



Supplementary Materials for

Niche Engineering Demonstrates a Latent Capacity for Fungal-Algal Mutualism

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This PDF file includes:

Materials and Methods
SupplementaryText
Figs. S1 to S7
Table S1
Captions for Movies S1 to S16
Captions for Datasets S1 to S7

Other Supplementary Materials for this manuscript includes the following:

Movies S1 to S16
Datasets S1 to S7 in a single zipped archive:

- Hom_DatasetS1.xlsx
- Hom_DatasetS2.xlsx
- Hom_DatasetS3.xlsx
- Hom_DatasetS4.xlsx
- Hom_DatasetS5.xlsx
- Hom_DatasetS6.xlsx
- Hom_DatasetS7.xlsx

Materials and Methods

Culture Medium and Pre-culturing

Cells were cultured in a defined basic medium designed to support the growth of various algal and fungal strains. A **Chlamydomonas-Yeast Basal (CYB)** medium (25 mOsm) lacking carbon or nitrogen sources, consists of the following components (per liter): 0.125 g MgSO₄•7H₂O, 0.074 g CaCl₂•H₂O, 0.4 g K₂HPO₄, 0.18 g NaH₂PO₄, 29.2 mg EDTA, 6.6 mg KOH, 1.2 mg MnCl₂•4H₂O, 0.3 mg ZnCl₂, 0.06 mg H₃BO₃, 0.2 mg CoCl₂•6H₂O, 0.4 mg CuCl₂•2H₂O, 0.05 mg Na₂MoO₄•2H₂O, 0.018 mg KBr, 0.003 mg KI, 0.0018 mg Na₃VO₄, 0.0018 mg Na₂SeO₃, and the following vitamins: 4 mg myo-inositol, 0.4 mg 4-amino benzoic acid, 0.4 mg calcium D-pantothenate, 0.4 mg niacin, 0.4 mg pyridoxine hydrochloride, 0.4 mg thiamine hydrochloride, 0.002 mg biotin, 0.002 mg cyanocobalamin, and 0.002 mg folic acid. The pH of this medium is 6.8. The designation “CYM:2% glucose + 10 mM KNO₂” is used to indicate a **Chlamydomonas-Yeast Medium (CYM)** with a CYB composition supplemented with 2% glucose and 10 mM potassium nitrite (final osmolarity of ~150 mOsm). All CYB, carbon, and nitrogen stock solutions were separately autoclaved (20 min at 120 °C), 0.2 µm-filter sterilized, and aged/equilibrated at room temperature for at least 24 hrs before use.

Prior to co-culturing experiments, fungal strains are cultivated in log-phase for at least 12 hrs on CYM:2% glucose + 37.4 mM NH₄Cl. *C. reinhardtii* strains are cultivated for at least 24 hrs on either CYM:20 mM Tris•HCl (pH 7.0) + 10 mM KNO₃ or a modified Tris-acetate medium, TAP* (equivalent to CYM:20 mM Tris•Acetate (pH 7.0) + “N-stock” (see below), but *without* CYB vitamins). Other algal strains were cultivated in CYM:N-stock. “N-stock” is a mixture of different nitrogen sources with a 1X composition (per liter) as follows: 0.6 g urea, 0.32 g KNO₃, 0.12 g acetamide, and 0.04 g NH₄Cl. Prior to co-culturing yeast and algae, cultures of individual organisms were centrifuged at 2000 x g (algae) or 4000 x g (yeast) for 3 min to pellet cells, and cells were gently resuspended in a volume of CYB equivalent to the pre-culture volume. The cells were washed two more times, with the final volume being half of the initial volume. Algal cells were allowed to recover for 30 min in CYB under light. Cell densities of these washed cultures were then measured using a Coulter counter (Z2 analyzer; Beckman Coulter, Inc., Danvers, MA) and a stock concentration of each cell type was prepared in fresh CYB medium for co-culture inoculation.

Co-culturing Experiments

Fluorescently-labelled prototrophic *S. cerevisiae* (S288c expressing monomeric teal fluorescent protein (26) under the ENO2 promoter at the HO locus) and intrinsically fluorescent *C. reinhardtii* (strain CC-1690) cells were co-cultured in a minimal salts CYB medium supplemented with glucose and/or potassium nitrite (see Culture Medium) in 1 mL volumes in 2 ml screw-cap tubes with silicone rubber gaskets (Neptune Scientific, San Diego, CA) [Figs. 1B, 3A, and S1] or in 250 µL volumes in the wells of 96-well round-bottom, polystyrene microtiter plates (well capacity ~330 µL) (Corning, product #3788; Corning Inc., Corning NY) [Figs. 2, 3B, S2, and S3]. Unless otherwise indicated, cells were inoculated at a density of 0.05x10⁶ cells/mL (dashed horizontal line in Figs. 1 and S1) and grown at room temperature. To maintain an air-tight environment, 2 mL

tubes were tightly capped and 96-well microtiter plates were sealed with a 76.2 μm thick inert polyolefin film with pressure-sensitive-release silicone adhesive (ThermalSealRTS, Excel Scientific, Victorville, CA), double-sealed with a 127 μm polyvinylidene fluoride film (CS Hyde Company, Lake Villa, IL), covered with a 1/8" thick custom glass plate, and vacuum sealed in a 76.2 μm thick nylon/polyethylene pouch (Doug Care Equipment, Springville, CA). Standing co-cultures were exposed to 45 or 110 $\mu\text{mol}/\text{m}^2/\text{s}$ (400-700 nm, photosynthetic active radiation) of continuous light from a bank of six F40T12 Soft White/ALTO tubes (Philips, Arlington, MA), appropriately attenuated using neutral density filters (Roscolux #398 Neutral Grey, 40% transmission; Rosco Laboratories, Central Falls, RI). A rotary tube mixer (model 346; Fisher Scientific, Waltham, MA) operated at 25 rpm was used for agitated co-culture experiments, (Fig. 1B, condition 6).

At appropriate experimental time points, co-cultured cells were thoroughly mixed by vigorous vortexing and inversion and counted by flow cytometry (LSR Fortessa; BD Biosciences, San Jose, CA); 10 μL at 0.5 $\mu\text{L}/\text{sec}$) using custom gates to discriminate yeast and algal cells based largely on CFP (ex=440 nm; em=470/20 nm) and APC-Cy7 (ex=635 nm; em=780/60 nm) fluorescence, respectively. A co-culture was sacrificed for each experimental measurement made. Cell counts were corrected for aggregate particle counts (typically less than 10%) by custom gating on particle side-scatter time-of-flight values (vs. side-scatter amplitude) and through multipeak-Gaussian fits of fluorescence amplitude histograms (27) assuming ≤ 4 aggregate populations using IgorPro (Wavemetrics, Lake Oswego, OR). Images of co-cultures were generated by adding 250 μL of a resuspended culture in a well of a 96-well round-bottom microtiter plate, centrifuging at 4000 x g for 3 min, and imaging from below using an Epson Perfection V700 Photo transparency scanner (Epson, Long Beach, CA). Other fungi and Chlamydomonas strains (see Table S1) were co-cultured and analyzed by similar methods. Maximum predicted cell densities for *C. reinhardtii* and *S. cerevisiae* over the gradient landscape of Fig. 2 were calculated assuming 0.54×10^{-12} moles of nitrogen per *C. reinhardtii* cell grown under autotrophic conditions (28, 29) and 2.5×10^{-12} moles of carbon per *S. cerevisiae* cell (30).

Co-cultures with filamentous fungi were started from small ($\leq 1 \text{ mm}^3$) hyphal fragments (dissected with sterile forceps), washed in CYB, and grown with $\sim 1 \times 10^5$ cells (total) of *C. reinhardtii* in 50 mL of CYB supplemented with 2% glucose and 10 mM KNO_2 medium in 150 mL polystyrene bottles (Corning, product #431153) under 45 $\mu\text{mol}/\text{m}^2/\text{s}$. For control experiments in which the dependence of fungi on *C. reinhardtii* for nitrogen is short-circuited, 20 mM NH_4Cl was added to the culture medium. For control experiments in which the dependence of *C. reinhardtii* on fungi for carbon is short-circuited, polystyrene bottles were only loosely capped, ensuring access to ambient CO_2 . For complementary control experiments in which the dependence of fungus on *C. reinhardtii* for nitrogen is short-circuited, wild-type, nitrite-positive utilizing fungal strains were used and an additional 10 mM KNO_2 was added to the culture medium. All culture inoculation and assembly steps were conducted aseptically in a tissue culture hood. Images and movies of symbiotic composite associations were acquired using a Zeiss Axio Zoom.V16, and processed using Zen 2012 (Zeiss, blue edition; Zeiss, Thornwood, NY) software.

