



Supplement of

Transfer of diazotroph-derived nitrogen to the planktonic food web across gradients of N_2 fixation activity and diversity in the western tropical South Pacific Ocean

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5 Figure 1: Relative increase of cells abundance associated to *Synechococcus*, *Prochlorococcus*, bacteria, diatoms, pico-eukaryotes and nano-eukaryotes (blue pattern) after 48 h for E1, E2 and E3. Error bars represent the standard deviation of triplicate counts and the propagated analytical errors. * Diatoms were not counted at T0. ** Diatoms and nano-eukaryotes were not analyzed for E3.



Figure 2: NanoSIMS images showing the ¹⁵N-enrichment after 48 h of incubation in the presence of ¹⁵N₂ for *Prochlorococcus* (a,b), pico-eukaryotes (c,d), heterotrophic bacteria (e,f), *Synechococcus* (g), and *Trichodesmium* (h). The ROIs are represented in white line.



Figure 3: Clustering of planktonic communities by flow cytometry on green fluorescence vs. forward scatter cytograms: heterotrophic bacteria (red), *Prochlorococcus* (blue), *Synechococcus* (green), and the pico-eukaryotes (pink)

Table 1: Number of ROIs analyzed for diazotrophs (*Trichodesmium* in E1 and E2, UCYN-B in E3), *Synechococcus*, *Prochlorococcus*, bacteria, diatoms, pico-eukaryotes and nano-eukaryotes, for E1, E2 and E3.

| Experiment | diazotrophs | Synechococcus | Prochlorococcus | bacteria | diatoms | pico-euk. | nano-euk. |
|------------|-------------|---------------|-----------------|----------|---------|-----------|-----------|
| E1 | 30 | 87 | 32 | 200 | 8 | 111 | 60 |
| E2 | 25 | 156 | 213 | 85 | 33 | 200 | 29 |
| E3 | 192 | 50 | 115 | 70 | 0 | 70 | 0 |