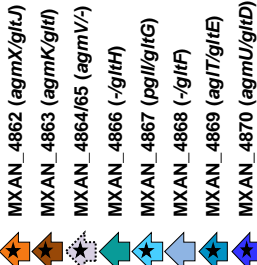


Myxococcus xanthus DK_1622 (Taxid: 246197/Deltaproteobacteria)

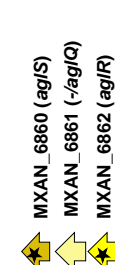
Cluster G1



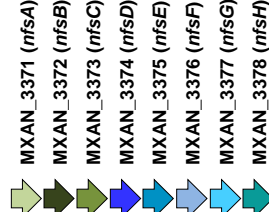
Cluster G2



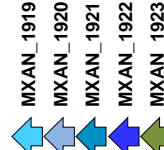
Cluster M1



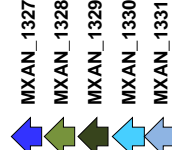
Cluster G3



Cluster G4



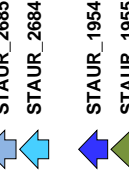
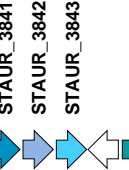
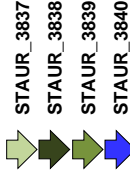
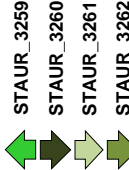
Cluster G5



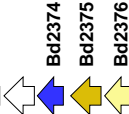
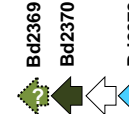
Cluster M2



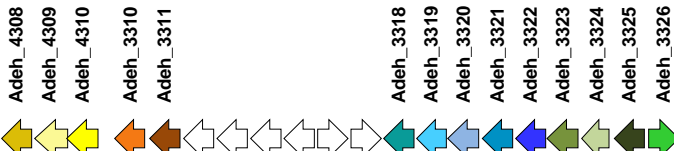
Stigmatella aurantiaca DW4/3-1 (Taxid: 378806/Deltaproteobacteria)



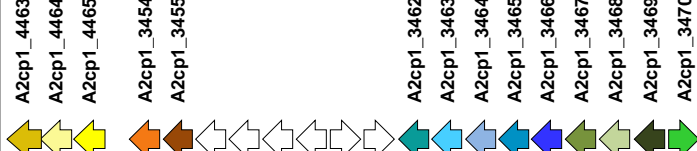
Bdellovibrio bacteriovorus HD100 (Taxid: 264462/Deltaproteobacteria)



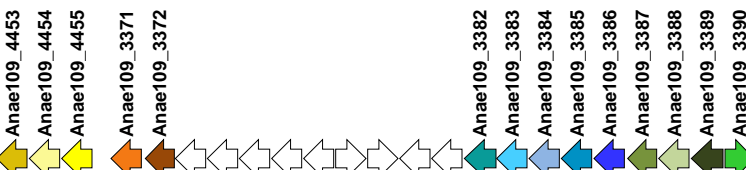
Anaeromyxobacter dehalogenans 2CP-C (Taxid: 290397/Deltaproteobacteria)



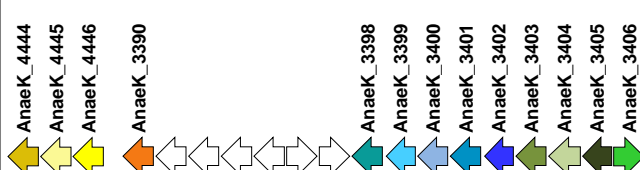
Anaeromyxobacter dehalogenans 2CP-1 (Taxid: 455488/Deltaproteobacteria)



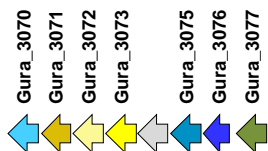
Anaeromyxobacter sp. Fw 109-5 (Taxid: 404589 /Deltaproteobacteria)



