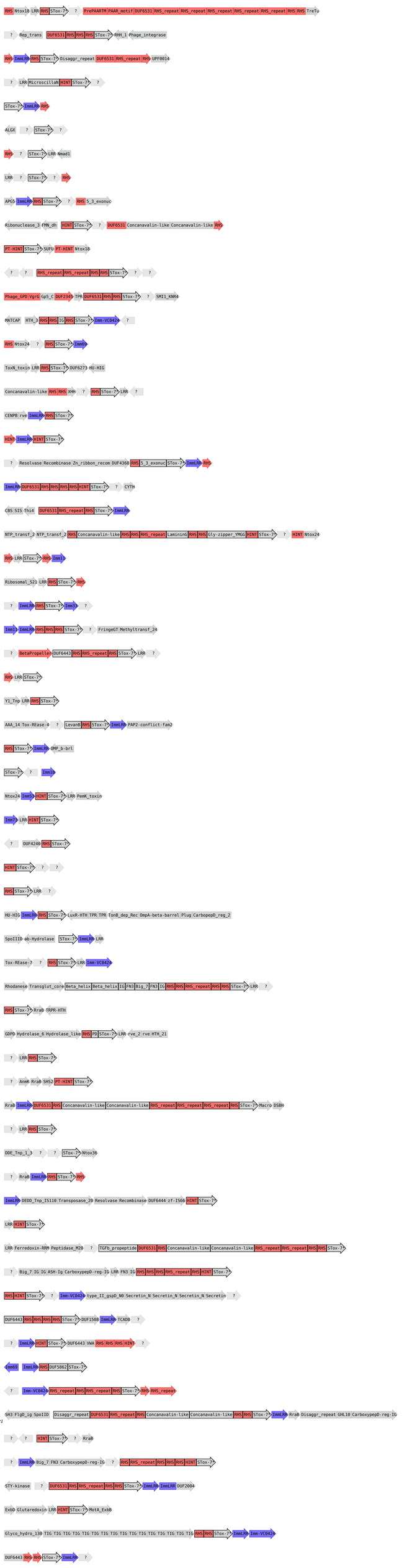
BBN78551.1
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Salmonella enterica
RHS Ntox18 LRR RHS STox-? ? PrePAARTM PAAR_motif DUF6531 RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS RHS TreTo
GJM17949.1
BGJQ01000005.1:14512-26246
Phycisphaeraceae bacterium
? Rep_trans DUF6531 RHS RHS RHS STox-? RHH_1 Phage_integrase
GKV68547.1
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Sporosarcina sp. NCCP-2716
RHS ImmLRb RHS STox-? Disaggr_repeat DUF6531 RHS_repeat RHS UPF0014
HAS47434.1
DMXI01000687.1:65067-70793
Microscillaceae bacterium
? LRR MicroscillaN HINT STox-? ?
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Peptococcaceae bacterium
STox-? ImmLRb RHS
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Chryseobacterium indologenes
ALGX ? STox-? ?
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Clostridiaceae bacterium
RHS ? STox-? LRR Nmad1
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JACSXZ010000003.1:9010-10842
Flavobacterium piscinae
LRR ? STox-? ? RHS
MBE5956217.1
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Lachnospiraceae bacterium
APG5 ImmLRb RHS STox-? ? RHS 5_3_exonuc
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Oscillospiraceae bacterium
Ribonuclease_3 PMN_dh HINT STox-? ? DUF6531 Concanavalin-like Concanavalin-like RHS
MBG8037269.1
WBL001000064.1:1-2914
Neisseria meningitidis
PT-HINT STox-? SUFU PT-HINT Ntox18
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Polyangiaceae bacterium
? ? RHS_repeat RHS_repeat RHS RHS STox-? ? ?
