1 Recipient-biased competition for a cross-fed nutrient is required 2 for coexistence of microbial mutualists 3 Alexandra L. McCully, Breah LaSarre, James B. McKinlay<sup>#</sup> 4 Department of Biology, Indiana University, Bloomington, IN 5 6 Running title: Biased competition within a mutualism <sup>#</sup>Corresponding author. 1001 E 3<sup>rd</sup> Street, Jordan Hall, Bloomington, IN 47405 7 8 Phone: 812-855-0359 9 Email: jmckinla@indiana.edu 10 Conflict of interest. 11 The authors declare no conflict of interest.

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**Abstract.** Many mutualistic microbial relationships are based on nutrient cross-feeding. Traditionally, cross-feeding is viewed as being unidirectional from the producer to the recipient. This is likely true when a producer's metabolic waste, such as fermentation products, provides carbon for a recipient. However, in some cases the cross-fed nutrient holds value for both the producer and the recipient. In such cases, there is potential for nutrient reacquisition by producer cells in a population, leading to competition against recipients. Here we investigate the consequences of inter-partner competition for cross-fed nutrients on mutualism dynamics using an anaerobic coculture pairing fermentative Escherichia coli and phototrophic Rhodopseudomonas palustris. In this coculture, E. coli excretes waste organic acids that provide carbon for R. palustris. In return, R. palustris cross-feeds E. coli ammonium (NH<sub>4</sub><sup>+</sup>), a valuable nitrogen compound that both species prefer. To explore the potential for inter-partner competition, we first used a kinetic model to simulate cocultures with varied affinities for NH<sub>4</sub><sup>+</sup> in each species. The model predicted that inter-partner competition for cross-fed NH<sub>4</sub> could profoundly impact population dynamics. We then experimentally tested the predictions by culturing mutants lacking NH<sub>4</sub><sup>+</sup> transporters in both NH<sub>4</sub><sup>+</sup> competition assays and cooperative cocultures. Both theoretical and experimental results indicated that the recipient must have a competitive advantage in acquiring valuable cross-fed NH<sub>4</sub><sup>+</sup> to avoid collapse of the mutualism. Thus, the very metabolites that form the basis for cooperative cross-feeding can also be subject to competition between mutualistic partners. **Significance.** Mutualistic relationships, particularly those based on nutrient cross-feeding, promote stability of diverse ecosystems and drive global biogeochemical cycles. Cross-fed nutrients within these systems can be either waste products valued only by one partner or nutrients that both partners value. Here, we explore how inter-partner competition for a communally-valuable cross-fed nutrient impacts mutualism dynamics. We discovered that mutualism stability necessitates that the recipient have a competitive advantage against the producer in obtaining the cross-fed nutrient. We propose that the requirement for recipient-biased competition is a general rule for mutualistic coexistence based on the transfer of communally valuable resources, microbial or otherwise.

### Introduction

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Mutualistic cross-feeding of resources between microbes can have broad impacts ranging from influencing host health (1, 2) to driving global biogeochemical cycles (3–6). Cross-fed metabolites are often regarded as nutrients due to the value they provide to a dependent partner, the recipient. However, for the partner producing the nutrient, the producer, a cross-fed nutrient's value can vary. On one extreme, the cross-fed metabolite is valued by the recipient but not the producer, as is the case for fermentative waste products (7–10). In other cases, a cross-fed metabolite holds value for both the recipient and the producer, as is the case for vitamin  $B_{12}$  (6, 11, 12) and ammonium (NH<sub>4</sub><sup>+</sup>) (13, 14). Such communallyvaluable cross-fed nutrients are subject to partial privatization (15), wherein the producer has mechanisms to retain a portion of the nutrient pool for itself. While most mutualism cross-feeding studies only consider unidirectional metabolite transfer from producer to recipient, we wondered whether these mechanisms for partial privatization could lead to competition between partner populations for communally-valuable cross-fed nutrients. It seems likely that such competition could influence mutualism stability, as is known to be the case for competition for exogenous limiting resources (8, 16–19). To the best of our knowledge inter-partner competition for cross-fed nutrients and its impact on mutualism dynamics have never been investigated. One example of cross-feeding that could involve competition between mutualistic partners is NH<sub>4</sub><sup>+</sup> excretion by N<sub>2</sub>-fixing bacteria (Fig. 1A), called N<sub>2</sub>-fixers (13, 14). During N<sub>2</sub> fixation, the enzyme nitrogenase converts N<sub>2</sub> gas into two NH<sub>3</sub> (20). In an aqueous environment, NH<sub>3</sub> is in equilibrium with NH<sub>4</sub><sup>+</sup>. At neutral pH, NH<sub>4</sub><sup>+</sup> is the predominant form but small amounts of NH<sub>3</sub> can potentially leave the cell by diffusion across the membrane (21) (Fig. 1B). This inherent 'leakiness' for NH<sub>3</sub> likely fosters NH<sub>4</sub><sup>+</sup> cross-feeding, as extracellular NH<sub>3</sub> is available to neighboring microbes. Importantly, these neighbors can include clonal N<sub>2</sub>-fixers, as NH<sub>3</sub>/NH<sub>4</sub><sup>+</sup> is a preferred nitrogen source for most microbes. At concentrations above 20 μM, NH<sub>3</sub> can be acquired by passive diffusion; below 20 μM, NH<sub>4</sub><sup>+</sup> is bound and transported as NH<sub>3</sub> by AmtB transporters (Fig. 1B) (22). AmtB-like transporters are conserved throughout all domains of life (23). There is growing evidence that AmtB is used by N<sub>2</sub>-fixers to recapture NH<sub>3</sub> lost

by passive diffusion, as  $\Delta AmtB$  mutants accumulate  $NH_4^+$  in culture supernatants whereas wild-type strains do not (24–26). Thus, during  $NH_4^+$  cross-feeding, AmtB likely facilitates both  $NH_4^+$  acquisition by the mutualistic partner and recapture of  $NH_4^+$  by the  $N_2$ -fixer.

Assessing the effects of inter-partner competition for a cross-fed nutrient would require a level of experimental control not possible in most natural settings. However, synthetic microbial communities, or cocultures, are well-suited to address such questions (27–29). We previously developed a bacterial coculture that features cross-feeding of waste products (organic acids) from *Escherichia coli*, and a communally-valuable nutrient (NH<sub>4</sub><sup>+</sup>) from *Rhodopseudomonas palustris* Nx (Fig. 1A) (26). Here, using both a kinetic model and genetic manipulation to alter the affinity of each species in the coculture for NH<sub>4</sub><sup>+</sup>, we demonstrate that inter-partner competition for cross-fed NH<sub>4</sub><sup>+</sup> plays a direct role in maintaining coexistence. Specifically, insufficient competition by *E. coli* for NH<sub>4</sub><sup>+</sup> resulted in a collapse of the mutualism. Mutualism collapse could be delayed or potentially avoided through higher net NH<sub>4</sub><sup>+</sup> excretion by *R. palustris* or increased *E. coli* population size. Our results suggest that, as a general rule, competition for a cross-fed nutrient in an obligate mutualism must be biased in favor of the recipient to avoid mutualism collapse and the potential extinction of both species.

