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NTDB id 1283 ACE17W RS02630 WP 010947729.1   ................................................................................    0
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NTDB id 1194 PAKAF RS01995 WP 003084552.1    ................................................................................    0
NTDB id 1202 PSJM300 01600 AFN76400.1        ................................................................................    0
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NTDB id 1203 PSJM300 01605 AFN76401.1        ..............................MEFEKLLRLMVEKGGSDLFITA...GVPPSMKVNGKILPVNKTPMSPE    45
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NTDB id 1307 DR RS10055 WP 010888596.1       .........................MTSGIDITDILRLAADKGASDIILAA...GLPPQFKLSGDYDAQGFEPLSAS    49
NTDB id 1086 NMV RS00245 WP 002218487.1      ..............................MQITDLLAFGAKNKASDLHLSS...GISPMIRVHGDMRRINLPEMSAE    45
NTDB id 1116 NGFG RS10615 WP 003690200.1     ..............................MQITDLLAFGAKNKASDLHLSS...GISPMIRVHGDMRRINLPEMSAE    45
NTDB id 1185 GTF74 RS11940 WP 000350195.1    ..............................MDIAELLEFSVKHNASDLHLSA...GVPPMVRIDGEVRKLGVPAFTHS    45
NTDB id 1176 A1552VC RS01160 WP 000350195.1  ..............................MDIAELLEFSVKHNASDLHLSA...GVPPMVRIDGEVRKLGVPAFTHS    45
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NTDB id 1060 ABD1 RS04380 WP 000355489.1     ..............................MDITELLAFSVKNGASDLHLSA...GMPPMIRVDGEVRRINLPALEHK    45
