



Dynamic change in an ocean desert: Microbial diversity and trophic transfer along the 110 °E meridional in the Indian Ocean

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ABSTRACT

The eastern Indian Ocean is among the most oligotrophic regions in the world and has been described as an ocean desert. Limited information exists on microbial community profiles from marker gene data, and an open question in this system is how energy is transported from the base of the food web to higher trophic levels. Here we show that, along a 3300 km long transect in the ultra-oligotrophic eastern Indian Ocean, both alpha and beta diversity metrics for prokaryotic and eukaryotic trophic groups revealed remarkably strong latitudinal trends. The latitudinal Shannon diversity pattern for autotrophic eukaryotes furthermore aligned with the isotopic $\delta^{13}\text{C}$ ratios of particulate organic carbon, fractionated zooplankton and hand-picked fish larvae, suggesting a close trophic linkage between autotrophic eukaryotes and higher trophic levels. Our data also showed an increasing contribution of eukaryotic mixotrophs and a high contribution of heterotrophic eukaryotes towards warmer waters. These findings highlight that not only the recycling of organic matter via bacterial regeneration is important in this system but that mixo- and heterotrophic eukaryotes play a major role in redistributing energy within the marine food web of these oligotrophic waters. Our data provide a baseline to understand how environmental changes such as warming surface waters might impact the open-ocean food web in this oligotrophic basin.

1. Introduction

Sea surface temperature in the tropical Indian Ocean (IO) has increased by an average of about 1 °C from 1950 to 2015 (Roxy et al., 2020) and is predicted to warm further under varying future emission scenarios (Zhongming et al., 2021). Ocean warming has been shown to enhance stratification (Hood et al., 2006) and reduce nutrient fluxes across the pycnocline (Boyd et al., 2016), and is predicted to alter nitrogen (N) budgets on time scales of decades (Huesemann et al., 2002). Additional expected effects are declining magnitudes of primary

productivity and carbon sequestration to deeper waters, with cascading effects on genetic and evolutionary traits of organisms (Boyd et al., 2010). Our study seeks to develop a deeper understanding of how community and trophic structures respond to physicochemical variability in open-ocean oligotrophic conditions along 110 °E. The aims are to provide a baseline and mechanistic framework for the food web structures (including NO_3^- fluxes over the pycnocline, stable isotope plankton stoichiometry and microbial community observations) for future studies.

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