### Results

Competition for cross-fed  $NH_4^+$  is predicted to shape mutualism population dynamics. Within our coculture (Fig. 1A), *E. coli* (*Ec*) ferments sugars into waste organic acids, providing essential carbon and electrons to *R. palustris* (*Rp*) Nx. *R. palustris* Nx is genetically engineered to excrete low micromolar amounts of  $NH_4^+$ , providing essential nitrogen for *E. coli* (26).  $NH_4^+$  excretion by *R. palustris* Nx is due to mutation of NifA, the master transcriptional regulator of nitrogenase, which results in constitutive nitrogenase activity even in the presence of normally inhibitory  $NH_4^+$  (30). In contrast to organic acids, which are only useful to *R. palustris*,  $NH_4^+$  produced by *R. palustris* Nx is essential for the growth of both species; *R. palustris* uses some  $NH_4^+$  for its own biosynthesis and excretes the rest, which serves as the nitrogen source for *E. coli*. However, *R. palustris* Nx can also take up  $NH_4^+$  (30). Thus, we hypothesized

that competition for cross-fed NH<sub>4</sub><sup>+</sup> between the *R. palustris* Nx producer population and the *E. coli* recipient population could influence mutualism dynamics.

We first explored whether competition for cross-fed  $NH_4^+$  could affect the mutualism using SyFFoN, a mathematical model describing our coculture (26, 31). SyFFoN simulates population and metabolic dynamics in batch cocultures using Monod equations with experimentally-determined parameter values. As previous versions described  $NH_4^+$  uptake kinetics only for *E. coli* (26, 31), we amended SyFFoN to include both an *R. palustris*  $NH_4^+$  uptake affinity ( $K_m$ ) and higher *R. palustris* maximum growth rate ( $\mu_{MAX}$ ) when  $NH_4^+$  is used (SI Appendix Table S1). We then simulated batch cocultures wherein the relative affinity for  $NH_4^+$  varied between the two species (Fig. 2). The model predicted that coexistence is maintained when the *R. palustris* affinity for  $NH_4^+$  is low relative to that of *E. coli* (Rp:Ec < 1); sufficient  $N_2$  is converted to  $NH_4^+$  to support *R. palustris* growth and enough  $NH_4^+$  is cross-fed to support *E. coli* growth. In contrast, when the *R. palustris* affinity for  $NH_4^+$  is high relative to that of *E. coli* (Rp:Ec > 1), *E. coli* growth is no longer supported because *E. coli* cannot compete for excreted  $NH_4^+$ . However, high *R. palustris* cell densities were still predicted (Fig. 2) due to persistent, low-level organic acid cross-feeding stemming from *E. coli* maintenance metabolism, which can support *R. palustris* growth even when *E. coli* is not growing (31).

Genetic disruption of AmtB  $NH_4^+$  transporters affects relative affinities for  $NH_4^+$ . Bacterial cells generally acquire  $NH_4^+$  through two mechanisms: passive diffusion of  $NH_3$ , or active uptake by AmtB transporters (Fig. 1B). We hypothesized that deleting *amtB* genes in either species would result in a lower affinity for  $NH_4^+$  in that species and thus could be used to test how relative  $NH_4^+$  affinity impacts coculture dynamics. We generated  $\Delta$ AmtB mutants of both *E. coli* and *R. palustris* and first characterized the effect of the mutations in monoculture. Deletion of *amtB* in *E. coli* had no effect on growth or fermentation profiles when  $NH_4Cl$  was in excess (SI Appendix Fig. S1), consistent with previous observations where  $\Delta$ AmtB growth defects were only apparent at  $NH_4^+$  concentrations below 20  $\mu$ M (22). In *R. palustris*  $\Delta$ AmtB monocultures with  $N_2$  as the nitrogen source, growth trends were equivalent to

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those of the parent strain; however, R. palustris ΔAmtB excreted more NH<sub>4</sub><sup>+</sup> than the parent strain and about a third of that excreted by R. palustris Nx (SI Appendix Fig. S1C and D). In line with our hypothesis,  $NH_4^+$  excretion by R. palustris  $\Delta$ AmtB could be due to a decreased ability to reacquire  $NH_4^+$ lost by diffusion, resulting in increased net NH<sub>4</sub><sup>+</sup> excretion. Alternatively, we considered that NH<sub>4</sub><sup>+</sup> excretion by R. palustris ΔAmtB could be due to improper nitrogenase regulation. In several other N<sub>2</sub>fixers, proper nitrogenase regulation requires AmtB, for example to induce post-translational nitrogenase inhibition (switch-off) in response to NH<sub>4</sub> $^+$  (25, 32). We tested whether R. palustris  $\Delta$ AmtB exhibits NH<sub>4</sub><sup>+</sup>-induced switch-off by adding NH<sub>4</sub>Cl to exponentially growing cultures and measuring H<sub>2</sub> production, an obligate product of the nitrogenase reaction (33), as a proxy for nitrogenase activity. Upon adding NH<sub>4</sub>Cl, H<sub>2</sub> production stopped in R. palustris  $\Delta$ AmtB cultures. In contrast, H<sub>2</sub> production only slowed slightly in R. palustris Nx cultures (SI Appendix Fig. S2), consistent with previous observations for NifA\* strains (34, 35). Additionally, like the parent strain, R. palustris ΔAmtB did not produce H<sub>2</sub> when grown with NH<sub>4</sub><sup>+</sup>, unlike R. palustris Nx (SI Appendix Fig. S3). These observations demonstrate that R. palustris ΔAmtB is competent for NH<sub>4</sub><sup>+</sup>-induced nitrogenase repression, and thus NH<sub>4</sub><sup>+</sup> excretion by R. palustris  $\triangle$ AmtB is likely due to a poor ability to reacquire NH<sub>4</sub><sup>+</sup> lost by diffusion. To test our hypothesis that deleting *amtB* would lower cellular affinity for NH<sub>4</sub><sup>+</sup>, we directly competed all possible E. coli and R. palustris strain combinations in competition assays where ample carbon was available for each species but the NH<sub>4</sub><sup>+</sup> concentration was kept low; specifically, a small amount of NH<sub>4</sub><sup>+</sup> was added every hour to bring the final NH<sub>4</sub><sup>+</sup> concentration to 5 μM (Fig. 3). In this competition assay, the species that is more competitive for NH<sub>4</sub><sup>+</sup> should reach a higher cell density than the other species. In all cases, WT E. coli was more competitive for NH<sub>4</sub><sup>+</sup> than R. palustris. However, each R. palustris strain was able to outcompete E. coli  $\triangle$ AmtB (Fig. 3), even though the R. palustris maximum growth rate is 4.6-times slower than that of E. coli (SI Appendix Fig. S1). Even R. palustris strains lacking AmtB outcompeted E. coli \( \Delta \text{AmtB} \) (Fig. 3), indicating that R. palustris has a higher affinity for NH<sub>4</sub><sup>+</sup> than E. coli independent of AmtB. These data confirmed that deletion of amtB was an effective means by which to lower the relative affinity for NH<sub>4</sub><sup>+</sup> in each mutualistic partner.

