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RECEIVED 10 February 2025
ACCEPTED 18 March 2025
PUBLISHED 01 April 2025

CITATION
Cohen NR, Ellis KA, Lampe RH, McNair H,
Twining BS, Maldonado MT, Brzezinski MA,
Kuzminov FI, Thamatrakoln K, Till CP,
Bruland KW, Sunda WG, Bargu S and
Marchetti A (2025) Corrigendum: Diatom
transcriptional and physiological responses
to changes in iron bioavailability
across ocean provinces.
Front. Mar. Sci. 12:1574405.
doi: 10.3389/fmars.2025.1574405

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Corrigendum: Diatom transcriptional and physiological responses to changes in iron bioavailability across ocean provinces

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KEYWORDS

diatoms, *Thalassiosira*, *Pseudo-nitzschia*, iron, metatranscriptomics, California Upwelling Zone, Northeast Pacific Ocean

A Corrigendum on

[Diatom transcriptional and physiological responses to changes in iron bioavailability across ocean provinces](#)

By Cohen NR, Ellis KA, Lampe RH, McNair H, Twining BS, Maldonado MT, et al. (2017). *Front. Mar. Sci.* 4:360. doi: 10.3389/fmars.2017.00360

In the published article, there was an error in [Figure 3](#) as published. The initial (T0) phytoplankton community composition at site C2 was incorrectly displayed, with prasinophytes erroneously appearing as relatively abundant taxa. The corrected [Figure 3](#) appears below.

A text correction has been made to **Results**, *Community Composition across Sites*, paragraph number 1. This sentence previously stated:

“In contrast, CUZ site C2 initially yielded a phytoplankton community transcript pool dominated equally by diatoms (30%) and prasinophytes (28%), with diatoms remaining a

dominant taxa following incubation (26–28%) and prasinophyte transcripts substantially decreasing from 28 to 3–8% in both Fe and DFB incubations.”

The sentence has been removed.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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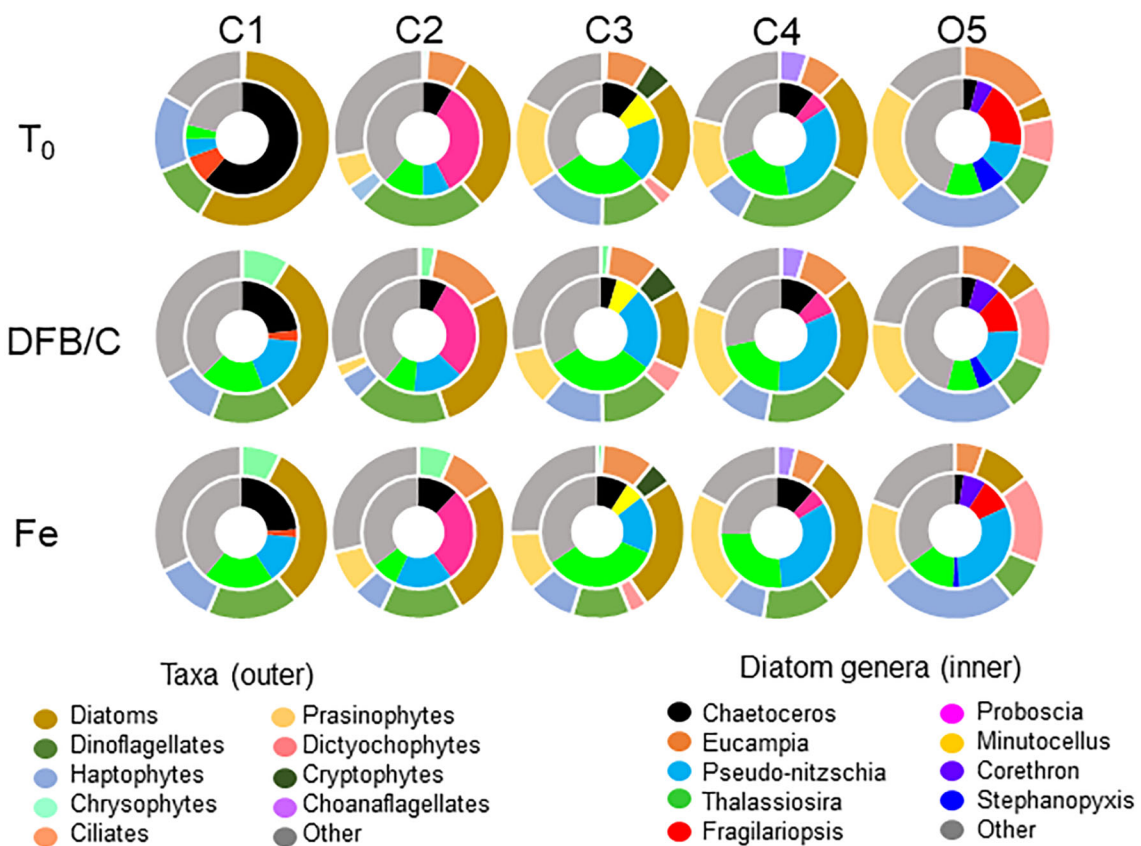


FIGURE 3

The average transcript proportions of phytoplankton taxa (outer charts) and diatom genera (inner charts) from initial seawater (T₀) and during the first time point (T₁; see Supplementary Table 1) within the Fe addition (Fe) and DFB addition (DFB) treatments at each site. Note that for site O5, the T₂ control (Ct) treatment is provided as the Fe-limited comparison.