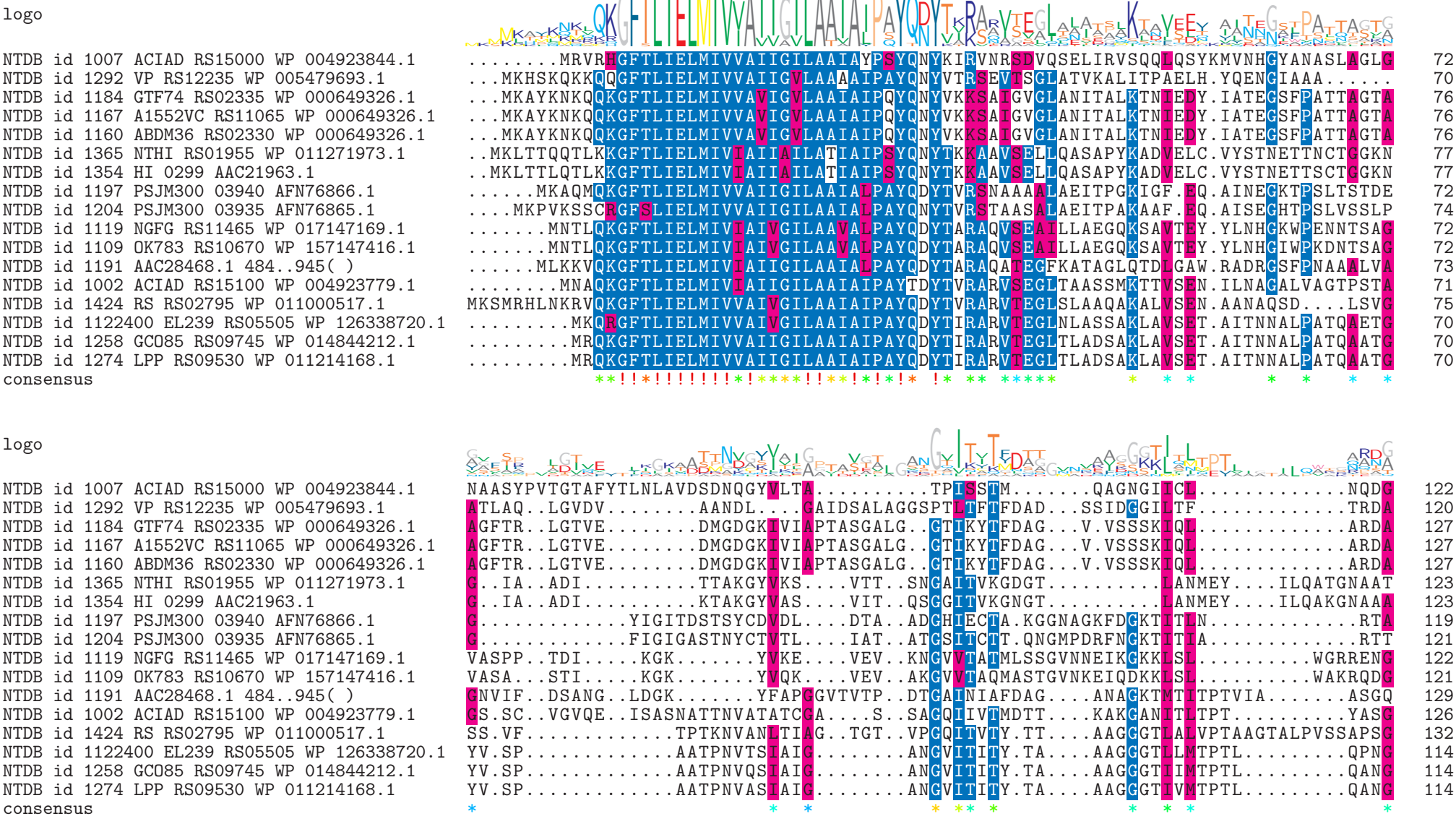
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NTDB id 1292 VP RS12235 WP 005479693.1 ...MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELH.YQENGIAAA..... 70
NTDB id 1184 GTF74 RS02335 WP 000649326.1 ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDY.IATEGSFPATTAGTA 76
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NTDB id 1197 PSJM300 03940 AFN76866.1 ......MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGF.EQ.AINEGKTPSLTSTDE 72
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NTDB id 1002 ACIAD RS15100 WP 004923779.1 ........MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSEN.ILNAGALVAGTPSTA 71
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consensus
logo
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NTDB id 1292 VP RS12235 WP 005479693.1 ATLAQ..LGVDV.........AANDL....GAIDSALAGGSPTLTFTFDAD...SSIDGGILTF.............TRDA 120
NTDB id 1184 GTF74 RS02335 WP 000649326.1 AGFTR..LGTVE.........DMGDGKIVIAPTASGALG..GTIKYTFDAG...V.VSSSKIQL.............ARDA 127
NTDB id 1167 A1552VC RS11065 WP 000649326.1 AGFTR..LGTVE.........DMGDGKIVIAPTASGALG..GTIKYTFDAG...V.VSSSKIQL.............ARDA 127
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NTDB id 1365 NTHI RS01955 WP 011271973.1 G..IA..ADI...........TTAKGYVKS....VTT..SNGAITVKGDGT...........LANMEY....ILQATGNAAT 123
NTDB id 1354 HI 0299 AAC21963.1 G..IA..ADI...........KTAKGYVAS....VIT..QSGGITVKGNGT...........LANMEY....ILQAKGNAAA 123
NTDB id 1197 PSJM300 03940 AFN76866.1 G..............YIGITDSTSYCDVDL....DTA..ADGHIECTA.KGGNAGKFDGKTITLN.............RTA 119
NTDB id 1204 PSJM300 03935 AFN76865.1 G..............FIGIGASTNYCTVTL....IAT..ATGSITCTT.QNGMPDRFNGKTITIA.............RTT 121
NTDB id 1119 NGFG RS11465 WP 017147169.1 VASPP..TDI.....KGK.......YVKE....VEV..KNGVVTATMLSSGVNNEIKGKKLSL...........WGRRENG 122
NTDB id 1109 OK783 RS10670 WP 157147416.1 VASA...STI.....KGK.......YVQK....VEV..AKGVVTAQMASTGVNKEIQDKKLSL...........WAKRQDG 121
NTDB id 1191 AAC28468.1 484..945( ) GNVIF..DSANG..LDGK.......YFAPGGVTVTP..DTGAINIAFDAG....ANAGKTMTITPTVIA.......ASGQ 129
NTDB id 1002 ACIAD RS15100 WP 004923779.1 GS.SC..VGVQE..ISASNATTNVATATCGA....S..SAGQIIVTMDTT....KAKGANITLTPT..........YASG 126
NTDB id 1424 RS RS02795 WP 011000517.1 SS.VF.............TPTKNVANLTIAG..TGT..VPGQITVTY.TT....AAGGGTLALVPTAAGTALPVSSAPSG 132
NTDB id 1122400 EL239 RS05505 WP 126338720.1 YV.SP.............AATPNVTSIAIG.........ANGVITITY.TA....AAGGGTLLMTPTL.........QPNG 114
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NTDB id 1274 LPP RS09530 WP 011214168.1 YV.SP.............AATPNVASIAIG.........ANGVITITY.TA....AAGGGTIVMTPTL.........QANG 114
consensus