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Altering relative NH<sub>4</sub><sup>+</sup> affinities affects mutualistic partner frequencies. We then examined how relative affinities for NH<sub>4</sub><sup>+</sup> influenced mutualism dynamics by comparing the growth trends of cocultures containing either WT E. coli or E. coli ΔAmtB, paired with either R. palustris ΔAmtB, R. palustris Nx, or R. palustris Nx $\Delta$ AmtB, the latter of which we previously determined to exhibit 3-fold higher NH<sub>4</sub><sup>+</sup>excretion levels than the Nx strain in monoculture (26). For each R. palustris partner, cocultures with E. coli ΔAmtB grew slower than cocultures with WT E. coli (Fig. 4A,B). E. coli ΔAmtB also constituted a lower percentage of the population and achieved lower cell densities compared to WT E. coli when paired with the same R. palustris strain (Fig. 4C). These lower frequencies were consistent with the competitive disadvantage of E. coli  $\triangle$ AmtB for excreted NH<sub>4</sub><sup>+</sup> (Fig. 3). For R. palustris strains lacking AmtB, the effects on population trends varied. Consistent with our previous work, R. palustris NxΔAmtB supported higher WT E. coli percentages and cell densities (Fig. 4C) (26). With high  $NH_4^+$  excretion levels from R. palustris  $Nx\Delta AmtB$ , faster E. coli growth leads to rapid organic acid accumulation, which acidifies the environment, inhibits R. palustris growth, and leaves organic acids unconsumed (Fig. 4D) (26). Surprisingly, although R. palustris ΔAmtB excreted less NH<sub>4</sub><sup>+</sup> than R. palustris Nx in monoculture, R. palustris ΔAmtB supported a higher WT E. coli population in coculture and consumable organic acids accumulated (Fig. 4C, D). These trends resemble those from cocultures with R. palustris NxΔAmtB (Fig. 4C, D), which has a high level of NH<sub>4</sub><sup>+</sup> excretion (SI Appendix Fig. S1D). Unlike Nx strains, which have constitutive nitrogenase activity due to a mutation in the transcriptional activator NifA (30), R. palustris  $\Delta$ AmtB has WT NifA. Thus, R. palustris  $\Delta$ AmtB can likely still regulate nitrogenase expression, and thereby its activity, in response to nitrogen starvation. We hypothesized that in coculture with WT E. coli, R. palustris ΔAmtB might experience heightened nitrogen starvation, as NH<sub>4</sub><sup>+</sup> consumption by WT E. coli would limit NH<sub>4</sub><sup>+</sup> reacquisition by R. palustris  $\Delta$ AmtB (in an *R. palustris*  $\Delta$ AmtB monoculture any lost NH<sub>4</sub><sup>+</sup> would remain available to *R. palustris*). We therefore tested whether coculture conditions stimulated higher nitrogenase activity using an acetylene reduction assay. In agreement with our hypothesis, R. palustris ΔAmtB had increased nitrogenase activity in coculture conditions compared to monocultures, whereas R. palustris Nx, which

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exhibits constitutive nitrogenase activity, showed similar levels in both conditions (SI Appendix Fig. S4). Thus, the relatively high WT E. coli population in coculture with R. palustris  $\Delta$ AmtB is likely due to both the competitive advantage for acquiring  $NH_4^+$  over R. palustris  $\Delta$ AmtB (Fig. 3) and higher  $NH_4^+$  crossfeeding levels due to increased nitrogenase activity. E. coli must have a competitive advantage for NH<sub>4</sub><sup>+</sup> acquisition to avoid mutualism collapse. We were surprised to observe that cocultures of R. palustris  $\Delta$ AmtB paired with E. coli  $\Delta$ AmtB showed little growth when started from a single colony of each species (Fig. 4A), a method that we routinely use to initiate cocultures (26, 31). We reasoned that the higher R. palustris  $\Delta$ AmtB affinity for NH<sub>4</sub><sup>+</sup> relative to E. coli ΔAmtB (Fig. 3) likely led to community collapse as predicted by SyFFoN (Fig. 2). Even though SyFFoN had predicted R. palustris growth when outcompeting E. coli for NH<sub>4</sub><sup>+</sup> (Fig. 2), SyFFoN likely underestimates the time required to achieve these densities, if they would be achieved at all, as SyFFoN does not take into account cell death, which is known to occur when E. coli growth is prevented (31). Consistent with the hypothesis that poor coculture growth was due to a competitive disadvantage of E.  $coli \Delta AmtB$  for  $NH_4^+$ , SyFFoN simulations indicated that starting with a larger E.  $coli \Delta AmtB$  population would increase the probability that any given E. coli cell would acquire NH<sub>4</sub><sup>+</sup> versus R. palustris and thereby overcome the competitive disadvantage of E. coli  $\Delta$ AmtB for NH<sub>4</sub><sup>+</sup> (SI Appendix Fig. S5). Indeed, we observed greater growth of both species when cocultures were inoculated with equal or higher relative densities of E. coli ΔAmtB versus R. palustris ΔAmtB (SI Appendix Fig. S5). The explanation that mutualism collapse was due to a competitive advantage of R. palustris  $\triangle$ AmtB over E. coli  $\triangle$ AmtB for NH<sub>4</sub><sup>+</sup> called into question why cocultures pairing E. coli  $\triangle$ AmtB with either R. palustris Nx or R. palustris NxΔAmtB did not collapse as well (Fig. 4), given that in all of these pairings E. coli ΔAmtB is at competitive disadvantage (Fig. 3). We hypothesized that a relatively high NH<sub>4</sub><sup>+</sup> excretion level by these latter R. palustris strains (SI Appendix Fig. S1D) could compensate for a low E. coli NH<sub>4</sub><sup>+</sup> affinity. To explore this hypothesis we simulated cocultures with the R. palustris affinity for  $NH_4^+$  set high relative to that of E. coli (Rp:Ec = 1000) and varied the R. palustris  $NH_4^+$  excretion level

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(Fig. 5). Indeed, increasing R. palustris NH<sub>4</sub><sup>+</sup> excretion was predicted to overcome a low E. coli affinity for NH<sub>4</sub><sup>+</sup> and support growth of both species (Fig. 5). The only exception was at the highest levels of NH<sub>4</sub><sup>+</sup> excretion, where R. palustris growth was predicted to be inhibited due to rapid E. coli growth and subsequent accumulation of organic acids that acidify the environment (Fig. 5) (26). These simulations suggested that R. palustris Nx and NxΔAmtB supported coculture growth with E. coli ΔAmtB due to higher NH<sub>4</sub><sup>+</sup> excretion levels (SI Appendix Fig. S1D), whereas a combination of low NH<sub>4</sub><sup>+</sup> excretion by R. palustris  $\triangle$ AmtB (SI Appendix Fig. S1D) and a low affinity for NH<sub>4</sub><sup>+</sup> by E. coli  $\triangle$ AmtB led to collapse of the mutualism in this pairing. So far, we had only considered the effect of severe discrepancies in NH<sub>4</sub><sup>+</sup> affinities between the two species (e.g., 1000-fold difference in  $K_m$  values in our simulations) as a mechanism leading to coculture collapse within the time period of a single culturing. However, we wondered if a subtle discrepancy in NH<sub>4</sub> affinities could lead to coculture collapse if given more time. We therefore simulated serial transfers of cocultures with partners having different relative NH<sub>4</sub><sup>+</sup> affinities (Fig. 6A, B). At equivalent NH<sub>4</sub><sup>+</sup> affinities (Fig. 6A), both species were predicted to be maintained over serial transfers. However, when the relative affinities approached a threshold (relative Rp:Ec = 2.75), cell densities of both species were predicted to decrease over serial transfers (Fig. 6B). This decline in coculture growth is due to E. coli being slowly but progressively outcompeted for NH<sub>4</sub><sup>+</sup> by R. palustris. As the difference between the R. palustris and E. coli populations expands, R. palustris cells have a greater chance of acquiring NH<sub>4</sub><sup>+</sup> than the smaller E. coli population, further starving E. coli and simultaneously cutting off R. palustris from its supply of organic acids from E. coli. The above prediction prompted us to investigate if cocultures pairing R. palustris Nx with E. coli ΔAmtB were stable through serial transfers. We focused on cocultures with R. palustris Nx rather than R. palustris NxΔAmtB because R. palustris Nx has AmtB and would therefore be most likely to outcompete E. coli ΔAmtB. Strikingly, after eight serial transfers of cocultures pairing R. palustris Nx with E. coli  $\Delta$ AmtB we observed coculture collapse (Fig. 6C). This observation is in stark contrast to cocultures of R. palustris Nx paired with WT E. coli, which we have serially transferred for over 100 times with no

extinction events (unpublished data). These results indicate that the recipient population must have a competitive advantage for a cross-fed nutrient versus the producer population to avoid mutualism collapse.

