

Trichodesmium growth rate and cell size related to pCO₂ and nutrients from experiments conducted at the University of Southern California from 2011-2013 (HiCO₂_AdaptCyano project)

Website: <https://www.bco-dmo.org/dataset/649904>

Data Type: experimental

Version: Final

Version Date: 2016-06-22

Project

» [Adaptation of key N₂-fixing cyanobacteria to changing CO₂](#) (HiCO₂_AdaptCyano)

| Contributors | Affiliation | Role |
|------------------------------------|-----------------------------------------------------|---------------------------------|
| Hutchins, David A. | University of Southern California (USC) | Principal Investigator, Contact |
| Webb, Eric A. | University of Southern California (USC) | Co-Principal Investigator |
| Allison, Dicky | Woods Hole Oceanographic Institution (WHOI BCO-DMO) | BCO-DMO Data Manager |
| York, Amber D. | Woods Hole Oceanographic Institution (WHOI BCO-DMO) | BCO-DMO Data Manager |

Table of Contents

- [Dataset Description](#)
 - [Methods & Sampling](#)
- [Data Files](#)
- [Parameters](#)
- [Instruments](#)
- [Deployments](#)
- [Project Information](#)
- [Funding](#)

Dataset Description

Trichodesmium erythraeum growth rates per day for all replicates in each treatment calculated from cell counts. *Trichodesmium erythraeum* was grown in biological triplicate under two CO₂ regimes (380 and 750 uatm pCO₂) and four different nutrient regimes (replete, phosphorus-limited, iron-limited, iron/phosphorus colimited).

These data were utilized in the following publications:

Walworth, N. G., Fu, F. X., Webb, E. A., Saito, M. A., Moran, D., McIlvin, M. R. & Hutchins, D. A. (2016a). Mechanisms of increased *Trichodesmium* fitness under iron and phosphorus co-limitation in the present and future ocean. *Nature Communications*, 7, 12081. doi: 10.1038/ncomms12081

Walworth, N. G., Lee, M. D., Suffridge, C., Qu, P., Fu, F., Saito, M. A., ... & Hutchins, D. A. (2018). Functional genomics and phylogenetic evidence suggest genus-wide cobalamin production by the globally distributed marine nitrogen fixer *Trichodesmium*. *Frontiers in Microbiology*, 9, 189. doi: [10.3389/fmicb.2018.00189](https://doi.org/10.3389/fmicb.2018.00189)

Genetic sequences from the *Trichodesmium erythraeum* used in this experiment can be accessed at The National Center for Biotechnology Information (NCBI) under the BioProject [PRJNA312342](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA312342).

Walworth, N. G., Lee, M. D., Fu, F. X., Hutchins, D. A., ... & Webb, E. A. (2016b). Molecular and physiological evidence of genetic assimilation to high CO₂ in the marine nitrogen fixer *Trichodesmium*. *Proceedings of the National Academy of Sciences*, 113(47), E7367-E7374. doi: [10.1073/pnas.1605202113](https://doi.org/10.1073/pnas.1605202113)

Related Dataset: [Trichodesmium proteomes and Hi CO₂ adaptation](#)

Methods & Sampling

The following are excerpts from the Nature Communication paper Walworth et al. 2016a. Please refer to this reference for more details of the methodology used to generate these data.

"We grew the nitrogen-fixing cyanobacterium *Trichodesmium* for 1 year under Fe/P co-limitation following 7 years of both low and high CO₂ selection. "

"Growth rates were calculated according to microscopic cell counts for reported values, or using *in vivo* chlorophyll fluorescence measurements with a Turner 10 AU fluorometer for semi-daily dilution calculations in real time during the experiments. Comparisons between cell counts and *in vivo*-based growth rates revealed no significant differences between the two methods of assessing biomass changes."

A picogram carbon/ μm filament was used for all replicates in each treatment (cell size proxy).

[[table of contents](#) | [back to top](#)]

Data Files

| File |
|--------------------------------------------------------------------------------------------------------------|
| cell_physio_size.csv (Comma Separated Values (.csv), 1.37 KB) MD5:8bce9923bfcc19905335a407352f7e91 |
| Primary data file for dataset ID 649904 |

[[table of contents](#) | [back to top](#)]

Parameters

| Parameter | Description | Units |
|----------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------|
| Treatments | CO ₂ and Nutrient regimes replicated three times; +P and -P speak of non-limiting or limiting phosphorus (20 μM and 0.5 μM phosphate respectively); +Fe or -Fe speak of limiting or non-limiting iron (500 nM iron); 380 or 750 indicate the partial pressure of CO ₂ | micromoles;nanomoles;uatm |
| Growth_rates | rates calculated from cell counts | reciprocal time - per day (d-1) - because the cell numbers cancel out |
| pg_C_per_um_filament | Picogram carbon per micron filament for all replicates in each treatment (cell size proxy) | picogram |