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NTDB id 1292 VP RS12235 WP 005479693.1        ATGWSCLPTA.P..................AIAVDGCS........   139
NTDB id 1184 GTF74 RS02335 WP 000649326.1     NGLWTCSTTVTS..................EIAPKGCTAGATIN.   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   NGLWTCSTTVTS..................EIAPKGCTAGATIN.   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    NGLWTCSTTVTS..................EIAPKGCTAGATIN.   153
NTDB id 1365 NTHI RS01955 WP 011271973.1      GVTWTTTCKGT..................DASLFPANFCGSVTQ..   149
NTDB id 1354 HI 0299 AAC21963.1               GVTWTTTCKGT..................DASLFPANFCGSVTK..   149
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NTDB id 1119 NGFG RS11465 WP 017147169.1      SVKWFCGQPVTRADDDTVADAKDGKEIDTKHLPSTCRDKASDAK   166
NTDB id 1109 OK783 RS10670 WP 157147416.1     SVKWFCGQPVTRTGDNDDTVADANNAIDTKHLPSTCRDESSAT.   164
NTDB id 1191 AAC28468.1 484..945( )           ISTWTCAPGAAN..............GIETRRLPTSCQ*.......   153
NTDB id 1002 ACIAD RS15100 WP 004923779.1     AVTWKCTTTS...................DKKYVPSECRG.......   147
NTDB id 1424 RS RS02795 WP 011000517.1        PIMWTCYAQNKAQAAS.SVAPSGTMSLAAKYAPAECR........   168
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NTDB id 1258 GCO85 RS09745 WP 014844212.1     DVTWTCT.GG.................TLLAKYRPASCRP.......   136
NTDB id 1274 LPP RS09530 WP 011214168.1       DVTWTCT.GG.................TLLAKYRPASCRP.......   136
consensus                                        !**                      *   *   *
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X non conserved

X similar

X ≥ 50% conserved