Transmission Electron Microscopy

Fragments of fungal-algal associations of 3 mm^3 were sterilely dissected and fixed at room temperature for 6-12 hrs in 2.5% glutaraldehyde + 2% paraformaldehyde (made fresh) in 1X CYB medium supplemented with 10 mM KNO_3 , 62.5 mM sodium cacodylate, and 16 mM sodium phosphate at pH ~7.2. Fixed fragments were transferred to 1X CYB supplemented with 10 mM KNO_3 and 63 mM HEPES, and stored at 4°C (< 72 hrs) until further processed. Fragments were subsequently osmium fixed, resin embedded, sectioned, and imaged by standard methods (31) using a Hitachi H-7650 transmission electron microscope (Central Electron Microscope Facility, University of Connecticut Health Center, Farmington, CT). Cell wall thicknesses reported in Fig. 3 (mean \pm SD) were determined by 25 independent measurements of randomly chosen cell wall cross-sections using the image processing suite Fiji (32).

Phylogenetic Trees

Evolutionary trees of Fig. 3 were adapted and modified from published analyses of ascomycetous fungi (33, 34) and algae (35). Dating of fungal branch points were obtained from (36-38) and for algal branch points from (39-41). References for annotated species characteristics (e.g., Crabtree-positive = ability to ferment in the presence of oxygen) can be found in Table S1.

Fig. S1

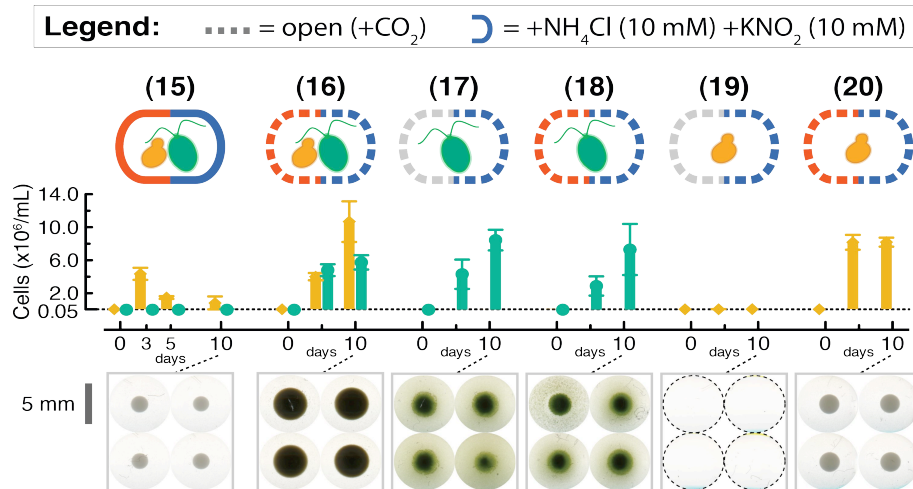
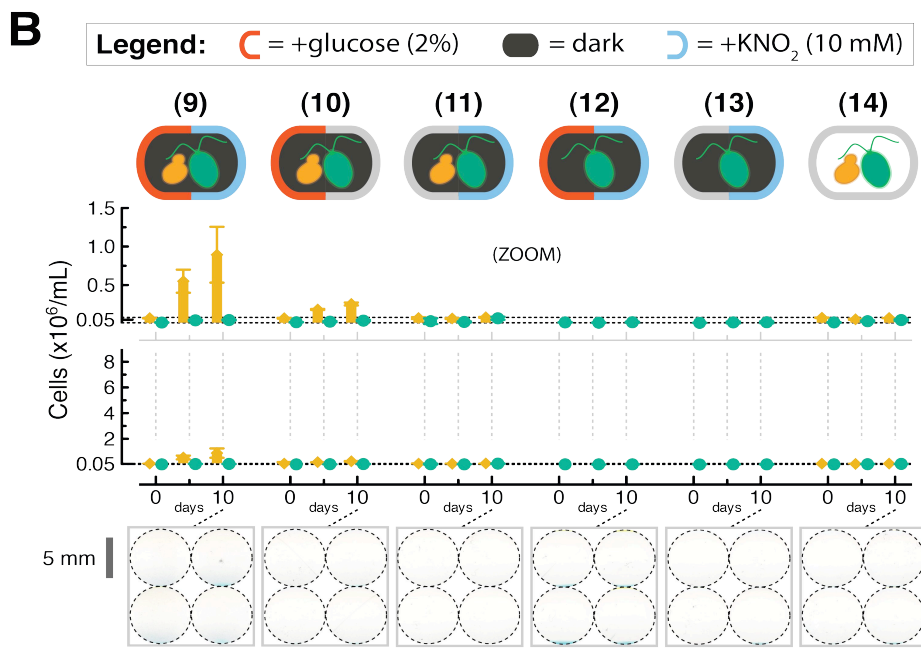
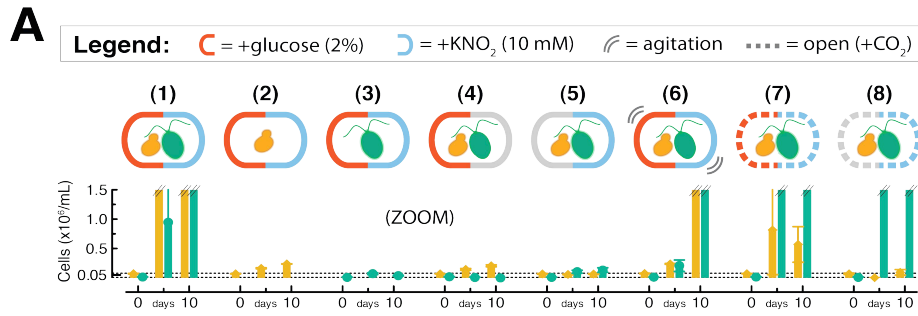


Fig. S1.

(A) Expanded view of Fig. 1B showing cell densities below 1.5×10^6 /mL. (B) Proliferation of *S. cerevisiae* and *C. reinhardtii* under additional co-culture conditions. See Fig. 1B legend for details. Conditions 9-13 were co-cultured in the absence of light. Although *C. reinhardtii* does not assimilate CO₂ or proliferate (substantially) on nitrite without light (14, 42), the alga generates sufficient light-independent nitrite reductase activity and ammonia production exists to support residual growth of *S. cerevisiae* in the dark. Conditions 15-20 include 10 mM ammonium chloride as a nitrogen source, which would allow *S. cerevisiae* to grow independently of *C. reinhardtii*. Condition 15 leads to loss of the alga; budding yeast rapidly out-proliferates the obligately dependent *C. reinhardtii* (although cell debris is accumulated; cf. image of pelleted cells for condition 15). Note different y-axis scales for different parts of the figure.

