

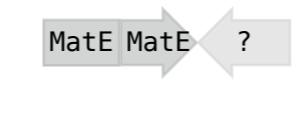
ART38309.1
KX576139.1:14993-21372
uncultured bacterium



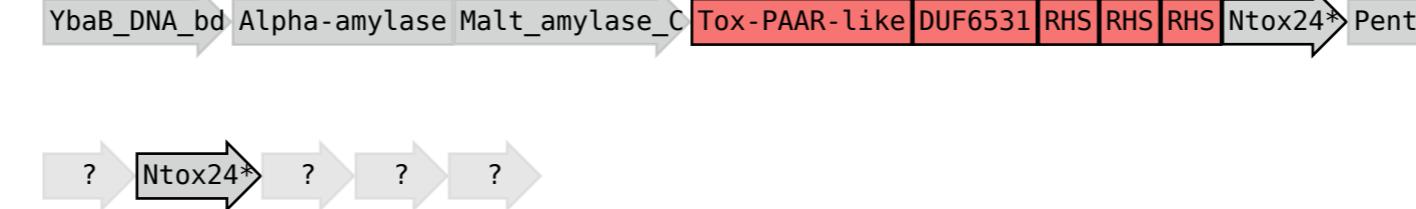
CBG89105.1
FN54502.1:2488973-2484627
Citrobacter rodentium ICC168



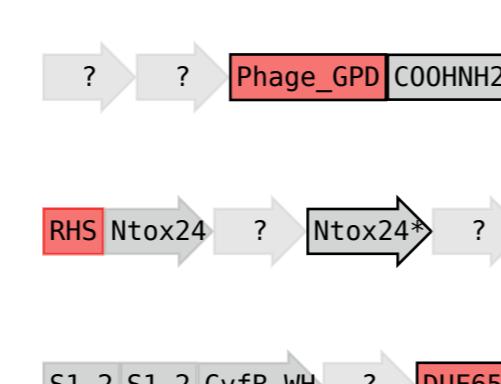
EBV0231387.1
AHAHDYI010000050.1:1-1247
Salmonella enterica subsp. *salamae* serovar *Sofia*



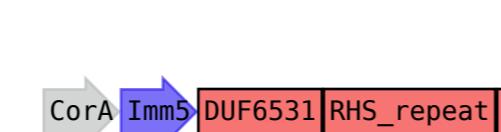
ECJ5866653.1
AATYYP01000002.1:238129-247314
Salmonella enterica subsp. *salamae*



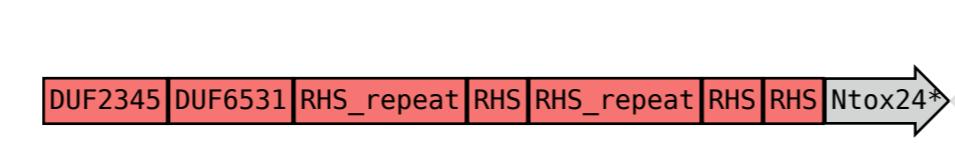
EF124099.1
GL349411.1:274086-282891
Neisseria sp. oral taxon 014 str. F0314



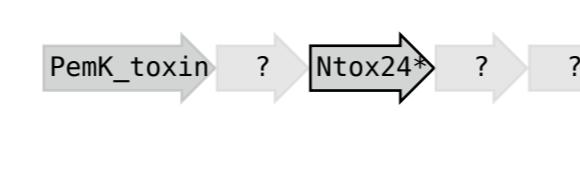
EXB27792.1
JEWN01000001.1:568794-571453
Acinetobacter baumannii 1437282



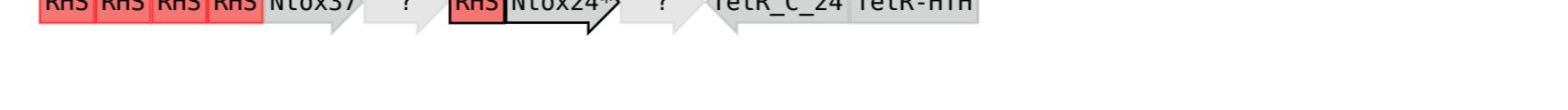
HBB702235.1
DADVN010000001.1:467770-475389
Escherichia coli



HCK4345624.1
DAIWR5010000006.1:1-3233
Pseudomonas aeruginosa



HCT39891.1
MPJB01000007.1:101247-110182
Moraxellaceae bacterium



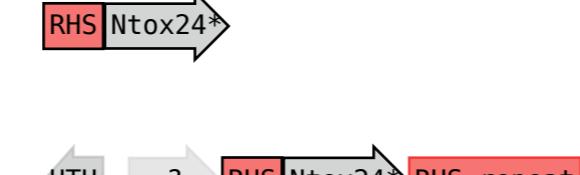
KAF1009578.1
WND001000001.1:194215-206011
Luteibacter sp.



