
Impact of CO₂ supply on the orientation of carbon metabolism in the diatom *Phaeodactylum tricornutum*

Bing Huang¹, Justine Marchand*¹, Brigitte Moreau¹, Grégory Carrier², Ewa Lukomska², Gaël Bougaran², Jean-Paul Cadoret³, Hu Hanhua⁴, and Benoît Schoefs¹

¹Laboratoire Mer, Molécules, Santé (MMS) – Université du Maine : EA2160 – IUML FR-3473 CNRS, Avenue Olivier Messiaen, 72000 LE MANS, France

²Laboratoire de Physiologie et Biotechnologie des Algues (PBA) – Institut Français de Recherche pour l'Exploitation de la Mer (IFREMER) – Rue de l'Île d'Yeu, BP 21105 Nantes cedex 3, France

³Greensea – Greensea – Promenade du sergent Navarro, 34140 Mèze, France

⁴Key Laboratory of Algal Biology – Institute of Hydrobiology, Chinese Academy of Sciences, Wuhan 430072, Chine

Résumé

Atmospheric carbon dioxide is expected to rise from current levels of 400 μatm to 700-1000 μatm by the end of this century, beyond the levels of the past 800,000 years of glacial-interglacial periods. Additional CO₂ into seawater could perturb the physiological processes of marine phytoplankton, including growth, photosynthesis and metabolic re-allocations. In particular, diatoms are biogeochemically important because they contribute up to 40% of the marine primary production and show relatively high carbon sequestration. The goal of this study was to evaluate the impact of the CO₂ supply on the marine diatom *Phaeodactylum tricornutum* using a continuous turbidostat bioreactor. Two different pCO₂ levels, 400 μatm (C-) and 1000 μatm (C+) were applied during a transition C+/C-/C+ of 60 days during which physiological (growth, photosynthesis, respiration, pigmentation) and molecular (RNAseq, qPCR) responses were followed. The results showed that growth rate and respiration were enhanced in C+. Parameters of the fluorescence showed that high CO₂ supply could improve the efficiency of the photochemical and biochemical phases of photosynthesis as illustrated by the decrease of the non-photochemical quenching (NPQ), suggesting a reduced demand in dissipation of excess of energy. Molecular data indicate that 534 genes are up-regulated while 377 down-regulated in the C- condition. Carbon acquisition is particularly reinforced in C- through the activation of the CO₂ concentration mechanism. Also several transcripts coding for proteins involved in cell cycle regulation were highlighted. The quantity of carbon accumulated in cells during the experiment seems to reflect a tendency to restore the initial equilibrium.

Mots-Clés: carbon dioxide, carbon metabolism, diatom, RNAseq, physiology

*Intervenant