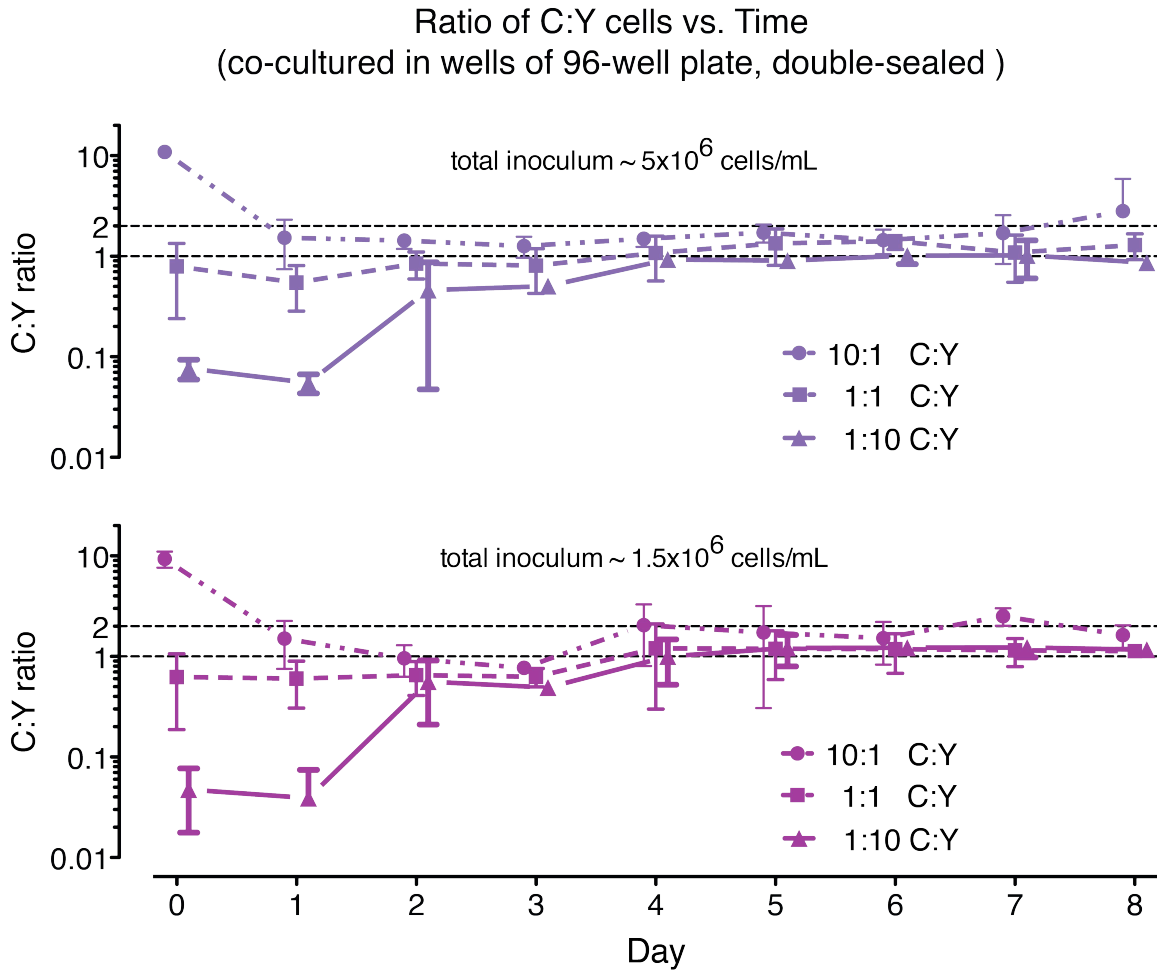


Fig. S2.

The ratio of *C. reinhardtii* to *S. cerevisiae* cells grown in sealed vessels (in CYM:2% glucose + 10 mM KNO₂ with 45 μmol/m²/s light) converges towards a stable ratio range over time (95% confidence intervals shown), with dashed lines for ratios of 1 and 2 drawn as a visual guide. Co-cultivation was done in wells of a vacuum-sealed, 96-well microtiter plate with initial inoculation ratios (nominally) of 10:1, 1:1, and 1:10 *C. reinhardtii* to *S. cerevisiae* (and at two different total cell inoculums: ~5x10⁶ (*top*), and ~1.5x10⁶ (*bottom*) cells/mL in 250 μL; N=4). The cell ratio converges to 1-2 C:Y within five days, essentially independent of initial inoculation ratio.

Fig. S3

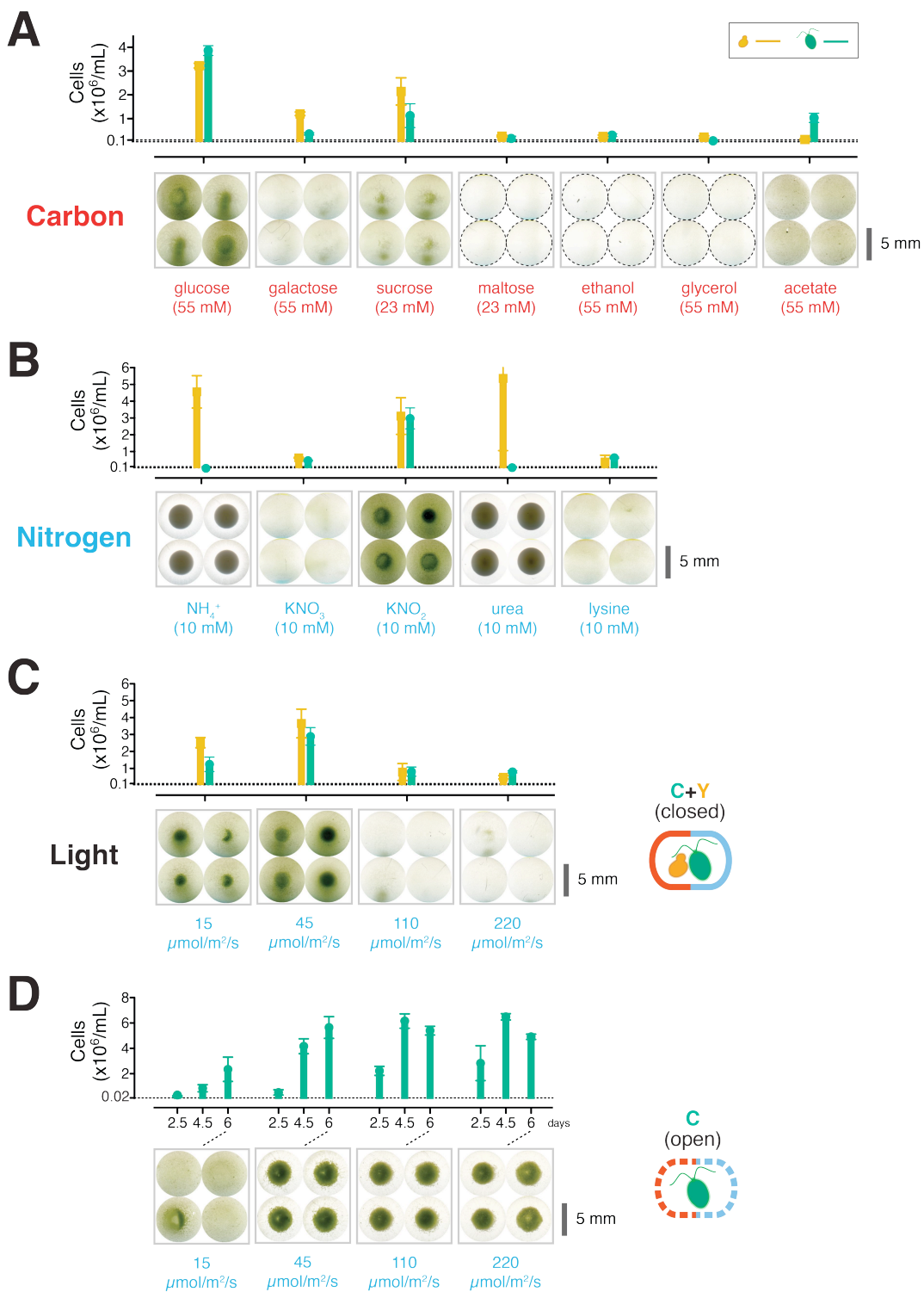


Fig. S3.

Mutualistic growth after 6 days of co-culture using different carbon and nitrogen resource nutrients, and light intensities. Unless otherwise indicated, the carbon source was 2% glucose (111 mM), the nitrogen source was 10 mM KNO₂, the irradiance was 45 μmol/m²/s, and *S. cerevisiae* and *C. reinhardtii* were inoculated at ~0.1x10⁶ cells/mL. **(A)** Carbon sources were used at the same concentration of 55 mM for comparison, with the exception of sucrose (disaccharide of glucose and fructose) and maltose (disaccharide of glucose), which were used at 23 mM. *C. reinhardtii* cannot use any of these carbon sources for growth, with the exception of acetate. Glucose is the preferred carbon source for *S. cerevisiae*, galactose is assimilated by respiro-fermentation, and ethanol, glycerol, and acetate are respired (30). Sucrose fermentation requires the induced expression of invertase, an extracellular enzyme that hydrolyzes sucrose. Although the *S. cerevisiae* strain used here has been corrected to enable maltose utilization (see Supplemental Table S1), the parental background may not be optimized to ferment it. **(B)** Nitrogen sources were used at the same concentration of 10 mM for comparison. *S. cerevisiae* can use NH₄⁺ and urea as nitrogen sources, and by day 6, leads to the extinction of *C. reinhardtii*. Lysine is an extremely poor nitrogen source for *S. cerevisiae* (43). In the absence of preferred nitrogen sources, *C. reinhardtii* expresses an extracellular L-amino-oxidase that hydrolyzes L-lysine to α-keto-ε-aminocaproic acid, H₂O₂, and NH₃ (44-46); NH₃ becomes a public good that is subsequently imported as a nitrogen source or diffuses away. Nitrate requires an additional two-electron NAD(P)H-mediated reduction to nitrite (by nitrate reductase, which is highly regulated) before it can be further assimilated by *C. reinhardtii* (47, 48), and metabolized nitrate leads to a complex light- and CO₂-dependent release of both nitrite and ammonium into the medium (49). Unlike nitrite or ammonium, nitrate can be stored and sequestered within the vacuoles of *C. reinhardtii*. **(C)** Light intensity can modulate the productivity of mutualism, with an optimum at ~45 μmol/m²/s, although all light levels examined lead to successful obligate mutualisms. **(D)** Proliferation of *C. reinhardtii* cells (alone) inoculated at ~0.02x10⁶ cells/mL and grown under the same conditions as in (C) except with access to atmospheric CO₂. The reduced mutualistic productivity at higher light levels in (C) are not due to a decreased ability of *C. reinhardtii* cells to proliferate; in fact, higher light levels lead to faster proliferation of *C. reinhardtii* cells (see days 2.5-4.5).