#### Discussion

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Here we demonstrate that mutualistic partners can compete for a cross-fed nutrient upon which the mutualistic interaction is based, in this case NH<sub>4</sub><sup>+</sup>. This competition can impact partner frequencies and mutualism stability. Efficient nutrient reacquisition by the producer can render nutrient excretion levels insufficient for cooperative growth, starving the recipient and leading to tragedy of the commons (36). Conversely, recipient-biased competition for a cross-fed nutrient drives cooperative directionality in nutrient exchange and thereby promotes mutualism stability. One implication of these results is that interpartner competition can influence the level of resource privatization. Within microbial interdependencies, partial privatization has primarily been thought to depend on mechanisms used by the producer to retain a portion of a communally-valuable resource (15). Our data indicate that for excreted resources having a transient availability to both mutualists, recipient acquisition mechanisms can also influence the level of producer privatization, as the competition impacts how much of a cross-fed resource will be shared versus re-acquired. In effect, recipient-biased competition avoids tragedy of the commons by enforcing partial privatization of a communally-valuable resource. The importance of the recipient having the upper hand in inter-partner competition likely applies to other synthetic cocultures and natural microbial mutualisms that are based on the cross-feeding of communally-valuable nutrients, including amino acids (37, 38) and vitamin B<sub>12</sub> (6, 11). The same rule could also apply to inter-kingdom and non-microbial examples of cross-feeding (e.g., plants and pollinators, nutrient transfer between plants and bacteria or fungi (39)) and cooperative feeding (e.g., honeyguide bird and human harvesting of bee hives (40), cooperative hunting between grouper fish and moray eels (41)). In such cases, increased privatization of a cross-fed or shared resource, for example through producer-biased competition, could threaten the mutualism upon which both species depend (15, 39, 42).

In our system, AmtB transporters were crucial determinants of inter-partner competition for NH<sub>4</sub><sup>+</sup>. We were intrigued to find that when both species lacked AmtB, R. palustris out-competed E. coli for NH<sub>4</sub> (Fig. 5), enough so to collapse the mutualism within a single culturing (Fig. 3). Whether by maximizing NH<sub>4</sub><sup>+</sup> retention or re-acquisition, R. palustris, and perhaps other N<sub>2</sub>-fixers, might have additional mechanisms aside from AmtB to minimize loss of NH<sub>4</sub><sup>+</sup> as NH<sub>3</sub>. These mechanisms could include a relatively low internal pH to favor NH<sub>4</sub><sup>+</sup> over NH<sub>3</sub>, negatively-charged surface features, or relatively high affinities by NH<sub>4</sub><sup>+</sup>-assimilating enzymes such as glutamine synthetase. There are several reasons why it would be beneficial for N<sub>2</sub>-fixers to minimize NH<sub>4</sub><sup>+</sup>loss. First, N<sub>2</sub> fixation is expensive, both in terms of the enzymes involved (43) and the reaction itself, costing 16 ATP to convert one N<sub>2</sub> into two NH<sub>3</sub> (33). Passive loss of NH<sub>3</sub> would only add to this cost, as more N<sub>2</sub> would have to be fixed to compensate. Second, loss of NH<sub>4</sub><sup>+</sup> could benefit nearby microbes competing against an N<sub>2</sub>-fixer for separate limiting nutrients (14, 44). The possibility that N<sub>2</sub>-fixers could have a superior ability to retain or acquire NH<sub>4</sub><sup>+</sup> independently of AmtB is not farfetched. Bacteria are known to exhibit differential abilities to compete for nutrients. For example, iron acquisition commonly involves iron-binding siderophores, but siderophores can be chemically distinct and thereby differ in their affinity for iron (45). Strategies to utilize siderophores as a shared resource are also numerous, leading to different cooperative or competitive outcomes in microbial communities (45, 46). One must consider that additional mechanisms for acquiring NH<sub>4</sub><sup>+</sup> beyond AmtB might likewise exist. As our results have raised the potential for interpartner competition for cross-fed resources themselves, understanding the physiological mechanisms that confer competitive advantages for nutrient acquisition between species will undoubtedly aid in describing the interplay between competition and cooperation within mutualisms.

### **Materials and Methods**

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**Strains and growth conditions.** Strains, plasmids, and primers are listed in SI Appendix Table S2. All R. palustris strains contained  $\Delta uppE$  and  $\Delta hupS$  mutations to facilitate accurate colony forming unit (CFU) measurements by preventing cell aggregation (47) and to prevent  $H_2$  uptake, respectively. E. coli was

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cultivated on Luria-Burtani (LB) agar and R. palustris on defined mineral (PM) (48) agar with 10 mM succinate. (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> was omitted from PM agar for determining R. palustris CFUs. Monocultures and cocultures were grown in 10-mL of defined M9-derived coculture medium (MDC) (26) in 27-mL anaerobic test tubes. To make the medium anaerobic, MDC was bubbled with N2, then tubes were sealed with rubber stoppers and aluminum crimps, and then autoclaved. After autoclaving, MDC was supplemented with cation solution (1 % v/v; 100 mM MgSO<sub>4</sub> and 10 mM CaCl<sub>2</sub>) and glucose (25 mM), unless indicated otherwise. E. coli monocultures were also supplemented with 15mM NH<sub>4</sub>Cl. All cultures were grown at 30°C laying horizontally under a 60 W incandescent bulb with shaking at 150 rpm. Starter cocultures were inoculated with 200 µL MDC containing a suspension of a single colony of each species. Test cocultures were inoculated using a 1% inoculum from starter cocultures. Serial transfers were also inoculated with a 1% inoculum. Kanamycin and gentamycin were added to a final concentration of 100 μg/ml for R. palustris and 15 μg/ml for E. coli where appropriate. Generation of R. palustris mutants. R. palustris mutants were derived from wild-type CGA009 (49). Generation of strains CGA4004, CGA4005, and CGA4021 was described previously (26). For generation of strain CGA4026 (R. palustris ΔAmtB) the WT nifA gene was amplified using primers JBM1 and JBM2, digested with XbaI and BamHI, and ligated into plasmid pJQ200SK to make pJQnifA16. This suicide vector was then introduced into CGA4021 by conjugation, and sequential selection and screening was performed as described (50) to replace nifA\* with WT nifA. Reintroduction of the WT nifA gene was confirmed by PCR and sequencing. Generation of the E. coli ΔAmtB mutant. P1 transduction (51) was used to introduce ΔamtB::Km from the Keio collection strain JW0441-1 (52) into MG1655. The ΔamtB::Km genotype of kanamycin-resistant colonies was confirmed by PCR and sequencing. **Analytical procedures.** Cell density was assayed by optical density at 660 nm ( $OD_{660}$ ) using a Genesys 20 visible spectrophotometer (Thermo-Fisher, Waltham, MA, USA). Growth curve readings were taken in culture tubes without sampling (i.e., Tube OD<sub>660</sub>). Specific growth rates were determined using readings between 0.1-1.0 OD<sub>660</sub> where there is linear correlation between cell density and OD<sub>660</sub>. Final

