ADN72587.1
CP002167.1:3347388-3352657
Escherichia coli UM146
DDE_Tnp_1_assoc
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Peptidase-C39
DUF2169
Phage_GPD
VgrG
Gp5_C
AGY57409.1
CP003587.1:1189896-1193251
Gloeobacter kilauweensis JS1
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ZnR
OrfB_IS605
HTH_OrfB_IS605
Peptidase_C70
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PfkB
CDL84571.1
CBXF010000109.1:21918-23654
Xenorhabdus szentirmaii DSM 16338
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Peptidase_C70
HKX00204.1
DVDU01000034.1:4687-7899
Oscillatoriaceae cyanobacterium M33_DOE_052
ADH_N
ADH_zinc_N
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Complex1_30kDa
Oxidored_q6
MBA39238442.1
JACDGJ010000870.1:1-2444
Nostocaceae cyanobacterium
NmrA
Peptidase-C39
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SbcC
MBA3991986.1
PMLND1000010.1:17585-25003
Cyanobacteria bacterium OSZ.3.42
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SNF
SWIM
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Cyanobacteria bacterium PR.023
PepSY
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Peptidase_C70
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JACONA010000435.1:258-1044
Hormoscilla sp. SP5CH51
Peptidase_C70
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MBC6472105.1
JACOMF010000091.1:722-2246
Hormoscilla sp. GM102CH51
Peptidase-C39
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Uma2
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JACOMF010000463.1:4566-9769
Hormoscilla sp. GM102CH51
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Peptidase_C39
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tRNA_SAD
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Hormoscilla sp. GM7CH51pb
Uma2
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Peptidase_C70
Uma2
DUF433
MBC8233259.1
JACNLI010000859.1:3706-6793
bacterium Bacteria.
ZnR
RDD-single
GAF
PP2C
His-Kinase
Peptidase-C39
Mal_decarbox_A1
His-Kinase
MBE8589983.1
JADOEM010000624.1:13465-17604
Pseudomonas cyclaminis
Tox-SHH
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Peptidase-C39
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HEPN
MBF2085751.1
JACYLP010000045.1:92268-96298
Thermoleptolyngbya sp. C42_A2020_037
Phagetail-S
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Peptidase_C70
Acetylase
cNMP-cyclase
cNMP-cyclase
MBI1271343.1
WGMS01000020.1:153238-155063
bacterium Bacteria.
HPPA
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MBI5175729.1
JACRFZ010000106.1:7395-13712
Candidatus Melainabacteria bacterium
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STY-kinase
MBK6685901.1
JADJ0V010000001.1:3926723-3939323
Deltaproteobacteria bacterium
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McrBC
DUF6531
RHS
Furin-like
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RHS
RHS
Peptidase-C39
McrB-AAA
McrC-NTD
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MBK8303596.1
JADJRV010000010.1:358-2004
Chloracidobacterium sp.
Peptidase-C39
TIR
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MBN3872442.1
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Nostoc sp. JL33
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RHS_repeat
RHS_repeat
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Peptidase-C39
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JAGHZP010000116.1:1-2848
Cyanobacteria bacterium SID2
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Peptidase-C39
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JAGNCZ010000002.1:157315-160580
bacterium Bacteria.
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DUF428
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JAGPDR010000199.1:1111-4674
bacterium Bacteria.
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STY-kinase
MBS5193497.1
JAFEAV010000082.1:20036-24771
Cyanobacteria bacterium SZAS LIN-3
Glyco_hydro_125
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Peptidase-C39
Deaminase
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Cyanobacteria bacterium REEB67
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Peptidase_C70
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JAFAZY010000290.1:217-2609
Pseudonocardia sp.
Peptidase-C39
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JAHHCI010000042.1:65906-70698
Scytonematopsis contorta HA4267-MV1
ISOCO7
RtcB
Peptidase_C39
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JAHRPM010000015.1:14572-20013
Candidatus Obscuribacterales bacterium
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JAHBSK010000001.1:136025-139850
Acidobacteria bacterium
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Peptidase-C39
CTP_transf_1
Prenyltransf
MBX3295921.1
JAHBLG010000010.1:29381-32232
Acidobacteria bacterium
AlkZ-like
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Peptidase-C39
dnstrm_HI1420
Gp49
MBX9690048.1
JAIEPM010000590.1:9894-13180
Candidatus Obscuribacterales bacterium
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Peptidase_C39
AAA_28
MBX9721591.1
JAIERA010000658.1:5415-15147
Candidatus Obscuribacterales bacterium
HTH_17
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Peptidase-C39
MshEN
DUF4097
MCA1597885.1
JAIVJW010000156.1:1-1925
Chloroflexota bacterium
HAD
Peptidase_C70
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Tautomerase_2
MCA9814417.1
JAGQMG010000020.1:12959-19059
Cyanobacteria bacterium HKST-UBA01
ab-Hydrolase
TPR_TPR
Peptidase_C70
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MCA9817803.1
JAGQMG010000307.1:1-5152
Cyanobacteria bacterium HKST-UBA01
Aminotran_3
Peptidase_C70
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Beta-lactamase
MCA9818507.1
JAGQMG010000475.1:1-2530
Cyanobacteria bacterium HKST-UBA01
Peptidase_C70
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RsaC
MCA9898966.1
JAGQGG010000113.1:29909-34981
Anaerolineales bacterium
HTH_11
WYL
Aminotran_1_2
RHS
Peptidase_C70
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BetaPropeller
MCC6980443.1
JADYYD010000384.1:2360-7662
Candidatus Melainabacteria bacterium
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ABC-ATPase
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MCC7384259.1
JAD2E501000062.1:138982-145557
Deltaproteobacteria bacterium
CP_ATPgrasp_2
ab-Hydrolase
Peptidase_C70
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AIRS_C
AIRS
MCK9815296.1
JALCX010000026.1:1-6635
Pseudomonas morbosilactucae
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Fil_haemagg_2
Fil_haemagg_2
Fil_haemagg_2
Fil_haemagg_2
DUF637
PT-VENN
Peptidase_C39
SBP_bac_3
BPD_transp_1
MCL5508530.1
JAMDDR010000445.1:58-4424
Chloroflexota bacterium
Uma2
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Peptidase_C70
RHS
ABC_tran
MCP2727200.1
JAMCPM010000009.1:847-2663
Limnofasciculus baicalensis BBK-W-15
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Rhd-YefM
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MCP9493104.1
JANDLT010000003.1:46196-51792
Pyrinomonadaceae bacterium MAG19_C2-C3
COX15-CtaA
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Peptidase-C39
UvdB
Beta-Prop
Peptidase_S9
NEP12619.1
JAAHLG010000188.1:317-2583
Symploca sp. SIO2C1
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Peptidase-C39
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TrapB
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NJK50651.1
JAAXUT010000128.1:192-1845
Candidatus Gracilibacteria bacterium
Peptidase-C39
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Phage_integrase
NJM54188.1
JAAURM010000397.1:130-3267
Blastocatellia bacterium
FlaToC
OmpR-HTH
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Peptidase_C39
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JABSPE010000703.1:38-719
Crocosphaera sp.