NTDB id 1040 HON27 RS13360 WP 000355489.1    ..............................MDITELLAFSVKNGASDLHLSA...GMPPMIRVDGEVRRINLPALEHK    45
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NTDB id 1086 NMV RS00245 WP 002218487.1      EVGNMVTSVMNDHQRKI..YQQNLEVDFSFELPN.VARFRVNAFNIGRGPAAVFRTIPSTVL..SLEELKAPSI....FQ   116
NTDB id 1116 NGFG RS10615 WP 003690200.1     EVGNMVTSVMNDHQRKI..YQQNLEVDFSFELPN.VARFRVNAFNTGRGPAAVFRTIPSTVL..SLEELKAPSI....FQ   116
NTDB id 1185 GTF74 RS11940 WP 000350195.1    DVHRLIFEIMNDAQRSE..YEEKLEVDFSFELPN.VGRFRVNAFHQARGCSAVFRTIPTVIP..TLEQLDAPEI....FS   116
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NTDB id 1202 PSJM300 01600 AFN76400.1        QVHALIYDIMNDKQRKD..FEEFLETDFSFEVPG.VARFRVNAFNQNRGAGAVFRTIPSKVL..TMEDLGMGEV....FR   116
NTDB id 1020 ACIAD RS04210 WP 011182164.1    EVHNLIYDIMNDKQRRD..YEEKLETDFSFEVPN.LARFRVNVFNQNRGAGAVFRTIPSQVL..TLEELGLGKI....FR   116
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NTDB id 1060 ABD1 RS04380 WP 000355489.1     DVHRLVYDIMNDKQRRD..YEEKLETDFSFEVPN.VARFRVNAFNQNRGAGAVFRTIPSKVL..TMEDLGLGQI....FK   116
NTDB id 1040 HON27 RS13360 WP 000355489.1    DVHRLVYDIMNDKQRRD..YEEKLETDFSFEVPN.VARFRVNAFNQNRGAGAVFRTIPSKVL..TMEDLGLGQI....FK   116
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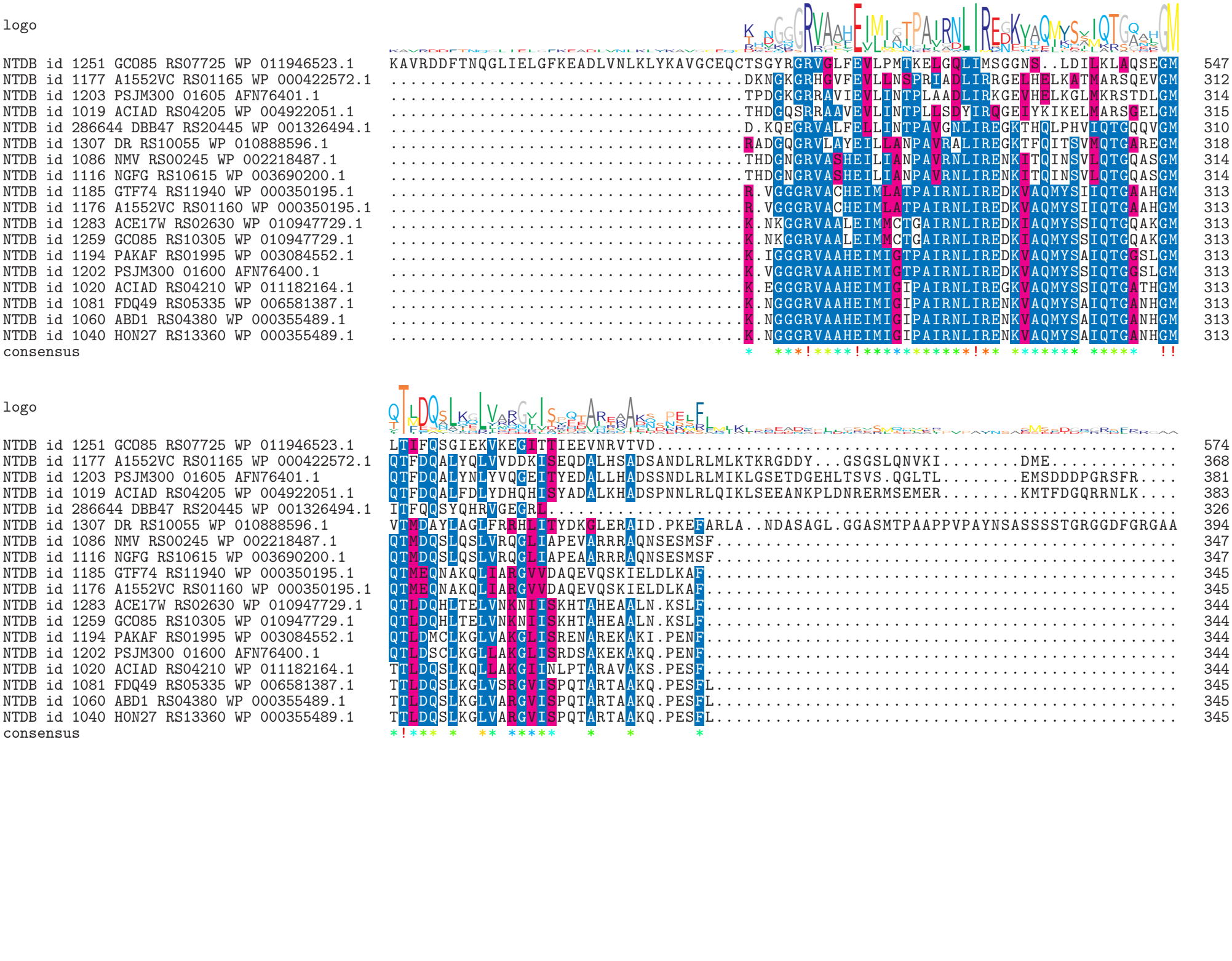


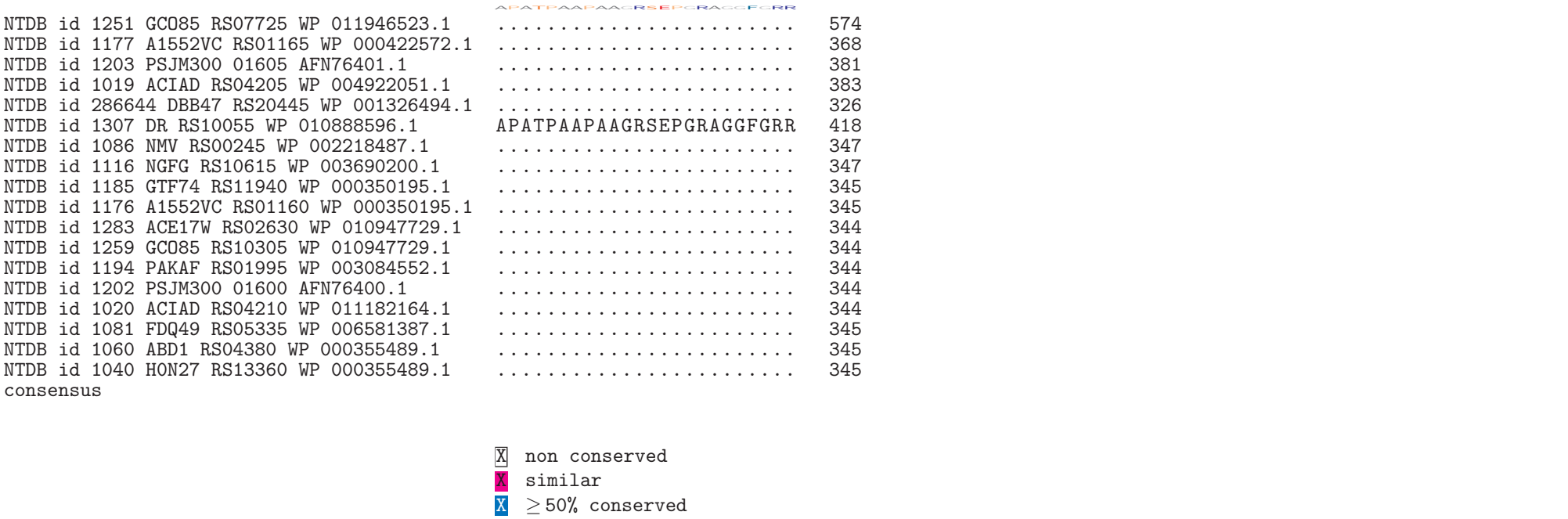