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 $OD_{660}$  measurements were taken in cuvettes and samples were diluted into the linear range as necessary. H<sub>2</sub> was quantified using a Shimadzu (Kyoto, Japan) gas chromatograph (GC) with a thermal conductivity detector as described (53). Glucose, organic acids, formate and ethanol were quantified using a Shimadzu high-performance liquid chromatograph (HPLC) as described (54). NH<sub>4</sub><sup>+</sup> was quantified using an indophenol colorimetric assay as described (26). Nitrogenase activity. Nitrogenase activity was measured using an acetylene reduction assay (43). Cells from 10-mL cultures were harvested and resuspended in 10-mL fresh MDC medium in 27-mL sealed tubes pre-flushed with argon gas. Suspensions were incubated in light for 1 h at 30°C to recover. Then, 250 µl of 100% acetylene gas was injected into the headspace to initiate the assay, and ethylene production was measured over time by gas chromatography as described (43). Ethylene levels were normalized to total R. palustris CFUs in the 10-ml volume. NH<sub>4</sub><sup>+</sup> competition assay. Fed-batch cultures were performed in custom anaerobic 75-ml serum vials with side sampling ports. Each vial contained a stir bar and 30-mL of MDC, and was sealed at both ends with rubber stoppers and aluminum crimps. Each vial was supplemented with 25 mM glucose, 1 % v/v cation solution and 20 mM sodium acetate. Starter monocultures of each species were grown to equivalent CFUs/mL in MDC tubes containing limiting nutrients (3 mM sodium acetate for R. palustris and 1.5 mM NH<sub>4</sub>Cl for E. coli), and 1 mL of each species was inoculated into the serum vials. These competition cocultures were incubated at 30°C under a 60 W incandescent bulb with stirring at 200 rpm (Thermo Scientific) for 96 h. Each serum vial was constantly flushed with Ar to maintain anaerobic conditions. NH<sub>4</sub>Cl was fed from a 500 μM NH<sub>4</sub>Cl stock using a peristaltic pump (Watson-Marlow) on an automatic timer (Intermatic DT620) at a rate of 0.33 mL/min once an hour for a final concentration of  $\sim 5 \mu M$  upon each addition. Samples were taken at 0 and 96 h for quantification of CFUs. Mathematical modeling. A Monod model describing bi-directional cross-feeding in batch cultures, called SyFFoN v3 (Syntrophy between Fermenter and Fixer of Nitrogen), was modified from our previous model (31) to allow for competition between E. coli and R. palustris for NH<sub>4</sub><sup>+</sup> as follows: (i) an equation for R. palustris growth rate on NH<sub>4</sub><sup>+</sup> was added to boost the R. palustris growth rate when

- acquiring  $NH_4^+$  and (ii) the ability for R. palustris to consume  $NH_4^+$  was added along with a  $K_m$  of R.
- *palustris* for NH<sub>4</sub><sup>+</sup> (K<sub>AR</sub>). Equations and default parameter values are in the SI Appendix and Table S1.
- 326 SyFFoN v3 runs in R studio and is available for download at: https://github.com/McKinlab/Coculture-
- 327 Mutualism.

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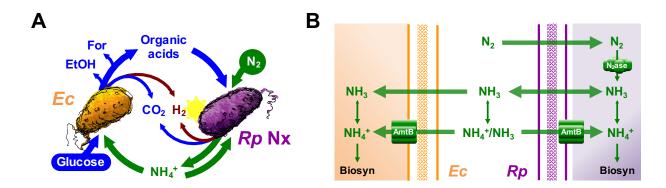
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## Figure Legends



**Fig. 1.** Mechanisms of NH<sub>4</sub><sup>+</sup> transfer within an obligate bacterial mutualism based on cross-feeding of essential nutrients. (A) *Escherichia coli* (*Ec*) anaerobically ferments glucose into organic acids, supplying *Rhodopseudomonas palustris* Nx (*Rp* Nx) with essential carbon. *R. palustris* Nx fixes N<sub>2</sub> gas and excretes NH<sub>4</sub><sup>+</sup>, supplying *E. coli* with essential nitrogen. For, formate; EtOH, ethanol. (B) NH<sub>4</sub><sup>+</sup> can be passively lost from cells as NH<sub>3</sub>. Both species encode high-affinity NH<sub>4</sub><sup>+</sup> transporters, AmtB, that facilitate NH<sub>4</sub><sup>+</sup> uptake. NH<sub>4</sub><sup>+</sup> is the predominant form at neutral pH, as indicated by an enlarged arrow head on double-sided arrows.

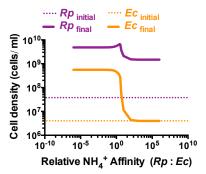
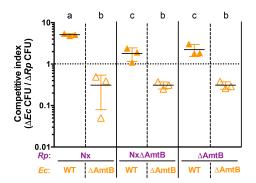


Fig. 2. Simulations suggest that *E. coli* must have a competitive advantage for  $NH_4^+$  acquisition relative to *R. palustris* to support mutualistic growth. Final cell densities (solid lines) of *R. palustris* (Rp, purple) and *E. coli* (Ec, orange) after 300 h in simulated batch cultures for a range of relative  $NH_4^+$  affinities. Initial cell densities are indicated by dotted lines. Relative  $NH_4^+$  affinity values represent the relative E. coli  $K_m$  for  $NH_4^+$  ( $K_A$ ) divided by that of R. palustris ( $K_{AR}$ ).



**Fig. 3. AmtB is important for competitive NH**<sub>4</sub><sup>+</sup> **acquisition.** Competitive indices for *E. coli* after 96 h in NH<sub>4</sub><sup>+</sup>-limited competition assay cocultures. Cocultures were inoculated with *E. coli* and *R. palustris* at equivalent cell densities with excess carbon available for both species (25 mM glucose for *E. coli* and 20 mM sodium acetate for *R. palustris*). NH<sub>4</sub><sup>+</sup> was added to cocultures to a final concentration of 0.5 μM every hour for 96 h. The dotted line indicates a competitive index value of 1, where both species are equally competitive for NH<sub>4</sub><sup>+</sup>. Filled triangles, WT *E. coli*; open triangles *E. coli* ΔAmtB. Error bars indicate SD, n=3. Different letters indicate statistical differences between *E. coli* competitive index values, p < 0.05, determined by one-way ANOVA with Tukey's multiple comparisons post test.

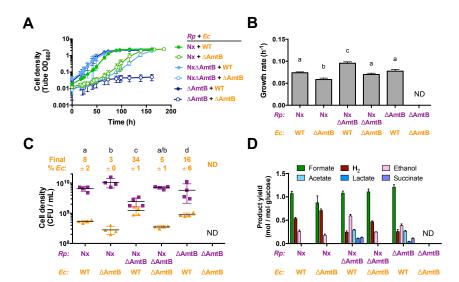


Fig. 4. NH<sub>4</sub><sup>+</sup> transporters influence population and metabolic trends of both partners in coculture.

Growth curves (A), growth rates (B), final cell densities after one culturing (C), and fermentation product yields (D) from cocultures of all combinations of mutants lacking AmtB in each species. Final cell

densities and fermentation product yields were taken after one week, within 24 h into stationary phase.

ND, not determined. Error bars indicate SD, n=4. Different letters indicate statistical differences, p < 0.05, determined by one-way ANOVA with Tukey's multiple comparisons post test.

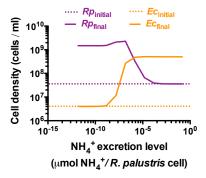


Fig. 5. Higher *R. palustris*  $NH_4^+$  excretion levels are predicted to compensate for a low *E. coli*  $NH_4^+$  affinity. 300 h batch cultures were simulated with a relative *R. palustris*: *E. coli* (*Ec*: *Rp*)  $K_m$  value for  $NH_4^+$  of 0.001 over different *R. palustris*  $NH_4^+$  excretion levels ( $R_A$ ). Final cell densities, solid lines; initial cell densities, dotted lines.

