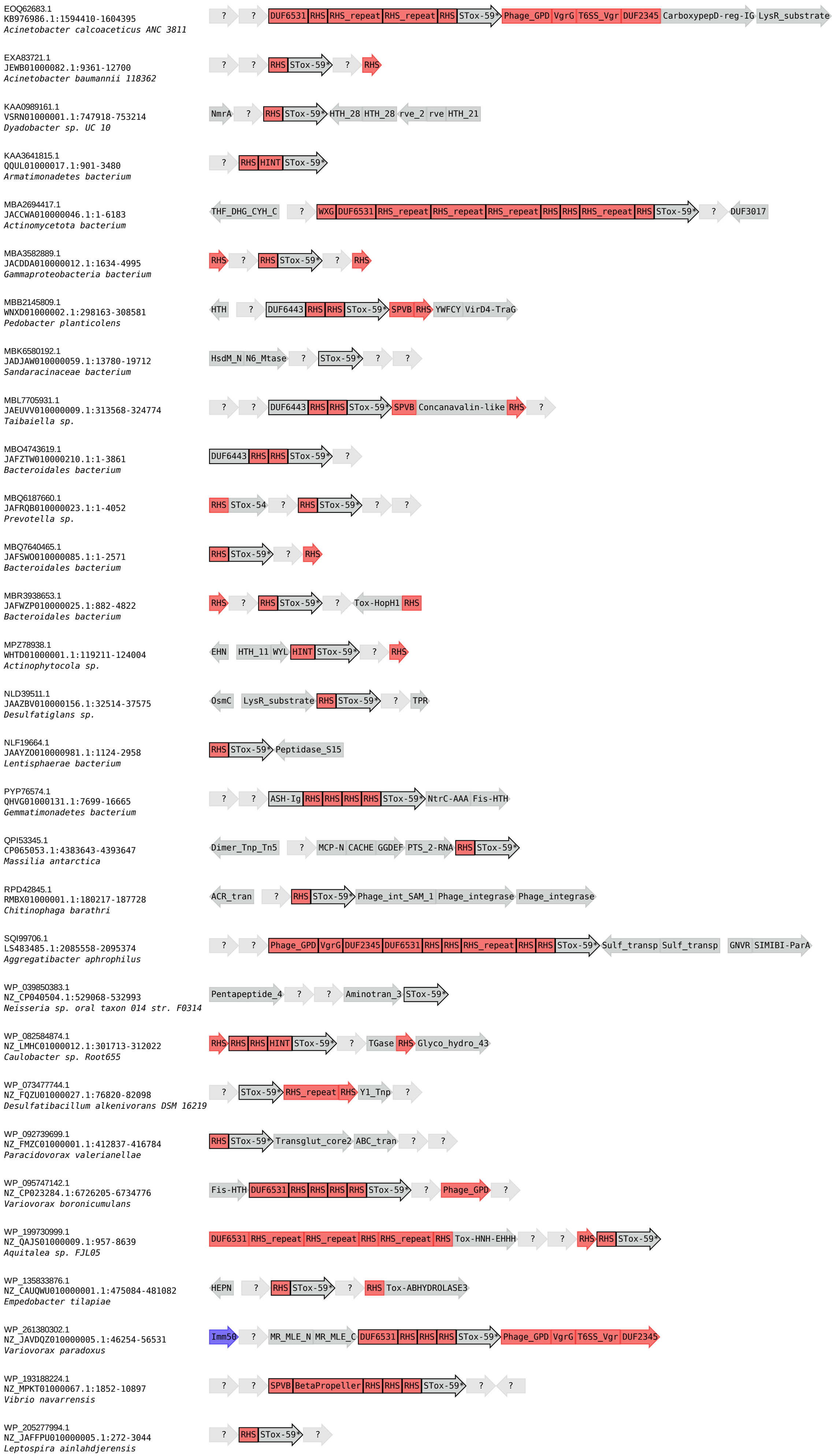
EOQ62683.1
KB976986.1:1594410-1604395
Acinetobacter calcoaceticus ANC 3811
? ? DUF6531 RHS RHS_repeat RHS_repeat RHS STox-59* Phage_GPD VgrG T6SS_Vgr DUF2345 CarboxypepD-reg-IG LysR_substrate
EXA83721.1
JEWB01000082.1:9361-12700
Acinetobacter baumannii 118362
? ? RHS STox-59* ? RHS
KAA0989161.1
VSRN01000001.1:747918-753214
Dyadobacter sp. UC 10
NmrA ? RHS STox-59* HTH_28 HTH_28 rve_2 rve HTH_21
KAA3641815.1
QQUL01000017.1:901-3480
Armatimonadetes bacterium
? RHS HINT STox-59*
MBA2694417.1
JACCWA010000046.1:1-6183
Actinomycetota bacterium
THF_DHG_CYH_C ? WXG DUF6531 RHS_repeat RHS_repeat RHS_repeat RHS RHS RHS_repeat RHS STox-59* ? DUF3017
MBA3582889.1
JACDDA010000012.1:1634-4995
Gammaproteobacteria bacterium
RHS ? RHS STox-59* ? RHS
MBB2145809.1
WNXD01000002.1:298163-308581
Pedobacter planticolens
HTH ? DUF6443 RHS RHS STox-59* SPVB RHS YWFCY VirD4-TraG
MBK6580192.1
JADJAW010000059.1:13780-19712
Sandaracinaceae bacterium
HsdM_N N6_Mtase ? STox-59* ? ?