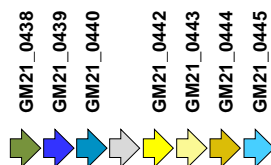
Anaeromyxobacter sp. K (Taxid: 447217/Deltaproteobacteria)



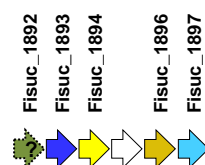
***Geobacter uraniireducens* Rf4**
(Taxid: 351605/Deltaproteobacteria)



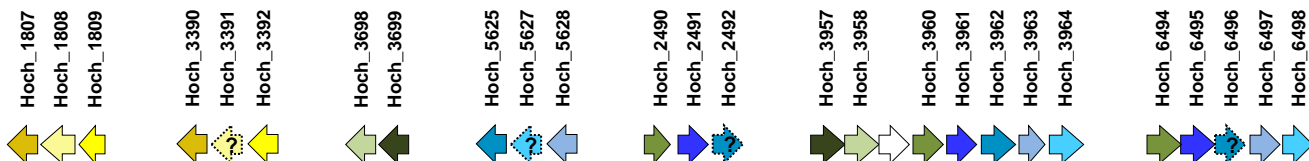
***Geobacter* sp. M21**
(Taxid: 443144/Deltaproteobacteria)



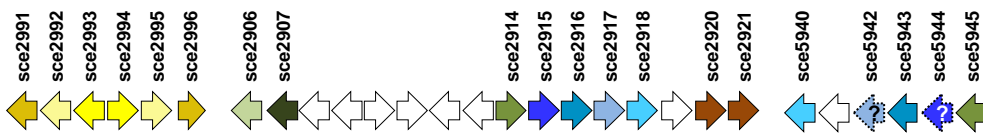
***Fibrobacter succinogenes* S85**
(Taxid:59374/Fibrobacteres)



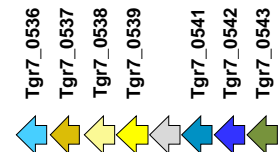
***Haliangium ochraceum* DSM 14365** (Taxid: 502025/Gammaproteobacteria)



***Sorangium cellulosum* Soce56** (Taxid: 448385/Deltaproteobacteria)



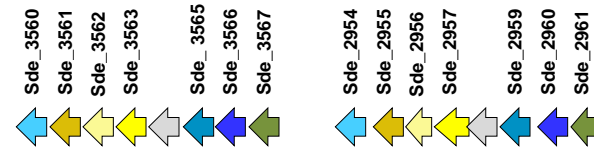
***Thioalkalivibrio* sp. HL-EbGR7**
(Taxid: 396588/Gammaproteobacteria)



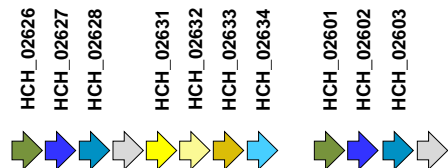
***Teredinibacter turnerae* T7901** (Taxid: 377629/Gammaproteobacteria)



***Sccharophagus degradans* 2-40**
(Taxid: 203122/Gammaproteobacteria)



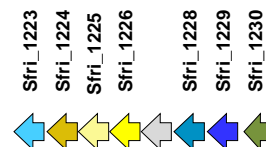
***Hahella chejuensis* KCTC 2396**
(Taxid: 349521/Gammaproteobacteria)



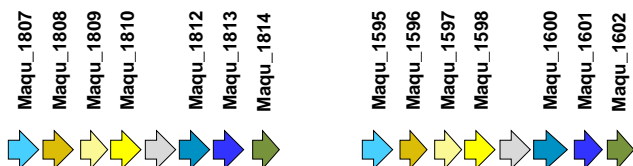
***Cellvibrio japonicus* Ueda107**
(Taxid:498211/Gammaproteobacteria)



***Shewanella frigidimarina* NCIMB 400**
(Taxid: 318167/Gammaproteobacteria)



***Marinobacter aquaeolei* VT8** (Taxid: 351348/Gammaproteobacteria)



***Methylibium petroleiphilum* PM1**
(Taxid: 420662/Betaproteobacteria)

