



HIGH CH₄ CONSUMPTION RATES BY *Sphingobacterium* sp.

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Keywords: Greenhouse gas, methane, *Sphingobacterium* sp.

Introduction. Methane (CH₄) is the second greenhouse gas after CO₂, with a global warming potential between 25-72. It is produced during enteric fermentation, natural gas extraction, oil refining, wastewater treatment and mines, mainly [1]. When the percentage of CH₄ is below the explosive limit (5% in air) biological treatments are a viable option to degrade it and methanotrophic bacteria have been studied [2,3]. However, CH₄ removal is restricted by its low solubility (28 mg L⁻¹), so the systems are normally limited by mass transfer. The objective was to characterize a new bacterium with high CH₄ consumption rates and to study CH₄ degradation in different bioreactors configurations.

Methods. The strain was isolated from a consortium of methanotrophic bacteria obtained of the wastewater treatment plant in UAM-Iztapalapa [2]. Bacterium was isolated by the streak plate method on mineral salts medium solidified with Gelrite Gellan Gum. Plates were incubated in CH₄ atmosphere at 30°C. Bacteriological agar and CH₄ degradation capacity was tested at different concentrations in closed flask systems. The consumption rates were determined using the Gompertz model.

The strain was previously identified by molecular biology techniques as *Sphingobacterium* sp. DDT-6 with a sequence identity of 97.5%. The isolate was tested in a stirred tank reactor, a two phase partitioning bioreactor and a membrane reactor under the same conditions reported by Zúñiga et al., 2011.

Table 1. CH₄ consumption, CO₂ production rates at different percentages of CH₄ in closed flasks

CH ₄ (%)	r _{CH₄} (g m ⁻³ h ⁻¹)	r _{CO₂} (g m ⁻³ h ⁻¹)	Q _{CH₄} (mg _{CH₄} g _x ⁻¹ h ⁻¹)
5	1.64 ± 0.02	0.51 ± 0.07	55 ± 6.1
10	8.06 ± 0.16	0.17 ± 0.01	203 ± 10.5
15	5.16 ± 0.68	0.19 ± 0.02	142 ± 11.1
20	3.99 ± 0.17	0.26 ± 0.01	79 ± 4.9
Control X	nd	0.17 ± 0.02	nd

Results.

Table 1 shows that the maximum specific degradation rate was 203 mg_{CH₄} (g_x h)⁻¹ at 10% CH₄ in closed flask experiments. Figure 1 shows the evolution of the specific CH₄ consumption rate in the stirred tank reactor. Although the highest and initial value was 350 mg_{CH₄}(g_xh)⁻¹, and in the steady state was around 150 mg_{CH₄}(g_xh)⁻¹, the first value is higher than the 300 mg_{CH₄}(g_xh)⁻¹ reached in membrane reactor. Finally in the two phase partitioning bioreactor 80 mg_{CH₄}(g_xh)⁻¹ was achieved

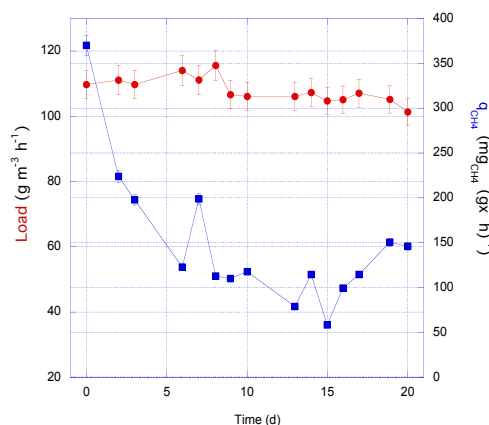


Fig.1 Specific CH₄ consumption rate of *Sphingobacterium* sp. in a stirred tank reactor.

Conclusions. *Sphingobacterium* sp. has CH₄ consumption rates twice higher than values reported for any methanotroph. This is the first report of CH₄ consumption by this bacterium.

Acknowledgements. The authors thanks to CONACyT and Sylvie Le Borgne for the support in the molecular biology identification.

References.

- Goodland R, Anhang J. (2009) Livestock and Climate Change. *World Watch*, 10-19.
- Rocha-Rios J, Bordel S, Hernández S, Revah S. (2009) Methane degradation in two-phase partition bioreactors, *Chem Eng J*. 289–292.
- Zúñiga C, Morales M, LeBorgne S, Revah S. (2011) Production of poly-β-hydroxybutyrate (PHB) by *Methylobacterium organophilum* isolated from a methanotrophic consortium in a two-phase partition