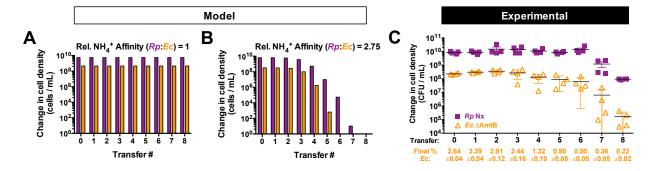


Fig. 6. A low *E. coli*  $NH_4^+$  affinity results in coculture collapse over serial transfers when paired with *R. palustris* Nx. (A,B) 300 h batch cultures were simulated and serial transferred used a 1% inoculum based on the cell density at 300 h for the previous culture. Relative  $NH_4^+$  affinity values represent the relative *E. coli*  $K_m$  for  $NH_4^+$   $(K_A)$  divided by that of *R. palustris*  $(K_{AR})$ . (C) Final cell densities of *R. palustris* Nx and *E. coli*  $\Delta AmtB$  of cocultures grown for one week, less than 24 h into stationary phase. A 1% inoculum was used for each subsequent serial transfer. Error bars indicate SD, n=4. Final *E. coli* cell percentages +/- SD for each transfer are shown.

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                                             Recipient-biased competition for a cross-fed nutrient is required
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                                                                            for coexistence of microbial mutualists
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                                                     Alexandra L. McCully, Breah LaSarre, James B. McKinlay
  4
                                                       Department of Biology, Indiana University, Bloomington
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                                                                                             SI Appendix.
  6
            SyFFoN v3 description.
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            Equations 1-4 were used to describe E. coli and R. palustris growth rates:
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            Eq. 1: E. coli growth rate; \mu_{Ec} = \mu_{EcMAX} \cdot [G/(K_G + G)] \cdot [A/(K_A + A)] \cdot [b_{Ec}/(b_{Ec} + 10^{(f+C)})]
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           Eq. 2: R. palustris growth rate (N<sub>2</sub>); \mu_{Rpn} = \mu_{RpMAX} \cdot [C/(K_C + C)] \cdot [N/(K_N + N)] \cdot [b_{Rp}/(b_{Rp} + 10^{(f+C)})]
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            Eq. 3: R. palustris growth rate (NH<sub>4</sub><sup>+</sup>); \mu_{Rpa}=
                                                                                   \mu_{RpMAX2} \bullet [C/(K_C + C)] \bullet [A/(K_{AR} + A)] \bullet [b_{Rp}/(b_{Rp} + 10^{(f+C)})]
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            Eq 4: Total R. palustris growth rate; \mu_{Rp} = \mu_{Rpn} + \mu_{Rpa}
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            Equations 5-14 were used to describe temporal changes in cell densities and extracellular compounds.
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            Numerical constants in product excretion equations are used to account for molar stoichiometric
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            conversions. Numerical constants used in sigmoidal functions are based on those values that resulted in
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            simulations resembling empirical trends. All R and r parameters are expressed in terms of glucose
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            consumed except for R<sub>A</sub> which is the amount of NH<sub>4</sub><sup>+</sup> produced per R. palustris cell (Table S1).
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           Eq. 5: Glucose; dG/dt = -\mu_{Ec} \cdot Ec/Y_G - \mu_{Ec} \cdot Ec \cdot (R_c + R_f + R_c + R_{CO2}) - Ec \cdot (G/(K_G + G)) \cdot (10/(10 + 1.09^{(1000 \cdot \mu Ec)})) \cdot (b_{Ec}/(b_{Ec} + 10^{(f+C)})) \cdot ((100/(100 + 6^C)) \cdot (100/(100 + 6^C)))
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                                         (r_{\text{C}} + r_{\text{f}} + r_{\text{e}} + r_{\text{CO2}}) + r_{\text{C} \text{ mono}} + r_{\text{f} \text{\_mono}} + r_{\text{e} \text{\_mono}} + r_{\text{CO2} \text{\_mono}} ) 
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            Eq. 6: N<sub>2</sub>; dN/dt = -\mu_{Rp} \cdot Rp \cdot 0.5 \cdot Ra \cdot (1 - (40/(40 + 1.29^{N})) - \mu_{Rp} \cdot Rp/Y_{N}
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            \begin{array}{l} \text{Eq. 7: Consumable organic acids; } \textit{dC/dt} = \text{Ec} \bullet \mu_{Ec} \bullet R_c \bullet 2 + \text{Ec} \bullet 2 \bullet (G/(K_G + G)) \\ \bullet (10/(10 + 1.09^{(1000 \bullet \ \mu Ec)})) \bullet (b_{Ec} \ / (b_{Ec} + 10^{(f + C)})) \bullet (r_C \bullet (100/(100 + 6^C)) + r_{C\_mono}) - (\mu_{Rp} \bullet Rp / Y_C) \\ - 0.25 \bullet Rp \bullet \mu_{Rp} \bullet Rh_{Rp} - 0.25 \bullet Rp \bullet r_{Hp} \bullet (C/(K_C + C)) \bullet (40/(40 + 1.29^N)) \bullet (b_{Rp} / (b_{Rp} + 10^{(f + C)})) \end{array} 
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            Eq. 8: Formate; d\mathbf{f}/d\mathbf{t} = (\text{Ec} \cdot \mu_{\text{Ec}} \cdot R_{\text{f}} \cdot 6) + \text{Ec} \cdot 6 \cdot (G/(K_{\text{G}} + G)) \cdot (10/(10 + 1.09^{(1000 \cdot \mu_{\text{Ec}})}))
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                          • (b_{Ec} / (b_{Ec} + 10^{(f+C)})) • (r_f \cdot (100/(100 + 6^C)) + r_{f,mono})
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            Eq. 9: NH_4^+; dA/dt = Rp \bullet \mu_{Rp} \bullet R_A \bullet (1 - (40/(40 + 1.29^N))) - \mu_{Ec} \bullet Ec/Y_A - (\mu_{Rp} \bullet Rp/Y_{AR}) \bullet (A/(K_{AR} + A))
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            Eq. 10: E. coli; d\mathbf{E}\mathbf{c}/d\mathbf{t} = \mu_{Ec} \cdot \mathbf{E}\mathbf{c}
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            Eq. 11: R. palustris; d\mathbf{R}\mathbf{p}/d\mathbf{t} = \mu_{\mathbf{R}\mathbf{p}} \cdot \mathbf{R}\mathbf{p}
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           Eq. 12: Ethanol; \textit{de/dt} = \text{Ec} \cdot 3 \cdot (\mu_{\text{Ec}} \cdot R_e + (G/(K_G + G)) \cdot (10/(10 + 1.09^{(1000 \cdot \mu_{\text{Ec}})})) \cdot (b_{\text{Ec}}/(b_{\text{Ec}} + 10^{(f + C)})) \cdot (r_e \cdot (100/(100 + 6^C)) + r_{e\_mono}))
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Eq. 13: CO_2; dCO_2/dt = Ec \cdot 6 \cdot (\mu_{Ec} \cdot R_{CO2} + (G/(K_G + G)) \cdot (10/(10 + 1.09^{(1000 \cdot \mu Ec)}))

• (b_{Ec}/(b_{Ec} + 10^{(f+C)})) \cdot (r_{co2} \cdot (100/(100 + 6^C)) + r_{co2\_mono}))

+ Rp \cdot 0.5 \cdot (\mu_{Rp} \cdot Rh_{Rp} + r_{Hp} \cdot (C/(K_C + C)) \cdot (40/(40 + 1.29^N)) \cdot (b_{Rp}/(b_{Rp} + 10^{(f+C)})))