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JADJYE010000018.1:101394-110030
Polyangiaceae bacterium
Phage_GPD VgrG Gp5_C DUF2345 TPR DUF6531 RHS RHS STox-? ? SMI1_KNR4
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JAFLDX010000004.1:24565-35578
Lysobacterales bacterium
MATCAP HTH_3 RHS RHS IG RHS STox-? Imm-VC0842 ?
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JAFKLP010000003.1:1-3949
Sphingobacteriales bacterium
RHS Ntox24 ? RHS STox-? Imm6
MBO6132145.1
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Treponema sp.
ToxN_toxin LRR RHS STox-? DUF6273 HU-HIG
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Clostridia bacterium
Concanavalin-like RHS RHS XHH ? RHS STox-? LRR ?
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Agrobacterium sp.
CENPB rve ImmLRb RHS STox-?
MBQ1327014.1
JAFOIV010000046.1:1-2807
Eubacterium sp.
HINT ImmLRb HINT STox-?
MBQ4560127.1
JAFQDE010000021.1:1486-5888
Tyzzerella sp.
? Resolvase Recombinase Zn_ribbon_recom DUF4368 RHS 5_3_exonuc STox-? ImmLRb RHS
MBQ7106792.1
JAFSDK010000038.1:172-9514
Clostridia bacterium
ImmLRb DUF6531 RHS RHS RHS RHS HINT STox-? ? CYTH
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Verrucomicrobia bacterium
CBS SIS Thi4 DUF6531 RHS_repeat RHS STox-? ImmLRb
MBR2138161.1
JAFVNC010000040.1:20159-31142
Bacilli bacterium
NTP_transf_2 NTP_transf_2 RHS Concanavalin-like RHS RHS RHS_repeat LaminiNG RHS RHS Gly-zipper_YMGG HINT STox-? ? HINT Ntox24
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Haemophilus haemolyticus
RHS LRR STox-? RHS Imm1
MBS7278195.1
JAHBAB010000164.1:16844-20271
Bacteroidales bacterium
Ribosomal_S21 LRR RHS STox-? RHS
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JAFAJM010000018.1:137748-142916
Chryseobacterium sp.
? ImmLRb RHS STox-? Imm3 ?
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JAGQOC010000059.1:1069-11388
Planctomycetaceae bacterium
Imm1 ImmLRb RHS RHS RHS STox-? ? FringeGT Methyltransf_24
MCC6327363.1
JADLDC010000072.1:972-10645
Acidobacteria bacterium
? BetaPropeller DUF6443 RHS RHS_repeat RHS STox-? LRR ?
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Sphingobacteriia bacterium
RHS LRR STox-?
MCH6255511.1
JAKYXN010000005.1:128240-131727
Puniceicoccaceae bacterium K14
Y1_Tnp LRR RHS STox-?
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JALEJC010000016.1:11814-17667
Acholeplasmataceae bacterium
AAA_14 Tox-REase-4 ? LevanB RHS STox-? ImmLRb PAP2-conflict-fam2
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Psychrosphaera sp.
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JAKLJZ010001494.1:1-1956
Polyangiaceae bacterium
STox-? ? Imm10
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Ruminococcus flavefaciens
Ntox24 Imm3 HINT STox-? LRR PemK_toxin
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JAMPAW010000315.1:436-3248
Clostridium sp.
Imm7 LRR HINT STox-?
MCP4162099.1
JAAENH010000338.1:1173-3072
Deltaproteobacteria bacterium
? DUF4240 RHS STox-?
NIT56939.1
JAACBR010000737.1:141-1817
Fodinibius sp.
HINT STox-? ? ?
NJK32366.1
JAAUUP010000261.1:1-2710
Deltaproteobacteria bacterium
RHS STox-? LRR ?
