

CO₂-fixation by Type X methanotrophs

Takeshi Naganuma[1], Hiroyuki Kimura[1]

[1] School of Biosphere Sci., Hiroshima Univ.

<http://home.hiroshima-u.ac.jp/hubol/naganuma.html>

Methanotrophs (methane-oxidizing bacteria), as well as thiotrophs (sulfur-oxidizing & CO₂-fixing bacteria), are important primary producers and symbionts for the deep-sea chemosynthesis-dependent animals such as vestimentiferan and pogonophoran tubeworms. Vestimentiferans host mostly thiotrophs, while pogonophorans harbor thiotrophs and/or methanotrophs. The apparent double symbiosis of pogonophorans may be explained by the involvement of Type X methanotrophs capable of both methane-oxidation and CO₂-fixation. We have detected the symbiotic bacteria that bear the genes encoding Type X methanotrophic 16S rRNA, methane mono-oxygenase (CH₄-oxidizing enzyme) and RuBisCO (CO₂-fixing enzyme). The Type X species should be advantageous in carbon assimilation, because 1) part of CH₄ is assimilated as biomass and 2) the rest is oxidized to CO₂ and assimilated by RuBisCO. Net release of CO₂ via CH₄-oxidation should be thus minimized. The apparent double metabolism strategy exhibited by Type X methanotrophs is discussed in terms of ecology and economics relevant to CO₂ dynamics.