Fig. S4

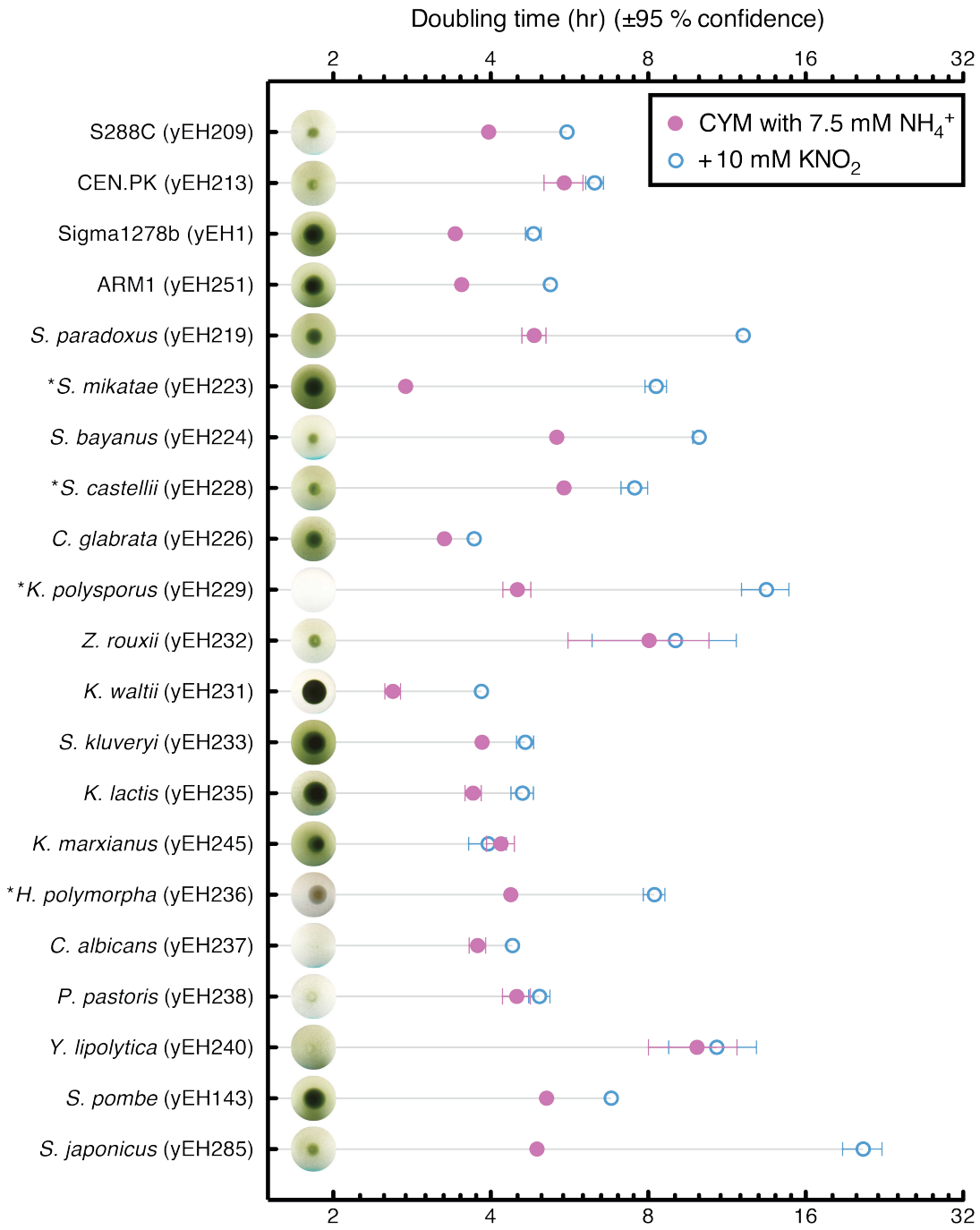


Fig. S4.

Doubling times (with 95% confidence intervals) of different yeasts across the fungal tree of Fig. 3 with (blue) and without (purple) 10 mM nitrite present in the growth medium. Yeasts (see Table S1 for full descriptions) were grown in CYM:2% glucose + 7.5 mM NH₄Cl with or without 10 mM KNO₂, and three replicate growth curves for each condition were generated using a Bioscreen C system (15 min interval measurements of 600 nm absorbance at 25 °C with shaking prior to measurement (Growth Curves USA, Piscataway, NJ) and an initial inoculum of 3x10⁴ cells/300 μL. Doubling times were determined by fitting phases of logarithmic growth by least-squares regression. Nitrite, an oxidant and common food preservative, generally inhibits growth, although to varying degrees depending on the yeast species and genetic background. Images of representative 9-day old cultures of yeast with *C. reinhardtii* (see Fig. 3A) have been superimposed for comparison with the success of mutualistic co-culture in glucose and nitrite. Strains that were originally isolated from soil, a habitat in common with *C. reinhardtii*, are indicated by an asterisk.

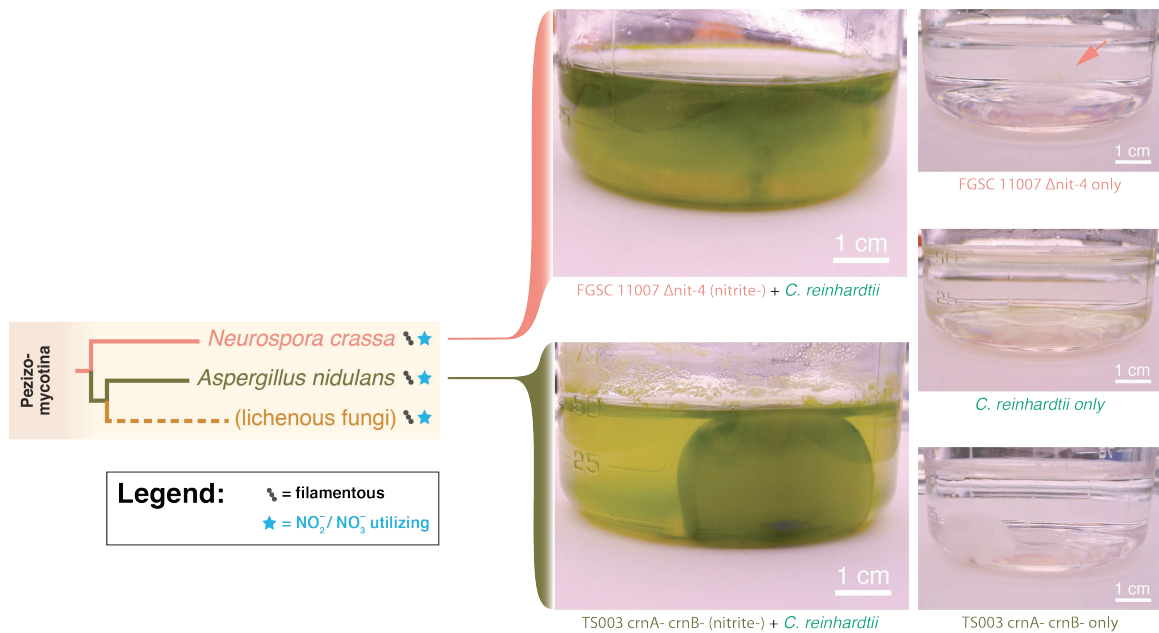






Fig. S5.

Mutualistic, symbiotic associations created between *C. reinhardtii* and model filamentous fungi carrying mutations that prevent nitrite utilization. *Left:* Annotated cladogram of showing the relationship between model filamentous fungi and lichenous fungi (cf. Fig. 3A). *Middle:* Representative 10 day-old co-cultures of *C. reinhardtii* and nitrite-utilization deficient mutants of *N. crassa* (FGSC 11007 Δnit-4; top) or *A. nidulans* (TS003 crnA- crnB-; bottom). *Right:* Corresponding monocultures of *N. crassa* (top), *C. reinhardtii* (middle), and *A. nidulans* (bottom). These model fungi as well as the yeasts shown in Fig. 3 all lack a high-affinity ammonia transporter gene (MEPα) that characterizes many lichen-forming fungi (50, 51); thus, such a transporter is not essential for our synthetic fungal-algal mutualisms.