[[table of contents](#) | [back to top](#)]

Instruments

| | |
|-----------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Dataset-specific Instrument Name | Turner 10 AU fluorometer |
| Generic Instrument Name | Turner Designs Fluorometer 10-AU |
| Generic Instrument Description | The Turner Designs 10-AU Field Fluorometer is used to measure Chlorophyll fluorescence. The 10AU Fluorometer can be set up for continuous-flow monitoring or discrete sample analyses. A variety of compounds can be measured using application-specific optical filters available from the manufacturer. (read more from Turner Designs, turnerdesigns.com, Sunnyvale, CA, USA) |

[[table of contents](#) | [back to top](#)]

Deployments

lab_Webb_Hutchins_Fu

| | |
|--------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Website | https://www.bco-dmo.org/deployment/59058 |
| Platform | Webb-Hutchins-Fu USC |
| Start Date | 2011-08-15 |
| End Date | 2013-03-31 |
| Description | Lab experiments of transcriptome samples (labeled 750) obtained from cultures grown in either projected year 2100 CO2 levels (~750ppm) or current 380ppm levels (labeled 380) for four years. |

[[table of contents](#) | [back to top](#)]

Project Information

Adaptation of key N2-fixing cyanobacteria to changing CO2 (HiCO2_AdaptCyano)

Coverage: Culture study at the University of Southern California, Los Angeles

Description from NSF award abstract:

This study will employ a novel combination of experimental evolution techniques and state-of-the-art molecular methods to yield unique insights into adaptive changes in the keystone marine cyanobacteria *Trichodesmium* and *Crocospaera* in response to selection by high CO₂. Several studies have suggested that N₂-fixation rates of the biogeochemically-critical cyanobacteria *Trichodesmium* and *Crocospaera* may increase dramatically in the future high CO₂ ocean, but these have all used the same limited set of cultured isolates and considered cells only briefly acclimated to elevated CO₂. The investigator's new results, however, demonstrate that a broad diversity of high- and low-CO₂ adapted ecotypes exists within each diazotroph genus. Furthermore, in a preliminary four year experimental evolution study with *Trichodesmium*, the PIs observed large adaptive responses following 500-700 generations of selection by high CO₂ but in a completely unexpected way. All of the six replicate high CO₂-adapted cell lines exhibited strong constitutive up-regulation of N₂ fixation rates. These very elevated N₂ fixation rates continued, even though the cultures have were switched back to low-CO₂ conditions for many months. Expression of the *nif* operon and N assimilatory genes was also up-regulated in these cell lines, as is expression of many intergenic regions of the genome.

The investigators hypothesize that constitutive up-regulation of cellular N₂ fixation systems may be a common adaptive response of both *Trichodesmium* and *Crocospaera* under extended selection by elevated CO₂. This project will test this hypothesis in a four-year experimental evolution study to determine the adaptive responses of both high- and low-CO₂ specialized ecotypes of these two diazotrophs to increased CO₂.

The investigators will grow representative high- and low-CO₂ adapted ecotypes from each genus in well-replicated cell lines at 380 ppm and 750 ppm CO₂ for up to 1000 generations. Periodically, they will perform "switch" experiments to measure N₂ and CO₂ fixation rates and growth rates of high CO₂-selected cell lines grown briefly (one week) at low CO₂, and vice versa. These switch experiments will allow screening for cell lines which exhibit adaptive changes in phenotypically-expressed rate parameters, such as those observed in the preliminary *Trichodesmium* study. Evolutionary mechanisms in the CO₂-selected cell lines will be examined by comparison of changes in their genomes, transcriptomes, and proteomes over time relative to reference genomes, using frozen samples archived monthly during the preceding selection period. Examination of these molecular and biochemical changes will be coordinated with an in-depth array of physiological and biogeochemical analyses. This combined approach will allow an evaluation of potential adaptive mechanisms in diazotrophic cyanobacteria ranging from indel, duplication, single nucleotide polymorphism, and transposition mutations to altered putative non-coding RNA expression, protein expression, and post-translational protein modifications, and then allow the investigators to link these mechanisms directly with their potential impacts on ecosystem-level biogeochemical processes like N₂ and CO₂ fixation. Finally, the research team will determine how long term selection by high CO₂ affects the iron and phosphorus requirements of *Trichodesmium* and *Crocospaera*, since constitutive up-regulation of N₂ fixation would also have major implications for limitation of diazotrophs by these two critical nutrients in the future high CO₂ ocean.

[[table of contents](#) | [back to top](#)]

Funding

| Funding Source | Award |
|----------------------------------------------------------|-----------------------------|
| NSF Division of Ocean Sciences (NSF OCE) | OCE-1260233 |

[[table of contents](#) | [back to top](#)]