NJN25814.1
JAAUQL010000021.1:3579-11834
Cyclobacteriaceae bacterium
HU-HIG ImmLRb RHS STox-? LuxR-HTH TPR TPR TonB_dep_Rec OmpA-beta-barrel Plug CarbopepD_reg_2
NLN03257.1
JAAYLF010000009.1:72-3706
Clostridiaceae bacterium
SpoIIID ab-Hydrolase STox-? ImmLRb LRR
OFC77819.1
LXLF01000015.1:41794-45600
Bacillus thuringiensis
Tox-REase-7 ? RHS STox-? LRR Imm-VC0842
OGV56663.1
MHBI01000069.1:3416-22049
Lentisphaerae bacterium GWF2_50_93
Rhodanese Transglut_core Beta_helix Beta_helix IG FN3 Big_7 FN3 IG RHS RHS RHS_repeat RHS RHS STox-? LRR ?
PZO47644.1
QBMO01000125.1:1-1658
Alphaproteobacteria bacterium
RHS STox-? RraB TRPR-HTH
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uncultured Lachnospira sp.
GDPD Hydrolase_6 Hydrolase_like RHS PD STox-? LRR rve_2 rve HTH_21
TAK85242.1
SCTN01000535.1:4856-8524
Aquabacterium sp.
? LRR RHS STox-?
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Neisseria meningitidis 69166
? AnmK RraB SHS2 PT-HINT STox-?
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Bhargavaea cecembensis DSE10
RraB ImmLRb DUF6531 RHS Concanavalin-like Concanavalin-like RHS_repeat RHS_repeat RHS STox-? Macro DSBH
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Halobacillus contractile SSD-17B
? LRR RHS STox-?
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Geobacillus sp. GHH01
DDE_Tnp_1_3 ? ? STox-? Ntox36
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Bhargavaea cecembensis
? RraB ImmLRb RHS STox-? RHS
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Gorillibacterium massiliense
ImmLRb DEDD_Tnp_IS110 Transposase_20 Resolvase Recombinase DUF6444 zf-IS66 HINT STox-?
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NZ_AZOP01000122.1:1-1275
Fervidicella metallireducens AeB
LRR HINT STox-?
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Virgibacillus pantothenticus
LRR Ferredoxin-RRM Peptidase_M20 ? TGFb_propeptide DUF6531 RHS Concanavalin-like Concanavalin-like RHS_repeat RHS_repeat RHS RHS STox-?
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Piscinibacter gummiphilus
? Big_7 IG IG ASM-Ig CarboxypepD-reg-IG LRR FN3 IG RHS RHS RHS RHS_repeat RHS HINT STox-?
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Hoeflea marina
RHS HINT STox-? ? Imm-VC0842 type_II_gspD_N0 Secretin_N Secretin_N Secretin_N Secretin ?
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Seonamhaeicola aphaedonensis
DUF6443 RHS RHS RHS RHS STox-? DUF1508 ImmLRb TCADB ?
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Hyunsoonleella pacifica
? ImmLRb HINT STox-? DUF6443 VWA RHS RHS RHS HINT ?
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Pedobacter hiemivivus
Imm6 ImmLRb RHS DUF5862 STox-?
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Asticcacaulis tiandongensis
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Peribacillus simplex NBRC 15720 = DSM 1321
SH3 FlgD_ig SpoIID Disaggr_repeat DUF6531 RHS_repeat RHS Concanavalin-like Concanavalin-like RHS RHS STox-? ImmLRb RraB Disaggr_repeat GHL10 CarboxypepD-reg-IG
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Pirellulimonas nuda
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Pseudoxanthomonas suwonensis
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Pseudenhygromyxa sp. WMMC2535
STY-kinase ? DUF6531 RHS RHS_repeat RHS RHS STox-? ImmLRb ImmLRR DUF2004
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Deefgea piscis
Exbb Glutaredoxin LRR HINT STox-? MotA_ExbB
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NZ_CP063849.1:7504757-7516799
Paludibaculum fermentans
Glyco_hydro_130 TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG TIG RHS RHS STox-? ImmLRb Imm-VC0842
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Chryseobacterium sp. C-71
DUF6443 RHS RHS STox-? ImmLRb ?