Peptidase-C39
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OCQ00298.1
MBR010000095.1:7-1721
Oscillatoriales cyanobacterium USR001
Peptidase-C39
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OJJ25270.1
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Roseofilum reptotaenium AO1-A
Peptidase_M42
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DUF4114
Big_13
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CarboxypepD-reg-IG
HYIG
IG
Hw_PIG
Cadherin
IG
DUF6531
RHS_repeat
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RHS_repeat
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Peptidase-C39
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Escherichia coli
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rve_3
HTH_21
HTH_1
Peptidase-C39
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Acetylase
TAD93340.1
RDXC01000114.1:1-2664
Oscillatoriales cyanobacterium
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Peptidase_C70
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McrB-AAA
WP_000026703.1
NZ_AGTL01000002.1:76008-80345
Escherichia coli O145:H28 str. 4865/96
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GST_N_2
GST_C_2
Peptidase-C39
PAAR_motif
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WP_004143342.1
NZ_GG694030.1:5122-11782
Cardiobacterium hominis ATCC 15826
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Peptidase-C39
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Tox-LHH
WP_011710971.1
NZ_QIGG01000062.1:32220-35463
Allorhizocola rhizosphaerae
YbaB_DNA_bd
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RHS
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OLH
WP_112096716.1
NZ_QVPN01000004.1:1-2126
Salmonella enterica
DDE_3
Peptidase-C39
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Reovirus_L2_7th
WP_210694319.1
NZ_JABVJY010000359.1:1-1401
Pseudomonas protegens
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Peptidase_C70
WP_141233598.1
NZ_HG424585.1:422-1334
Lysobacter antibioticus HS124
Peptidase-C39
WP_206600854.1
NZ_OFFU01000042.1:15077-16081
Pseudomonas ovata
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WP_183876264.1
NZ_JACIAM010000019.1:15558-26661
Rhizobium sp. BK491
TPS
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Toast-rack
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FilM
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Fil_haemagg_2
Fil_haemagg_2
Peptidase-C39
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DUF423b
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DUF746
WP_184832423.1
NZ_JACHMN010000001.1:1001479-1011610
Allocatellilglobosispora scoriae
LuxR-HTH_Receiver
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WXG
Gln_amidase
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YbaB_DNA_bd
WP_185556563.1
NZ_JACKZD010000100.1:8140-19027
Nostoc sp. 2MC
MA
SIMIBI-Para
Helicase_C
Peptidase-C39
DNA_methylase
cpREC
HTH-2W-DrnB
WP_190295726.1
NZ_JACJPA010000030.1:1863-14680
Trichocoleus sp. FACHB-591
TPR
HTyD_D23
DALR_1
DUF4114
Peptidase_M8
PPC
Apolipoprotein
Peptidase-C39
G2P
WP_190652582.1
NZ_CP108258.1:124016-128934
Leptolyngbya sp. FACHB-1624
MPtase
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Peptidase_C39_2
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AP-ATPase
WP_191314413.1
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Amycolatopsis bullii
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NACHT
BTAD
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PG
Peptidase-C39
T7SS_ESX_Esp
YbaB_DNA_bd
WP_193852495.1
NZ_CAMPOB010000080.1:1165-3656
Xenorhabdus sp. BG5
Tnn_NPRO1
Peptidase-C39
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Transposase_20
DEDD_Tnp_IS110
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Coleofasciculus sp. LEGE 07081
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Peptidase_C70
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WP_193979372.1
NZ_JAOEWW010000011.1:52983-63451
[Phormidium] sp. LEGE 05292
DNA_methylase
Peptidase-C39
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cpREC
HTH-2W-DrnB
Helicase_C
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Planktothrix spongiae mayteni
YbaB_DNA_bd
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WP_206817842.1
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Chroococcus sp. FPU101
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Peptidase-C39
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WP_241456403.1
NZ_JAKZHO010000008.1:7856-21829
Undibacterium sp. TS12
ZnR
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RHS
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RHS_repeat
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Peptidase-C39
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Sigma70_r4
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Argonema galeatum A003/A1
23S_rRNA_IVP
KH_dom-like
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MMR_HSR1
Peptidase_C70
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Argonema galeatum A003/A1
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Peptidase-C39
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