KNE86404.1
AEWB02000003.1:2-6867
Aggregatibacter aphrophilus ATCC 33389



KPA17598.1
JPD01000570.1:13621-17102
Candidatus Magnetomorium sp. HK-1



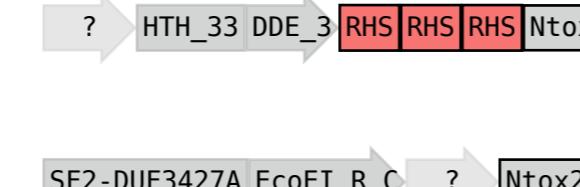
KRW62034.1
LKK01000001.1:213151-218934
Pseudomonas sp. TTU2014-080ASC



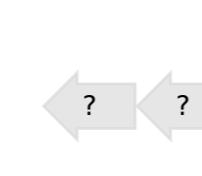
MBA2114127.1
AGAWGCY020000055.1:244881-247892
Salmonella enterica



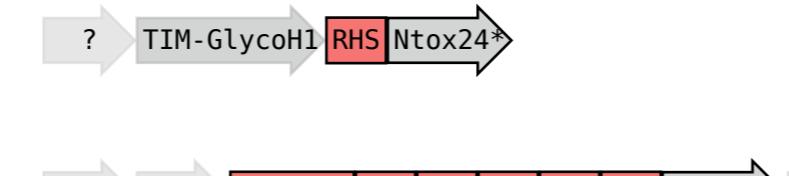
MBD8658560.1
JACYUQ010000016.1:1-1251
Oxalobacteraceae sp. CFBP 13730



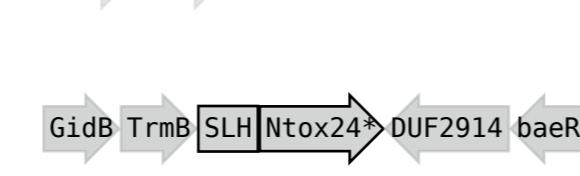
MBE0535677.1
JACZK0010000083.1:49481-52877
Phycisphaerae bacterium



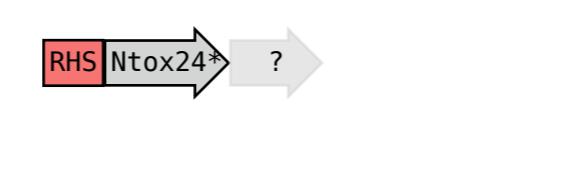
MBI3408315.1
JACPZP010000048.1:404-1479
Planctomyctetota bacterium



MBI4802861.1
JACQYT010000092.1:60128-63985
Elusimicrobia bacterium



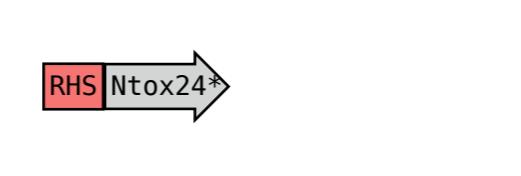
MBI9019390.1
JAETIOG010000170.1:3837-7772
Phycisphaerae bacterium



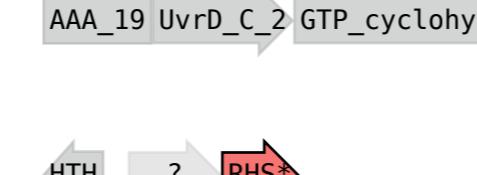
MBI9260594.1
JADJXL010000010.1:942-7084
Polyangiaceae bacterium



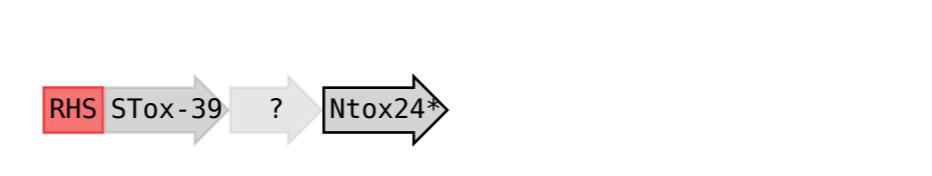
MBI9262824.1
JADJXL010000018.1:646214-648402
Polyangliaceae bacterium



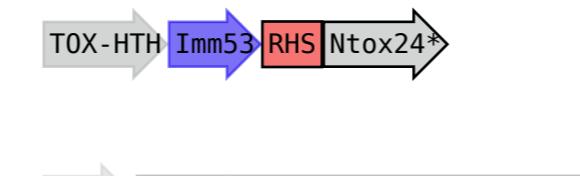
MBQ094965.1
JAFTSE010000043.1:7383-9049
Akkermansia sp.



