



Prokaryotic respiration and production in the meso- and bathypelagic realm of the eastern and western North Atlantic basin

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ABSTRACT: We measured prokaryotic production and respiration in the major water masses of the North Atlantic down to a depth of ~4,000 m by following the progression of the two branches of North Atlantic Deep Water (NADW) in the oceanic conveyor belt. Prokaryotic abundance decreased exponentially with depth from 3 to 0.4×10^6 cells mL⁻¹ in the eastern basin and from 3.6 to 0.3×10^6 cells mL⁻¹ in the western basin. Prokaryotic production measured via ³H-leucine incorporation showed a similar pattern to that of prokaryotic abundance and decreased with depth from 9.2 to $1.1 \mu\text{mol C m}^{-3} \text{ d}^{-1}$ in the eastern and from 20.6 to $1.2 \mu\text{mol C m}^{-3} \text{ d}^{-1}$ in the western basin. Prokaryotic respiration, measured via oxygen consumption, ranged from about 300 to $60 \mu\text{mol C m}^{-3} \text{ d}^{-1}$ from ~100 m depth to the NADW. Prokaryotic growth efficiencies of ~2% in the deep waters (depth range 1,200-4,000 m) indicate that the prokaryotic carbon demand exceeds dissolved organic matter input and surface primary production by 2 orders of magnitude. Cell-specific prokaryotic production was rather constant throughout the water column, ranging from 15 to 32×10^{-3} fmol C cell⁻¹ d⁻¹ in the eastern and from 35 to 58×10^{-3} fmol C cell⁻¹ d⁻¹ in the western basin. Along with increasing cell-specific respiration towards the deep water masses and the relatively short turnover time of the prokaryotic community in the dark ocean (34-54 d), prokaryotic activity in the meso- and bathypelagic North Atlantic might be higher than previously assumed.

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