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NTDB id 1019 ACIAD RS04205 WP 004922051.1    ELAMSKRGIIIFSGATGTGKSTSLASMIQHRNQHSKGHIITIEDPIEFIHEHAGCIITQREVGIDTDSFEIALKNTLRQA  197
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NTDB id 1116 NGFG RS10615 WP 003690200.1     KIAESPRGMVLVTGPTGSGKSTTLAAMINYINETQPAHILTIEDPIEFVHQSKKSLINQRELHQHTLSFANALSSALRED  196
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NTDB id 1176 A1552VC RS01160 WP 000350195.1  KIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHILTIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALRED  196
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NTDB id 1040 HON27 RS13360 WP 000355489.1    DICDYPRGIVLVTGPTGSGKSTTLAAMLDYINENRYDHILTVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALRED  196
consensus                                     **  **!******!*!!*!!*********  **!  *  *!***!!!*!*** ******************  *!***!!**
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NTDB id 1177 A1552VC RS01165 WP 000422572.1  PDMILIGEIRSRETMEYAMTFAETGHLCMATLHANNANQALERILHLVPKEQREQFLLDLSLNLKGVIAQQLLR......  268
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consensus                                     !!*!**!!*!*  !****!***!*!!!!***!!!***!*****!**************** ******  *****!  !**
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consensus
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consensus


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consensus
X non conserved
X similar
X ≥ 50% conserved