Fig. S6

Legend:  = +glucose (2%)  = +KNO₂ (10 mM) ← = feeds nitrogen
 = open (+CO₂)  = +KNO₂ (10 mM) + NH₄Cl (20 mM) → = feeds carbon
  = +KNO₂ (20 mM) ↔ = reciprocal exchange

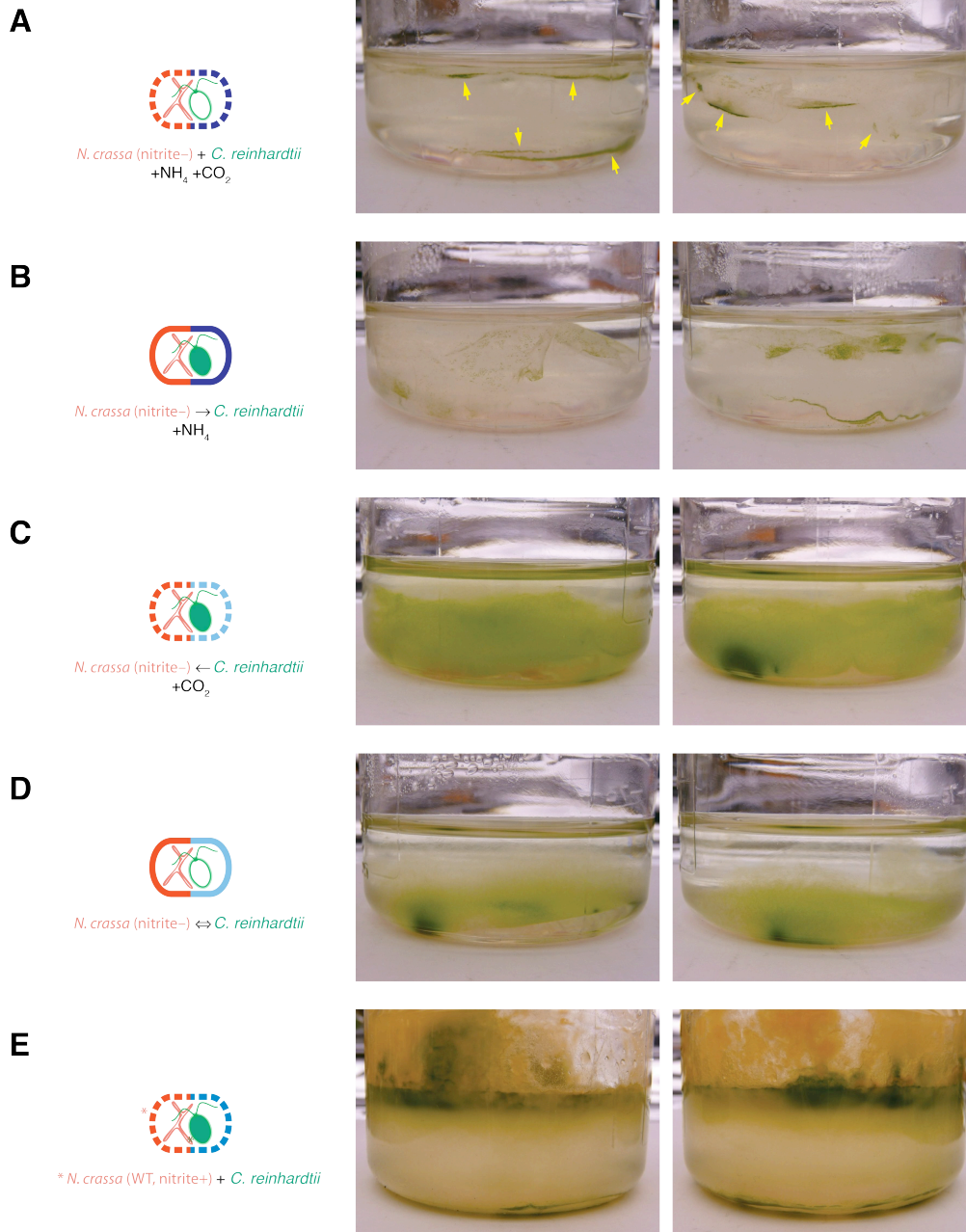


Fig. S6.

Metabolic “short-circuit” experiments demonstrate that the physical association of *C. reinhardtii* to *N. crassa* occurs regardless of any obligate metabolic dependency (see corresponding Movies S7-S11). (A) *N. crassa* mutant (FGSC 11007 $\Delta nit-4$, a non-nitrite-utilizing mutant) grown in co-culture for 8 days with *C. reinhardtii* (CC-1690) in CYM:2% glucose + 10 mM KNO_2 + 20 mM NH_4Cl and access to ambient CO_2 ; under these conditions, neither fungus nor alga requires the other for nitrogen or carbon. *N. crassa* grows faster than *C. reinhardtii*, leading to predominantly white fungal masses, some of whose edges are prominently green (yellow arrows) due to the attachment of *C. reinhardtii* (cf. Movie S7). (B) *N. crassa nit-* mutant grown in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 10 mM KNO_2 + 20 mM NH_4Cl and without access to ambient CO_2 ; under these conditions, the fungus does not require the alga for nitrogen but the alga is dependent on the fungus for CO_2 . As in (A), *N. crassa* grows faster than and independent of *C. reinhardtii*, leading to a predominantly white material, with attached *C. reinhardtii* leading to green spots (left panel) and green peripheral edges (cf. Movie S8). (C) *N. crassa nit-* mutant grown in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 10 mM KNO_2 and access to ambient CO_2 ; under these conditions, the fungus requires the alga for nitrogen but the alga does not require the fungus for CO_2 , producing green masses of the two organisms although the algal density is uniform (cf. Movie S9). (D) *N. crassa nit-* mutant grown in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 10 mM KNO_2 in a sealed environment; under these conditions, both fungus and alga are obligately dependent on one another for nitrogen and carbon (cf. Fig. S5 and Movies S1-S3). This also leads to a balanced, green growth, although the overall yield is lower than (C), where *C. reinhardtii* has access to ambient CO_2 (cf. Movie S10). (E) *N. crassa* (FGSC 2489) wild-type, nitrite-utilizing strain in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 20 mM KNO_2 and access to ambient CO_2 ; under these conditions, as in (A), neither fungus nor alga requires the other for nitrogen or carbon. As in (A), *N. crassa* grows faster than *C. reinhardtii*, leading to a predominantly white fungal material below the gas-liquid interface with algal growth primarily at this interface, where there is likely greater access to ambient CO_2 (cf. Movie S11). The deep orange “fuzz” along the sides of the vessel is due sporulating *N. crassa*, an indication that nitrogen has been significantly depleted in the culture medium (52).

Fig. S7.

Metabolic “short-circuit” experiments demonstrate that the physical association of *C. reinhardtii* to *A. nidulans* occurs regardless of any obligate metabolic dependency (see corresponding Movies S12-S16, and Fig. 4). **(A)** *A. nidulans* (TS003 *crnA- crnB-*, a non-nitrite-utilizing mutant) grown in co-culture for 8 days with *C. reinhardtii* (CC-1690) in CYM:2% glucose + 10 mM KNO₂ + 20 mM NH₄Cl and access to ambient CO₂. Under these conditions, neither fungus nor alga require the other for nitrogen or carbon, and yet composite “balls” of the two species form with a greener core and whiter periphery (cf. Movie S12). **(B)** *A. nidulans nit-* mutant grown in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 10 mM KNO₂ + 20 mM NH₄Cl and without access to ambient CO₂; under these conditions, the fungus does not require the alga for nitrogen but the alga is dependent on the fungus for CO₂. As in (A), composite balls form whose center is greener than their periphery; some *C. reinhardtii* accretion can be seen on top of the balls in the right panel (cf. Movie S13). **(C)** *A. nidulans nit-* mutant grown in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 10 mM KNO₂ and access to ambient CO₂. Under these conditions, the fungus requires the alga for nitrogen but the alga does not require the fungus for CO₂; this results in greener and more compact balls (cf. Movie S14). **(D)** *N. crassa nit-* mutant grown in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 10 mM KNO₂ in a sealed environment; under these conditions, both fungus and alga are obligately dependent on one another for nitrogen and carbon (cf. Fig. S5 and Movies S4-S6). This leads to large green aggregates with concentrations of *C. reinhardtii* on their surface (cf. Movie S15). **(E)** *A. nidulans* (FGSC A4) wild-type, nitrite-utilizing strain in co-culture for 8 days with *C. reinhardtii* in CYM:2% glucose + 20 mM KNO₂ and access to ambient CO₂; under these conditions, as in (A), neither fungus nor alga requires the other for nitrogen or carbon. Green *C. reinhardtii* are embedded amongst *A. nidulans* hyphae (cf. Movie S16), although the lack of *A. nidulans* dependence on *C. reinhardtii* leads to variable degrees of green coloration. Dark brown spots (left panel) are sites of *A. nidulans* spore formation, an indication that carbon or nitrogen has been significantly depleted in the culture medium (53).