MBR0968506.1
JAFCKV010000039.1:8479-16122
Bryodyrhizobium diazoeficiens



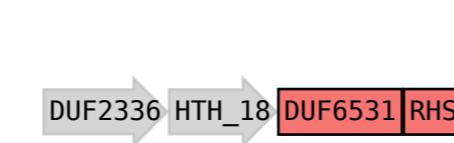
MBX3251153.1
JAHBW010000188.1:153-5862
Xylococcales bacterium



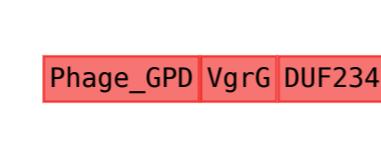
MBZ2010428.1
JAIOJB010000156.1:1-2034
Planctomyctetota bacterium



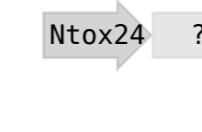
MCC2102186.1
JAGRKJ010000127.1:1-1922
Hyphomicrobiales bacterium



MCC7330937.1
JADYZ0010000157.1:1-1204
Gammaproteobacteria bacterium



NGO37790.1
JAACKYAO10000001.1:95-724
Lismisphaera ngatamarikiensis



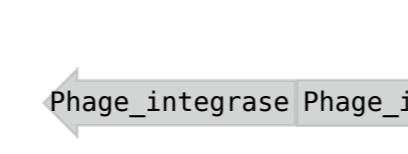
NOT88217.1
JABFRN010000010.1:1900644-1909610
Lysobacter sp.



NYZ69815.1
JACKCB010000154.1:825-3022
Spartanivicinus marinus



PIT08410.1
MDVA01000033.1:714-2578
Snodgrassella communis



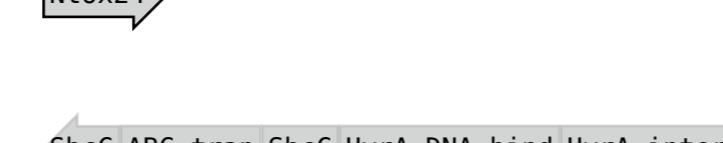
PNC69062.1
PKJR01000008.1:1795-3583
Akkermansi muciniphila



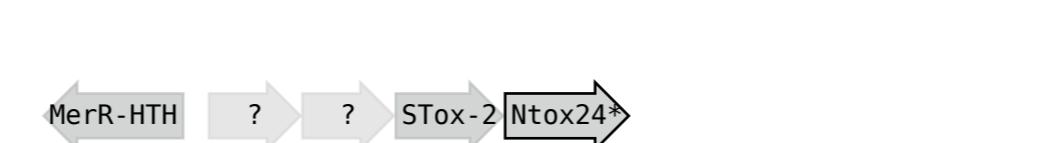
PWB18185.1
QET01000007.1:159099-176441
Comamonas sp. JNW



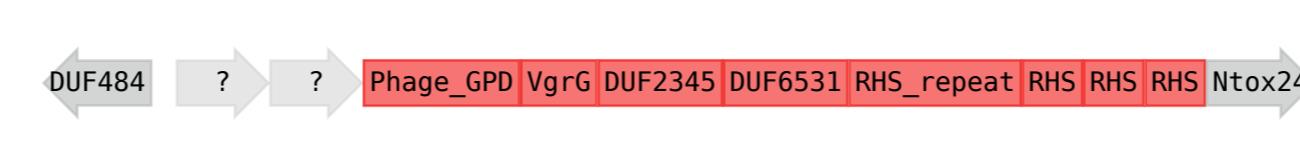
QPB31975.1
CP038008.1:1250793-1259792
Citrobacter rodentium



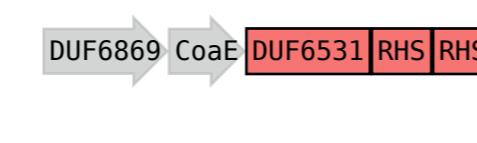
QGM9373.1
CP044331.1:2830072-2839534
Methylcytistis parvus



