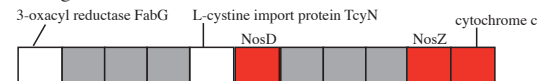


Flavobacteria

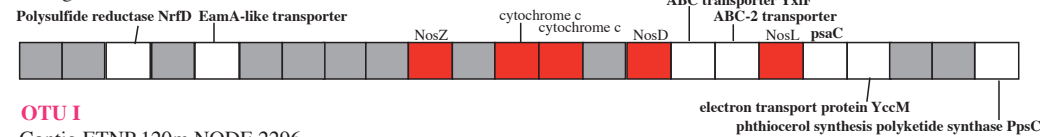
Contig ETNP 120m NODE 73975



Hypothetical

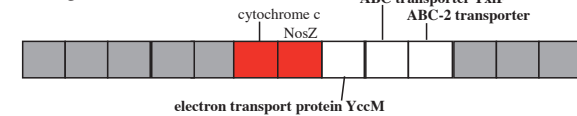
S adapted

Contig ETNP 120m NODE 137405

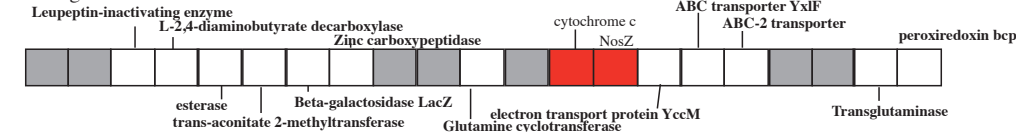


OTU I

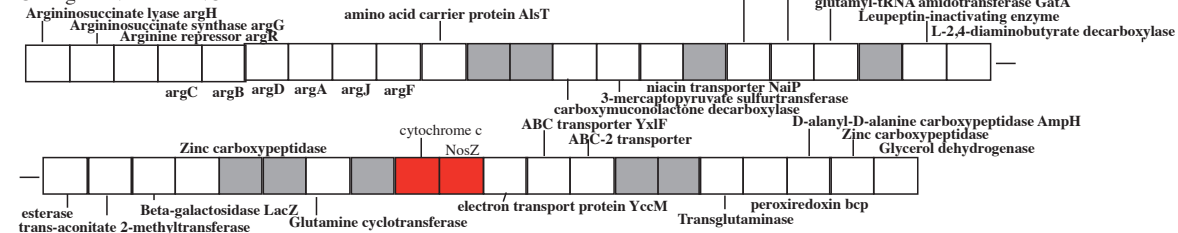
Contig ETNP 120m NODE 2296



Contig ETNP 140m NODE 23444

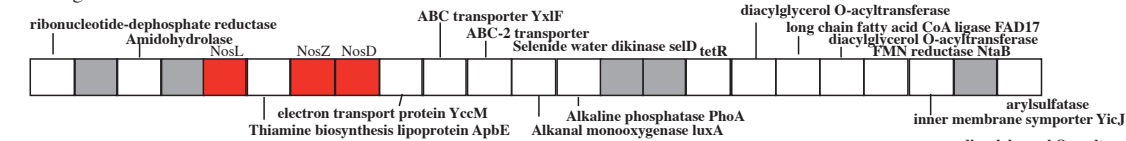


Contig ETNP 160m NODE 333



OTU II

Contig ETNP 160m NODE 833140



Contig ETNP 180m NODE 320139

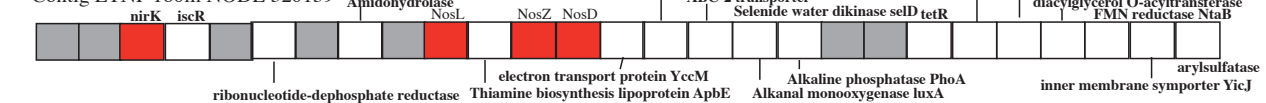


Figure S12. Schematic of selected contigs containing *nosZ* gene. Hypothetical proteins are shown in gray. Proteins related to N_2O reduction are shown in red. Contigs are clustered by their phylogenetic affiliation.