Eq. 14: H_2; dH/dt = Rp \cdot (\mu_{Rp} \cdot R_{HRp} + r_{Hp} \cdot (C/(K_C + C)) \cdot (40/(40 + 1.29^N)) \cdot (b_{Rp}/(b_{Rp} + 10^{(f+C)}))) + Ec \cdot (\mu_{Ec} \cdot R_{HEc} + (G/(K_G + G)) \cdot (10/(10 + 1.09^{(1000 \cdot \mu Ec)})) \cdot (b_{Ec}/(b_{Ec} + 10^{(f+C)})) \cdot (r_H \cdot (100/(100 + 6^C)) + r_{H\_mono}))
```

Where,

μ is the specific growth rate of the indicated species (h<sup>-1</sup>).

 $\mu_{MAX}$  is the maximum specific growth rate of the indicated species (h<sup>-1</sup>).

G, A, C, N, f, e, H and CO2 are the concentrations (mM) of glucose, NH<sub>4</sub><sup>+</sup>, consumable organic acids, N<sub>2</sub>, formate, ethanol, H<sub>2</sub>, and CO<sub>2</sub>, respectively. All gasses are assumed to be fully dissolved. Consumable organic acids are those that *R. palustris* can consume, namely, lactate (3 carbons), acetate (2 carbons), and succinate (4 carbons). All consumable organic acids were simulated to have three carbons for convenience. Only net accumulation of formate, ethanol, CO<sub>2</sub> and H<sub>2</sub> are described in accordance with observed trends.

K is the half saturation constant for the indicated substrate (mM).

Ec and Rp are the cell densities (cells/ml) of E. coli and R. palustris, respectively.

b is the ability of a species to resist the inhibiting effects of acid (mM).

Y is the *E. coli* or *R. palustris* cell yield from the indicated substrate (cells / μmol glucose). Y values were determined in MDC with the indicated substrate as the limiting nutrient.

R is the fraction of glucose converted into the indicated compound per  $E.\ coli$  cell during growth (µmol of glucose  $/\ E.\ coli$  cell), except for  $R_A$ . Values were adjusted to accurately simulate product yields measured in cocultures and in MDC with and without added NH<sub>4</sub>Cl.

 $R_A$  is the ratio of  $NH_4^+$  produced per *R. palustris* cell during growth (µmol / *R. palustris* cell). The default value was based on that which accurately simulated empirical trends.

r is the growth-independent rate of glucose converted into the indicated compound ( $\mu$ mol / cell / h).

Default values are based on those which accurately simulated empirical trends in coculture.

 $r_{mono}$  is the growth-independent rate of glucose converted into the indicated compound by *E. coli* when consumable organic acids accumulate. Default values are based on linear regression of products accumulated over time in nitrogen-free cell suspensions of *E. coli* (4).

# Table S1. Default parameter values used in the model unless stated otherwise

Parameter	Value	Description (Units); Source	
$\mu_{EcMAX}$	0.2800	E. coli max growth rate (h <sup>-1</sup> ); Monoculture	
$\mu_{RpMAX}$	0.0772	R. palustris max growth rate (h <sup>-1</sup> ); Monoculture	
$\mu_{RpMAX2}$	0.0152	Boost on <i>R. palustris</i> growth rate in presence of NH <sub>4</sub> <sup>+</sup> (h <sup>-1</sup> );	
		Monoculture <sup>a</sup>	
G	25	Glucose (mM)	
A C	0.00005	NH <sub>4</sub> <sup>+</sup> (mM); from initial (NH <sub>4</sub> ) <sub>6</sub> Mo <sub>7</sub> O <sub>24</sub> ·4H <sub>2</sub> O concentration	
С	0	Consumable organic acids (those that <i>R. palustris</i> was observed to consume: lactate, acetate, and succinate; mM)	
N	70	N <sub>2</sub> (assumed to be fully dissolved; mM)	
f	0	Formate (mM)	
e	0	Ethanol (mM)	
CO2	0	Carbon dioxide (mM)	
K <sub>G</sub>	0.02	E. coli affinity (Michaelis-Menten constant $(K_m)$ ) for glucose $(mM)$ ;	
KG	0.02	(1)	
K <sub>C</sub>	0.01	R. palustris affinity (K <sub>m</sub> ) for consumable organic acids (mM);	
110	0.01	Assumed (Kin) for consumate organic acids (inivi),	
K <sub>A</sub>	0.01	E. coli affinity for NH <sub>4</sub> (mM); (2)	
K <sub>AR</sub>	0.01	R. palustris affinity for NH <sub>4</sub> <sup>+</sup> (mM); Assumed <sup>b</sup>	
K <sub>N</sub>	6	R. palustris affinity $(K_m)$ for $N_2$ $(mM)$	
Ec	$0.4 \times 10^7$	E. coli cell density (cells / ml)	
Rp	$3.6 \times 10^7$	R. palustris cell density (cells / ml)	
$b_{Ec}$	$\frac{3.6 \times 10^7}{10^{43}}$	Resistance of <i>E. coli</i> to low pH (mM)	
	$10^{32}$	Resistance of <i>E. con</i> to low pH (IIIM)  Resistance of <i>R. palustris</i> to low pH (mM)	
$Y_G$	$8 \times 10^{7}$	Glucose-limited <i>E. coli</i> growth yield (cells / µmol glucose); Glucose-	
1 G		limited E. coli culture	
$Y_A$	1 x 10 <sup>9</sup>	NH <sub>4</sub> <sup>+</sup> -limited <i>E. coli</i> growth yield (cells / μmol NH <sub>4</sub> <sup>+</sup> ); NH <sub>4</sub> <sup>+</sup> -limited <i>E. coli</i> culture	
$Y_{C}$	$2.5 \times 10^8$	Organic acid-limited <i>R. palustris</i> growth yield (cells / µmol organic	
1 C		acid); Acetate-limited <i>R. palustris</i> growth yield (cens / µmor organic acid); Acetate-limited <i>R. palustris</i> culture	
$Y_N$	$5 \times 10^8$	$N_2$ -limited $R$ . palustris growth yield cells / $\mu$ mol $N_2$ ; $N_2$ -limited $R$ .	
		palustris culture	
$R_{C}$	1.9 x 10 <sup>-8</sup>	Fraction of glucose converted to organic acids (µmol glucose / cell)	
$R_{\mathrm{f}}$	8 x 10 <sup>-9</sup>	Fraction of glucose converted to formate (µmol glucose / cell)	
R <sub>e</sub>	4.5 x 10 <sup>-9</sup>	Fraction of glucose converted to ethanol (µmol glucose / cell)	
$R_{CO2}$	5 x 10 <sup>-10</sup>	Fraction of glucose converted to CO <sub>2</sub> (μmol glucose / cell)	
$R_{HRp}$	2 x 10 <sup>-9</sup>	R. palustris H <sub>2</sub> production (μmol H <sub>2</sub> / R. palustris cell)	
$R_{HEc}$	5 x 10 <sup>-9</sup>	E. coli H <sub>2</sub> production (μmol H <sub>2</sub> / E. coli cell)	
$R_A$	0.15 x 10 <sup>-9</sup>	R. palustris NH <sub>4</sub> <sup>+</sup> production (μmol NH <sub>4</sub> <sup>+</sup> / cell)	
$r_{\rm C}$	300 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to consumable organic acids (μmol glucose / cell / h) (3)	
$r_{\mathrm{f}}$	47 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to	
	11	formate (µmol glucose / cell / h) (3)	
$r_{e}$	15 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to	
		ethanol (µmol glucose / cell / h) (3)	