Table S1. Strains used in this study.

Species	StrainID	Strain Background	Genotype/Description	Origin/point of isolation	Notes	Source
<i>Saccharomyces cerevisiae</i>	yEH209	S288c: FY4	{S288c: FY4} MATa mal13::MAL63c9(RM11-1a) hoΔ::prENO2-ymTFP1-tmADH1:PTEF-KanMX6-TTEF (haploid)	rotting fig (domesticated lab strain)	Crabtree positive (54)	This study; Fred Winston (Harvard Medical School)
<i>Saccharomyces cerevisiae</i>	yEH213	CEN.PK	{CEN.PK122 segregant} MATa hoΔ::prENO2-ymTFP1-tmADH1:PTEF-KanMX6-TTEF (haploid)	laboratory constructed	Crabtree positive (54)	This study; Maitreya Dunham (U. Washington); (55)
<i>Saccharomyces cerevisiae</i>	yEH1	S1278b	{S1278b} Wild-type MATa (haploid)	laboratory selected	Crabtree positive (54)	Gerald Fink (F10560-5A); http://wiki.yeastgenome.org/index.php/History_of_Sigma
<i>Saccharomyces cerevisiae</i>	yEH251	Ba25 (Mortimer designation)	Wild vineyard isolate (A-RM1) (diploid)	Emilia Romagna vineyard (Italy)	Crabtree positive (54)	Robert Mortimer by way of Andrew Murray
<i>Saccharomyces paradoxus</i>	yEH219	NRRL Y-17217/NCYC 2600/CBS 432	{Type strain} hoΔ::prENO2-ymTFP1-tmADH1:KanMX6 (diploid)	tree exudate/sap (Russia)	Crabtree positive (54)	This study; Dawn Thompson/Aviv Regev (Broad Institute); (56)
<i>Saccharomyces mikatae</i>	yEH223	NBRC 1815/IFO 1815	Wild-type type strain (diploid)	soil (Japan)		Dawn Thompson/Aviv Regev (Broad Institute)
<i>Saccharomyces bayanus</i>	yEH224	NRRL Y-11845/CBS 7001	Wild-type, type strain of <i>Saccharomyces abulensis</i> (diploid)	insect, <i>Mesophylax adopersus</i> (Avila, Spain)	Crabtree positive (54)	ARS Culture Collection
<i>Candida glabrata</i>	yEH226	NRRL Y-65/CBS 138/ATCC 2001	Wild-type type strain (haploid)	human feces (USA)	Crabtree positive (57, 58)	ARS Culture Collection
<i>Saccharomyces castellii</i>	yEH228	NRRL Y-12630/CB 4309	Wild-type type strain also known as <i>Naumovozyma castellii</i>	soil (Finland)	Crabtree positive (57)	ARS Culture Collection
<i>Kluyveromyces polysporus</i>	yEH229	DSM 70294/CBS 2163	Wild-type type strain, also known as <i>Vanderwaltozyma polyspora</i> (haploid)	soil (South Africa)		Dawn Thompson/Aviv Regev (Broad Institute)
<i>Zygosaccharomyces rouxii</i>	yEH232	NRRL Y-229/CBS 732	Wild-type type strain (haploid)	must of black grapes (Italy)	Weakly Crabtree positive (57)	ARS Culture Collection
<i>Saccharomyces kluyveri</i>	yEH233	NRRL Y-12651/CBS 3082	Wild-type type strain, also known as <i>Lachancea kluyveri</i> (diploid)	insect, <i>Drosophila pinicola</i> (California, USA)	Weakly Crabtree positive (59)	Dawn Thompson/Aviv Regev (Broad Institute)
<i>Kluyveromyces waltii</i>	yEH231	NCYC 2644/CBS 640	Wild-type type strain, also known as <i>Lachancea waltii</i>	exudate of holly, <i>Ilex integra</i> (Japan)		Dawn Thompson/Aviv Regev (Broad Institute)
<i>Kluyveromyces marxianus</i>	yEH245	NRRL Y-8281/CBS 712	Wild-type type strain	unknown	Crabtree negative (60)	ARS Culture Collection
<i>Kluyveromyces lactis</i>	yEH235	CLIB 209/NRRL Y-1140/CBS 2359	Wild-type MATa (haploid)	dairy cream (Illinois, USA)	Crabtree negative (57)	Dawn Thompson/Aviv Regev (Broad Institute)
<i>Hansenula polymorpha</i>	yEH236	NRRL Y-5445/CBS 4732	Wild-type type strain, also known as <i>Pichia augusta</i> and <i>Ogataea polymorpha</i> (diploid)	soil, irrigated with distillery effluent (Pernambuco, Brazil)	Crabtree negative (60)	ARS Culture Collection
<i>Candida albicans</i>	yEH237	SC5314/ATCC MYA-2876	Wild-type diploid (diploid)	human with candidiasis (USA)	Crabtree negative (60)	Dawn Thompson/Aviv Regev (Broad Institute); (60)
<i>Pichia pastoris</i>	yEH238	NRRL Y-1603/CBS 704/DSM 70382	Wild-type type strain, also known as <i>Komagataella pastoris</i> and <i>Zygosaccharomyces pastoris</i> (haploid)	exudate of chestnut tree, <i>Castanea</i> sp. (near Lyon, France)	Crabtree negative (61)	ARS Culture Collection
<i>Yarrowia lipolytica</i>	yEH240	NRRL YB-423/CBS 6124	Wild-type anamorph of <i>Candida lipolytica</i> , type strain of <i>Endomycopsis lipolytica</i> (haploid)	fiber tailings from corn processing plant (Illinois)		ARS Culture Collection
<i>S. pombe</i>	yEH143	975/FWP9	{strain 975} Wild-type h+ (haploid)	rancid French wine (Switzerland)	Crabtree positive (60)	Fred Winston (Harvard Med School); http://www-bcf.usc.edu/~forsburg/main.html
<i>S. japonicus</i>	yEH285	yFS275/ATCC 10660	Wild-type	strawberry wine (Japan)		Nick Rhind (U. Massachusetts Medical Center)
<i>Neurospora crassa</i>	FEH20	FGSC 11007	<i>Mat A Δnit-4:NCU08294</i> (homokaryon knockout of nitrite/nitrate assimilation transcription factor) (haploid)	laboratory constructed	Crabtree negative (62)	Fungal Genetics Stock Center
<i>Neurospora crassa</i>	FEH10	FGSC 2489/74-OR23-1VA	Wild-type (Oak Ridge) inbred reference strain, <i>Mat A</i> (haploid)	laboratory selected	Crabtree negative (62)	Fungal Genetics Stock Center; http://www.fgsc.net/fgn51/fgn51perkins.html
<i>Aspergillus nidulans</i>	FEH6	TS003	<i>alpha/HMG biA1 pabaA1 argB2 crnA747 crnB110 veA1 argB-plasmid-complemented</i> (defect in nitrate/nitrite transporters crnA/B, leading to very poor growth on nitrate/nitrite) (haploid)	laboratory constructed	Crabtree negative (63); obligate aerobe	Thorsten Schinko/Joseph Strauss; (64)
<i>Aspergillus nidulans</i>	FEH8	FGSC A4	Wild-type (Glasgow) reference strain, <i>alpha/HMG veA+</i> (haploid)	laboratory selected	Crabtree negative (63); obligate aerobe	Fungal Genetics Stock Center; http://www.fgsc.net/scripts/StrainSearchReturnPage.asp?OrgID=4
<i>Chlamydomonas reinhardtii</i>	CC-1690	21gr (Massachusetts)	{21gr} Wild-type (nitrate+) (haploid)	originally, soil from potato field (Amherst, MA, USA; laboratory selected by Ruth Sager and domesticated)		Chlamydomonas Resource Center; (65)
<i>Chlamydomonas reinhardtii</i>	CC-2290	S1 D2 (Minnesota)	{S1 D2} Wild isolate (nitrate+) (haploid)	soil (Plymouth, MN, USA)		Chlamydomonas Resource Center; (66)
<i>Chlamydomonas reinhardtii</i>	CC-2342	CC-2342 (Pennsylvania)	Wild isolate, mt- [Jarvik #6] (nitrate+) (haploid)	soil (Pittsburgh, PA)		Chlamydomonas Resource Center; (67)
<i>Chlamydomonas reinhardtii</i>	CC-2931	CC-2931 (North Carolina)	Wild isolate, mt- [Harris #6] (nitrate+) (haploid)	soil from residential garden (Durham, NC, USA)		Chlamydomonas Resource Center
<i>Chlamydomonas reinhardtii</i>	CC-2935	CC-2935 (Quebec)	Wild isolate, mt- [Bell/LEE-1] (nitrate+) (haploid)	soil from cultivated field (Farnham, Québec, Canada)		Chlamydomonas Resource Center; (68)
<i>Chlamydomonas incerta</i>	CC-3871	CC-3871/SAG 7.73 (Cuba)	Wild-type, mt-; closely related to but cannot mate with <i>C. reinhardtii</i> ; missing <i>fus1</i> homolog necessary for gamete adhesion, resulting in prezygotic isolation (nitrate+) (haploid)	soil from botanical garden (Havana, Cuba; presumed)		Chlamydomonas Resource Center; (69)
<i>Chlamydomonas moewusii</i>	CC-55	CC-55/UTEX 97 (New York)	Wild-type, mt+; also known as <i>Chlamydomonas eugametos</i> var. <i>moewusii</i> (nitrate+) (haploid)	freshwater (New York, USA)		Chlamydomonas Resource Center; (70)
<i>Chlamydomonas monoica</i>	CC-1904	CC-1904/UTEX 2020 (Alabama)	Wild-type, homothallic; also known as <i>C. moewusii</i> var. <i>monoica</i> ; considered part of the <i>C. noctigama</i> clade but exhibits different mating characteristics (nitrate+) (haploid)	Black Warrior River (Alabama, USA)		Chlamydomonas Resource Center; (71)
<i>Chlorella vulgaris</i>	SAG 211-12	SAG 211-12/UTEX 30 (Geneva)	Wild-type (glucose+, nitrate+) (haploid)	freshwater (Geneva, Switzerland)		Sammlung von Algenkulturen