MBL7705931.1
JAEUVV010000009.1:313568-324774
Taibaiella sp.
? ? DUF6443 RHS RHS STox-59* SPVB Concanavalin-like RHS ?
MBO4743619.1
JAFZTW010000210.1:1-3861
Bacteroidales bacterium
DUF6443 RHS RHS STox-59* ?
MBQ6187660.1
JAFRQB010000023.1:1-4052
Prevotella sp.
RHS STox-54 ? RHS STox-59* ? ?
MBQ7640465.1
JAFSWO010000085.1:1-2571
Bacteroidales bacterium
RHS STox-59* ? RHS
MBR3938653.1
JAFWZP010000025.1:882-4822
Bacteroidales bacterium
RHS ? RHS STox-59* ? Tox-HopH1 RHS
MPZ78938.1
WHTD01000001.1:119211-124004
Actinophytocola sp.
EHN HTH_11 WYL HINT STox-59* ? RHS
NLD39511.1
JAAZBV010000156.1:32514-37575
Desulfatiglans sp.
OsmC LysR_substrate RHS STox-59* ? TPR
NLF19664.1
JAAYZO010000981.1:1124-2958
Lentisphaerae bacterium
RHS STox-59* Peptidase_S15
PYP76574.1
QHVG01000131.1:7699-16665
Gemmatimonadetes bacterium
? ? ASH-Ig RHS RHS RHS RHS STox-59* NtrC-AAA Fis-HTH
QPI53345.1
CP065053.1:4383643-4393647
Massilia antarctica
Dimer_Tnp_Tn5 ? MCP-N CACHE GGDEF PTS_2-RNA RHS STox-59*
RPD42845.1
RMBX01000001.1:180217-187728
Chitinophaga barathri
ACR_tran ? RHS STox-59* Phage_int_SAM_1 Phage_integrase Phage_integrase
SQI99706.1
LS483485.1:2085558-2095374
Aggregatibacter aphrophilus
? ? Phage_GPD VgrG DUF2345 DUF6531 RHS RHS RHS_repeat RHS RHS STox-59* Sulf_transp Sulf_transp GNVR SIMIBI-ParA
WP_039850383.1
NZ_CP040504.1:529068-532993
Neisseria sp. oral taxon 014 str. F0314
Pentapeptide_4 ? ? Aminotran_3 STox-59*
WP_082584874.1
NZ_LMHC01000012.1:301713-312022
Caulobacter sp. Root655
RHS RHS RHS HINT STox-59* ? TGase RHS Glyco_hydro_43
WP_073477744.1
NZ_FQZU01000027.1:76820-82098
Desulfatibacillum alkenivorans DSM 16219
? STox-59* RHS_repeat RHS Y1_Tnp ?
WP_092739699.1
NZ_FMZC01000001.1:412837-416784
Paracidovorax valerianellae
RHS STox-59* Transglut_core2 ABC_tran ? ?
WP_095747142.1
NZ_CP023284.1:6726205-6734776
Variovorax boronicumulans
Fis-HTH DUF6531 RHS RHS RHS RHS STox-59* ? Phage_GPD ?
WP_199730999.1
NZ_QAJS01000009.1:957-8639
Aquitalea sp. FJL05
DUF6531 RHS_repeat RHS_repeat RHS RHS_repeat RHS Tox-HNH-EHHH ? ? RHS RHS STox-59*
WP_135833876.1
NZ_CAUQWU010000001.1:475084-481082
Empedobacter tilapiae
HEPN ? RHS STox-59* ? RHS Tox-ABHYDROLASE3
WP_261380302.1
NZ_JAVDQZ010000005.1:46254-56531
Variovorax paradoxus
Imm50 ? MR_MLE_N MR_MLE_C DUF6531 RHS RHS RHS STox-59* Phage_GPD VgrG T6SS_Vgr DUF2345
WP_193188224.1
NZ_MPKT01000067.1:1852-10897
Vibrio navarrensis
? ? SPVB BetaPropeller RHS RHS RHS STox-59* ? ?
WP_205277994.1
NZ_JAFFPU010000005.1:272-3044
Leptospira ainlahdjerensis
? RHS STox-59* ?