$r_{\rm CO2}$	2 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to CO <sub>2</sub>	
		(μmol glucose / cell / h) (3)	

$r_{\mathrm{H}}$	2 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of H <sub>2</sub> production (μmol H <sub>2</sub> /
		cell / h) (3)
$r_{C\_mono}$	1.2 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to
_		consumable organic acids when consumable organic acids accumulate
		(μmol glucose / cell / h); (4)
r <sub>f mono</sub>	$0.83 \times 10^{-11}$	E. coli specific growth-independent rate of glucose conversion to
_		formate when consumable organic acids accumulate (µmol glucose /
		cell /h); (4)
r <sub>e_mono</sub>	0.5 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to
		ethanol when consumable organic acids accumulate (µmol glucose /
		cell /h); (4)
r <sub>co2 mono</sub>	1.3 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to CO <sub>2</sub>
		when consumable organic acids accumulate (µmol glucose / cell / h);
		(4)
r <sub>H mono</sub>	0.83 x 10 <sup>-11</sup>	E. coli specific growth-independent rate of glucose conversion to H <sub>2</sub>
_		when consumable organic acids accumulate (µmol glucose / cell / h);
		(4)
r <sub>Hp</sub>	27 x 10 <sup>-11</sup>	R. palustris specific growth-independent rate of H <sub>2</sub> production (µmol
r		$H_2$ / cell / h)

<sup>&</sup>lt;sup>a</sup> Increased growth rate in presence of  $NH_4^+$  versus  $N_2$  based on the difference in experimentally determined growth rates in R. palustris monocultures grown with either  $NH_4^+$  or  $N_2$  as a nitrogen source. <sup>b</sup>  $K_{AR}$  was assumed to be equivalent to the published E.  $coli\ K_m\ (2)$  for  $NH_4^+\ (K_A)$ .

# Table S2. Strains, plasmids, and primers used in this study

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Strain or	Description or Sequence (5'-3');	Source or Purpose		
plasmid	<b>Designation</b>			
R. palustris strains				
CGA009	Wild-type strain; spontaneous Cm <sup>R</sup> derivative	(5)		
	of CGA001			
CGA4004	CGA009 ΔhupS Δrpa2750; Parent	(4)		
CGA4005	CGA4004 <i>nifA*</i> ; <u>Nx</u>	(4)		
CGA4021	CGA4005 ΔamtB1 ΔamtB2; NxΔAmtB	(4)		
CGA4026	CGA4004 ΔamtB1 ΔamtB2; <u>Δ</u> AmtB	This study		
E. coli strains				
MG1655	Wild-type K12 strain, <u>WT</u>	(6)		
K-12 JW0441-1	Keio collection ∆amtB::Km	(7)		
MG1655ΔAmtB	MG1655 Δ <i>amtB::Km</i> ; <u>ΔAmtB</u>	This study		
Plasmids				
pJQnifA16	Gm <sup>R</sup> ; WT <i>nifA</i> gene flanked by XbaI/BamHI cloned into pJQ200SK	This study		
Primers				
ALM6f	TTCGTCGCTGAATTGCAACG	amtB upstream flanking region		
		(E. coli)		
ALM6r	TCAGGAAGGGGTGATGCGTA	amtB downstream flanking		
		region (E. coli)		
JBM1	CG <u>TCTAGA</u> CCGGCGCATCGC	<i>nifA16</i> upstream primer; XbaI		
JBM6	GG <u>GGATCC</u> TGGTTCGCAGAGG	<i>nifA16</i> downstream primer;		
		<u>BamHI</u>		

## SI Appendix Figures.

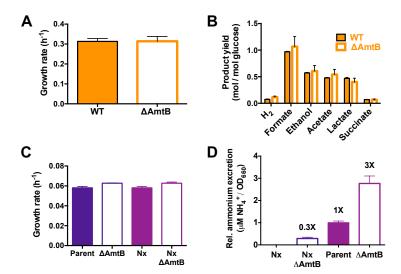


Fig. S1. *E. coli*  $\Delta$ AmtB and *R. palustris*  $\Delta$ AmtB monoculture growth and metabolic trends. (A,B) Growth rates (A) and fermentation product yields (B) from WT *E. coli* (filled) or without (open) *E. coli*  $\Delta$ AmtB monocultures grown in MDC with 25 mM glucose and 15 mM NH<sub>4</sub>Cl. Fermentation profiles were generated from stationary monocultures. Error bars indicate SD, n=3. (C,D) Growth curves (C) and relative NH<sub>4</sub><sup>+</sup> excretion (D) of *R. palustris* monocultures grown in MDC with 3 mM sodium acetate and a 100% N<sub>2</sub> headspace. Error bars indicate SD, n=4.

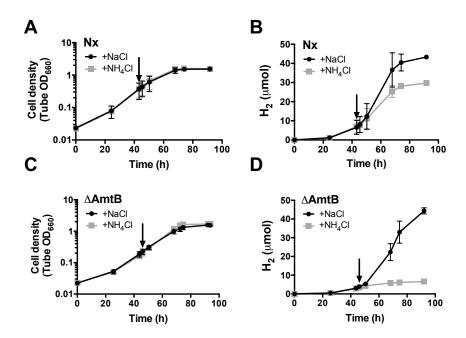


Fig. S2. R. palustris  $\Delta$ AmtB responds to  $NH_4^+$ -induced shutoff of nitrogenase. The effect of either  $NH_4Cl$  or NaCl on growth (A,C) and  $H_2$  production (B,D) in R. palustris Nx or R. palustris  $\Delta$ AmtB monocultures. R. palustris monocultures were grown in MDC with 20 mM sodium acetate and a 100%  $N_2$  headspace until mid-exponential phase and then supplemented with either 15 mM  $NH_4Cl$  or 15 mM NaCl at the time indicated by the arrow.

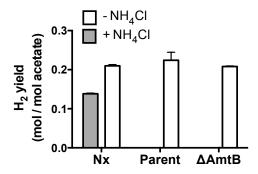
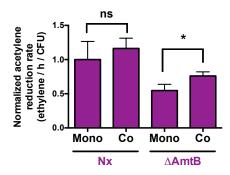


Fig. S3. Unlike *R. palustris* Nx, *R. palustris*  $\Delta$ AmtB does not produce H<sub>2</sub> when grown with NH<sub>4</sub><sup>+</sup>. *R. palustris* monocultures were grown in MDC with 20 mM sodium acetate and a 100% N<sub>2</sub> headspace with (grey) or without (white) 15mM NH<sub>4</sub>Cl. Samples for determining H<sub>2</sub> yields were taken one week after inoculation, within 24 hours into stationary phase. Error bars indicate SD, n=3.



**Fig. S4.** *R. palustris* Δ**AmtB nitrogenase activity increases in coculture.** Normalized nitrogenase activity of *R. palustris* in monoculture (Mono) or coculture (Co) measured by an acetylene reduction assay. Ethylene levels were divided by total *R. palustris* CFUs in the test tube and then normalized to the *R. palustris* Nx monoculture value. Error bars indicate SD, n=4. \*, statistical difference between monoculture and coculture conditions, p < 0.05, determined using multiple two-tailed t-tests; ns, no significant difference.

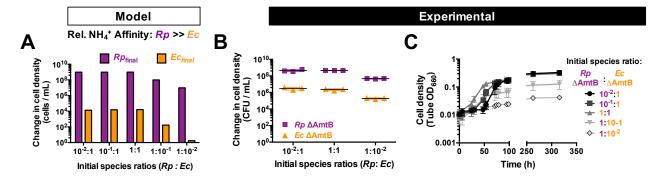


Fig. S5. Higher initial cell densities of *E. coli* ΔAmtB can partially compensate for a low *E. coli* NH<sub>4</sub><sup>+</sup> affinity. Simulations (A) and empirical data (B,C) showing the effect of initial *E. coli* (*Ec*) cell density on population and coculture growth trends when *E. coli* has a lower affinity for NH<sub>4</sub><sup>+</sup> compared to *R. palustris* (Rp). (A) 300 h batch cultures were simulated with a relative R. palustris: E. coli (Rp : Ec) K<sub>m</sub> value for NH<sub>4</sub><sup>+</sup> of 0.001. (B, C) Change in cell densities after one week of growth (B) and growth curves (C) of cocultures inoculated at different species ratios. (A-C) A ratio value of 1 represents 2.7 x 10<sup>6</sup> CFUs/mL, which was experimentally measured from the starting inoculum for both species before diluting to achieve the indicated ratios. Error bars indicate SD, n=3.

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