Additional Notes:

- 1) Phylogenetic relationships of ascomycetous fungi, see (33, 34)
- 2) Dating of fungal branch points, see (35-37)
- 3) Ploidy of yeasts, see (33)
- 4) Phylogenetic relationship of algal strain, see (38)
- 5) Dating of algal branch points, see (39-41)

Supplementary Movies S1-S3: Z-stack series of a representative symbiotic associations formed between *N. crassa* (*nit*-) and *C. reinhardtii* cultures under conditions that enforce mutualism. These movies serve to complement Fig. 4A in showing that the co-localization of *C. reinhardtii* on *N. crassa* hyphae is not merely due to a “hovering” of swimming algae in the vicinity; *C. reinhardtii* appear to be attached to hyphae, are resistant to gentle hydrodynamic displacement by liquid flow, and are predominantly immotile/de-flagellated.

Movie S1.

Low magnification (13X) z-stack movie series of a representative mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 *Δnit-4*, a non-nitrite-utilizing mutant) and *C. reinhardtii* (CC-1690) showing two types of organizational structure: (i) an even, dense distribution of *C. reinhardtii* cells (green) embedded within a dense meshwork of white, fine-haired fungal hyphae (top right), and (ii) less dense, peripheral regions where algal cells are concentrated into patches (left and middle). Each plane was imaged for 0.2 seconds and is 32 μm apart in the z-axis. Fungi and algae were co-cultured in 50 mL of CYM:2% glucose + 10 mM KNO₂ medium in an air-tight, 150 mL polystyrene container (Corning) under constant light (irradiance=45 μmol/m²/s) for 7 days (15). A segment of the resulting aggregate was sterilely dissected, transferred to a plastic petri dish, bathed in CYB medium, and imaged from the top using a Zeiss Axio Zoom.V16 stereomicroscope.

Movie S2.

Z-stack image series at 50X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 *Δnit-4*) and *C. reinhardtii* (CC-1690). Each plane was imaged for 0.2 seconds and is 3 μm apart in the z-axis. Numerous *C. reinhardtii* cells (green dots) can be seen coating strands of *N. crassa* hyphae (white filaments). The imaged sample was occasionally gently tapped to induce movement of the dissected fragment and liquid flow, and to demonstrate the attachment of *C. reinhardtii* cells to fungal hyphae. Fungi and algae were co-cultured and imaged as in Movie S1.

Movie S3.

Close-up of a representative mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 *Δnit-4*) and *C. reinhardtii* (CC-1690). The movie shows a z-stack, in which each plane was imaged for 0.2 seconds and is 1 μm apart in the z-axis. The images are shown at 100X magnification and 2X real time playback. They show *C. reinhardtii* cells (green) attached to *N. crassa* hyphae (white filaments) like beads on a string. Several *C. reinhardtii* cells can be seen freely swimming or jittering while attached to fungal hyphae (some via their flagella, e.g., see upper left, time = 10-12 sec). Fungi and algae were co-cultured and imaged as in Movie S1.

Supplementary Movies S4-S6: Z-stack series of a representative symbiotic associations formed between *A. nidulans* (*nit*-) (a distant relative of lichenous fungi) and *C. reinhardtii* under conditions that enforce mutualism. These movies serve to complement Fig. 4B in showing that the co-localization of *C. reinhardtii* on *A.*

***nidulans* hyphae is not merely due to a “hovering” of swimming algae in the vicinity; *C. reinhardtii* appear to be attached to hyphae in small clusters, are resistant to gentle hydrodynamic displacement by liquid flow, and are not visibly motile.**

Movie S4.

Low magnification (20X) z-stack series of a representative mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-, a non-nitrite-utilizing mutant) and *C. reinhardtii* (CC-1690) showing a patchy distribution of *C. reinhardtii* (green) cells on top of a homogeneously green tissue center (top), a predominantly white hyphal periphery (lower right), and a greening edge (lower left). Each plane was imaged for 0.2 seconds and is 15 μm apart in the z-axis. Fungi and algae were co-cultured and imaged as in Movie S1.

Movie S5.

Z-stack image series at 50X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-) and *C. reinhardtii* (CC-1690) (region in Movie S2, expanded). Each plane was imaged for 0.2 seconds and is 3 μm apart in the z-axis. *C. reinhardtii* cells (green) have a patchy distribution and are associated with *A. nidulans* hyphae (which are more densely packed than *N. crassa* hyphae) like clusters of grapes. Fungi and algae were co-cultured and imaged as in Movie S1.

Movie S6.

Close-up of a representative mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-) and *C. reinhardtii* (CC-1690). The movie shows a z-stack, in which each plane was imaged for 0.2 seconds and is 2 μm apart in the z-axis. The images are shown at 100X magnification and 2X real time playback. They show *C. reinhardtii* cells (green) attached on top of and between densely packed *A. nidulans* hyphae (white filaments) like clusters of grapes on a vine. Several *C. reinhardtii* cells can be seen swimming within the fungal hyphal network and near the periphery of the algal-fungal association (e.g., time = 10-14 sec). Fungi and algae were co-cultured and imaged as in Movie S1.

Supplementary Movies S7-S11: Representative associations formed between *N. crassa* and *C. reinhardtii* in series of combinatorial control experiments, demonstrating that physical association between fungus and alga is independent of any metabolic dependency (see Fig. S6).

Movie S7.

Z-stack image series at 50X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 *Anit-4*, a non-nitrite-utilizing mutant) and *C. reinhardtii* (CC-1690) grown together in a loosely capped 150 mL polystyrene container (Corning) containing 50 mL of CYM:2% glucose + 10 mM KNO_2 + 20 mM NH_4Cl medium, under constant light (irradiance=45 $\mu\text{mol}/\text{m}^2/\text{s}$) for 7 days (15). Under these conditions, *N.*

crassa (*nit*-) does not require *C. reinhardtii* for nitrogen nor does *C. reinhardtii* require *N. crassa* for carbon (cf. Fig. S6A); despite the lack of metabolic dependency, both fungus and alga still physically associate. In addition to regions of densely packed green *C. reinhardtii* cells within a *N. crassa* hyphal network, several *C. reinhardtii* cells or cluster of cells can be seen attached to moving *N. crassa* hyphae (white filaments). The fragment was imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 4 μm apart in the z-axis.

Movie S8.

Z-stack image series at 112X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 $\Delta nit-4$) and *C. reinhardtii* (CC-1690) co-cultured in 50 mL of CYM:2% glucose + 10 mM KNO_2 + 20 mM NH_4Cl medium in an air-tight, 150 mL polystyrene container (Corning) under constant light (irradiance=45 $\mu\text{mol}/\text{m}^2/\text{s}$) for 7 days (15). Under these conditions, *N. crassa* (*nit*-) does not require *C. reinhardtii* for nitrogen, although *C. reinhardtii* is obligately dependent on *N. crassa* for carbon (this feeding dependency is indicated in the movie as *N. crassa* \rightarrow *C. reinhardtii*) (cf. Fig. S6B). The fragment was imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 1 μm apart in the z-axis.