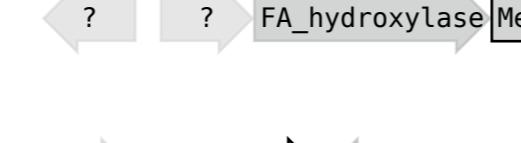
QIA75982.1
CP040863.1:131498-140654
Rodentibacter caecumuris



RPO23810.1
NSK01000012.1:204272-209690
Pseudomonas aeruginosa



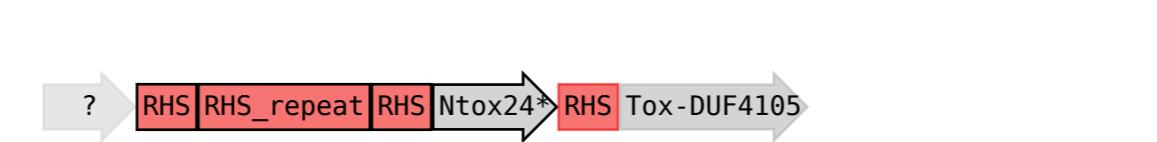
RYZ80261.1
SECL01000005.1:10341-13209
Moraxellaceae bacterium



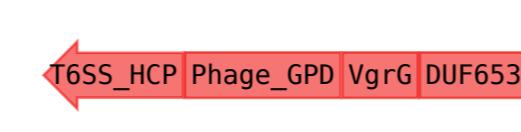
SDG76201.1
PMY001000025.1:2253-9405
Paraburkholderia lycopersici



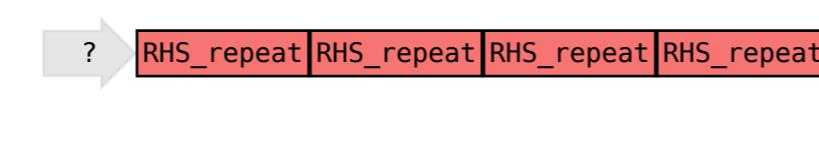
SEG15533.1
FNKV01000037.1:15318-18966
Nitrososphaera multififormis ATCC 25196



TAN40871.1
SCQE01000093.1:11134-13031
Nitrospirae bacterium



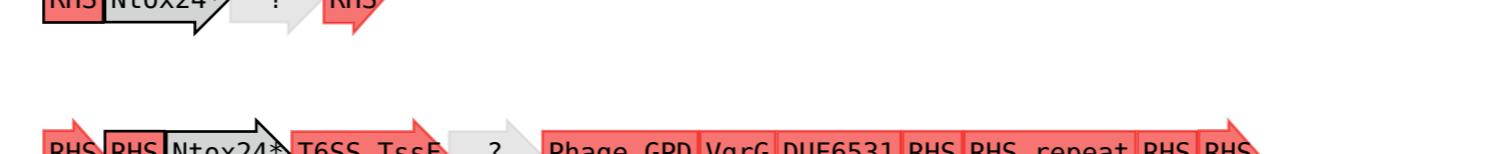
TWT45278.1
SMU01000001.1:2110025-2115989
Phycisphaerae bacterium RAS1



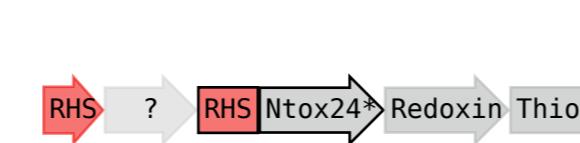
VTX59384.1
CAFBLM010000039.1:45-521
uncultured *Aggregatibacter* sp.



WP_212749466.1
NZ_KB849164.1:1220164-1223481
Acinetobacter bohemicus ANC 3994



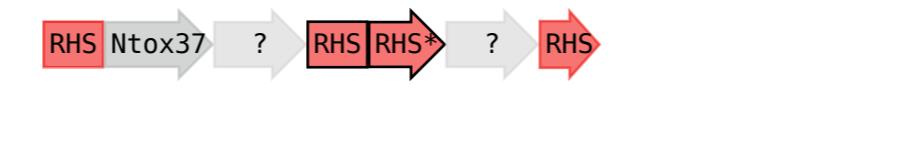
WP_050603340.1
NZ_CP012067.1:1207888-1216856
Aggregatibacter aphrophilus



WP_224229621.1
NZ_CPM04340.1:1185470-1195341
Xanthomonas oryzae pv. *oryzicola*



WP_049876215.1
NC_010162.1:2542886-2547658
Megalodessulfovibrio gigas DSM 1382 = ATCC 19364



WP_067550200.1
NC_010162.1:2484882-2491817
Sorangium celulosum So ce56