Movie S9.

Z-stack image series at 112X magnification and 2X real time playback of a representative fragment of mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 $\Delta nit-4$) and *C. reinhardtii* (CC-1690) co-cultured in a loosely capped 150 mL polystyrene container (Corning) containing 50 mL of CYM:2% glucose + 10 mM KNO_2 medium, under constant light (irradiance=45 $\mu\text{mol}/\text{m}^2/\text{s}$) for 7 days (15). Under these conditions, *N. crassa* (*nit*-) requires *C. reinhardtii* for nitrogen (this feeding dependency is indicated in the movie as *N. crassa* \leftarrow *C. reinhardtii*), while *C. reinhardtii* does not require *N. crassa* for carbon (cf. Fig. S6C). Patchy clusters of *C. reinhardtii* cells are seen within a *N. crassa* hyphal network of white filaments. The fragment was imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 2 μm apart in the z-axis.

Movie S10.

Z-stack image series at 112X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *N. crassa* (FGSC 11007 $\Delta nit-4$) and *C. reinhardtii* (CC-1690) co-cultured in 50 mL of CYM:2% glucose + 10 mM KNO_2 medium in an airtight 150 mL polystyrene container (Corning), under constant light (irradiance=45 $\mu\text{mol}/\text{m}^2/\text{s}$) for 7 days (15). Under these conditions, *N. crassa* (*nit*-) requires *C. reinhardtii* for nitrogen and *C. reinhardtii* requires *N. crassa* for carbon (this feeding dependency is indicated in the movie as *N. crassa* \leftrightarrow *C. reinhardtii*) (cf. Fig. S6D); these conditions are the same as in Movies S1-S3 and imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 1.4 μm apart in the z-axis.

Movie S11.

Z-stack image series at 112X magnification of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between a wild-type nitrite-utilizing *N. crassa* strain (FGSC 2489) and *C. reinhardtii* (CC-1690) co-cultured in a loosely capped 150 mL polystyrene container (Corning) containing 50 mL of CYM:2% glucose + 20 mM KNO₂ medium, under constant light (irradiance=45 μmol/m²/s) for 7 days (15). Wild-type *N. crassa* does not require *C. reinhardtii* for nitrogen nor does *C. reinhardtii* require *N. crassa* for carbon (cf. Fig. S5E); this is a similar metabolic arrangement as in Movie S7, achieved using a nitrite-utilizing fungal strain. Despite the lack of metabolic dependency, both fungus and alga still physically associate. This fragment was imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 3 μm apart in the z-axis.

Supplementary Movies S12-S16: Representative associations formed between *A. nidulans* (a distant relative of lichenous fungi) and *C. reinhardtii* in series of control experiments, demonstrating that physical association between fungus and alga is independent of any metabolic dependency (see Fig. S7). Cf. images of algal attachments to fungal hyphae in lichens (72).

Movie S12.

Z-stack image series at 90X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-, a non-nitrite-utilizing mutant) and *C. reinhardtii* (CC-1690) grown together in a loosely capped 150 mL polystyrene container (Corning) containing 50 mL of CYM:2% glucose + 10 mM KNO₂ + 20 mM NH₄Cl medium, under constant light (irradiance=45 μmol/m²/s) for 7 days (15). Under these conditions, *A. nidulans* (*nit*-) does not require *C. reinhardtii* for nitrogen nor does *C. reinhardtii* require *A. nidulans* for carbon (cf. Fig. S7A); despite the lack of metabolic dependency, both fungus and alga still physically associate. *C. reinhardtii* cells (green) cluster onto *A. nidulans* hyphae (white filaments) like “grapes on a vine.” Fungi and algae were imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 10 μm apart in the z-axis.

Movie S13.

Z-stack image series at 112X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-, *nit*-) and *C. reinhardtii* (CC-1690) co-cultured in 50 mL of CYM:2% glucose + 10 mM KNO₂ + 20 mM NH₄Cl medium in an air-tight, 150 mL polystyrene container (Corning) under constant light (irradiance=45 μmol/m²/s) for 7 days (15). Under these conditions, *A. nidulans* (*nit*-) does not require *C. reinhardtii* for nitrogen, although *C. reinhardtii* is obligately dependent on *A. nidulans* for carbon (this feeding dependency is indicated in the movie as *A. nidulans* → *C. reinhardtii*) (cf. Fig. S7B). As in Movie S12, *C. reinhardtii* cells (green) cluster onto *A. nidulans* hyphae (white filaments). Fungi and algae were imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 10 μm apart in the z-axis.

Movie S14.

Z-stack image series at 112X magnification and 2X real time playback of a representative fragment of mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-, *nit*-) and *C. reinhardtii* (CC-1690) co-cultured in a loosely capped 150 mL polystyrene container (Corning) containing 50 mL of CYM:2% glucose + 10 mM KNO₂ medium, under constant light (irradiance=45 μmol/m²/s) for 7 days (15). Under these conditions, *A. nidulans* (*nit*-) requires *C. reinhardtii* for nitrogen (this feeding dependency is indicated in the movie as *A. nidulans* ← *C. reinhardtii*), while *C. reinhardtii* does not require *N. crassa* for carbon (cf. Fig. S7C). In contrast to Movies S12 and S13, under these conditions, *C. reinhardtii* cells are much more densely packed in association with *A. nidulans* hyphal network of white filaments (cf. smaller, compact tissue ball in Fig. S7C), akin to the greener, interior sections of fragments in Movies S4-S6. Fungi and algae were imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 10 μm apart in the z-axis.

Movie S15.

Z-stack image series at 112X magnification and 2X real time playback of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between *A. nidulans* (TS003 *crnA*- *crnB*-, *nit*-) and *C. reinhardtii* (CC-1690) co-cultured in 50 mL of CYM:2% glucose + 10 mM KNO₂ medium in an airtight 150 mL polystyrene container (Corning) containing, under constant light (irradiance=45 μmol/m²/s) for 7 days (15). Under these conditions, *A. nidulans* (*nit*-) requires *C. reinhardtii* for nitrogen and *C. reinhardtii* requires *N. crassa* for carbon (this dependency is indicated in the movie as *A. nidulans* ↔ *C. reinhardtii*) (cf. Fig. S7D); these conditions are the same as in Movies S4-S6 and imaged as described in Movie S1, although the illumination arrangement is different. The clustering of *C. reinhardtii* cells on *A. nidulans* white hyphal filaments is dense, but not as extreme as in Movie S14 (*A. nidulans* ← *C. reinhardtii*; cf. Movies S4). Each plane was imaged for 0.2 seconds and is 10 μm apart in the z-axis.

Movie S16.

Z-stack image series at 111X magnification of a representative peripheral fragment of mutualistic symbiotic association formed by the interaction between a wild-type nitrite-utilizing *A. nidulans* strain (FGSC A4) and *C. reinhardtii* (CC-1690) co-cultured in a loosely capped 150 mL polystyrene container (Corning) containing 50 mL of CYM:2% glucose + 20 mM KNO₂ medium, under constant light (irradiance=45 μmol/m²/s) for 7 days (15). Wild-type *A. nidulans* does not require *C. reinhardtii* for nitrogen nor does *C. reinhardtii* require *A. nidulans* for carbon (cf. Fig S7E); this is a similar metabolic arrangement as in Movie S12, achieved using a nitrite-utilizing fungal strain. Despite the lack of metabolic dependency, both fungus and alga still physically associate. However, *C. reinhardtii* cells are less densely clustered compared to Movie S12, and substantially less so than in Movie S14 (*A. nidulans* ← *C. reinhardtii*). Thus, the degree of dependency of *A. nidulans* on *C. reinhardtii* for nitrogen appears to dictate the density of *C. reinhardtii* cells embedded within the *A. nidulans* hyphal network. This fragment was

imaged as described in Movie S1, although the illumination arrangement is different. Each plane was imaged for 0.2 seconds and is 20 μm apart in the z-axis.

Additional Dataset S1 (separate file in zip archive)

Data table for Fig. 1B graphs.

Additional Dataset S2 (separate file in zip archive)

Data table for Fig. 2.

Additional Dataset S3 (separate file in zip archive)

Data table for Fig. 3 graphs.

Additional Dataset S4 (separate file in zip archive)

Data table for Fig. S1B graphs.

Additional Dataset S5 (separate file in zip archive)

Data table for Fig. S2.

Additional Dataset S6 (separate file in zip archive)

Data table for Fig. S3 graphs.

Additional Dataset S7 (separate file in zip archive)

Data table for Fig. S4.