

Phototrophic lactate utilization by *Rhodopseudomonas palustris* is stimulated by co-utilization with additional substrates

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ABSTRACT

The phototrophic purple nonsulfur bacterium *Rhodopseudomonas palustris* is known for its metabolic versatility and is of interest for various industrial and environmental applications. Despite decades of research on *R. palustris* growth under diverse conditions, patterns of *R. palustris* growth and carbon utilization with mixtures of carbon substrates remain largely unknown. *R. palustris* readily utilizes most short chain organic acids but cannot readily use lactate as a sole carbon source. Here we investigated the influence of mixed-substrate utilization on phototrophic lactate consumption by *R. palustris*. We found that lactate was simultaneously utilized with a variety of other organic acids and glycerol in time frames that were insufficient for *R. palustris* growth on lactate alone. Thus, lactate utilization by *R. palustris* was expedited by its co-utilization with additional substrates. Separately, experiments using carbon pairs that did not contain lactate revealed acetate-mediated inhibition of glycerol utilization in *R. palustris*. This inhibition was specific to the acetate-glycerol pair, as *R. palustris* simultaneously utilized acetate or glycerol when either was paired with succinate or lactate. Overall, our results demonstrate that (i) *R. palustris* commonly employs simultaneous mixed-substrate utilization, (ii) mixed-substrate utilization expands the spectrum of readily utilized organic acids in this species, and (iii) *R. palustris* has the capacity to exert carbon catabolite control in a substrate-specific manner.

IMPORTANCE

Bacterial carbon source utilization is frequently assessed using cultures provided single carbon sources. However, the utilization of carbon mixtures by bacteria (i.e., mixed-substrate utilization) is of both fundamental and practical importance; it is central to bacterial physiology and ecology, and it influences the utility of bacteria as biotechnology. Here we investigated mixed-substrate utilization by the model organism *Rhodopseudomonas palustris*. Using mixtures of organic acids and glycerol, we show that *R. palustris* exhibits an expanded range of usable carbon substrates

when provided in mixtures. Specifically, co-utilization enabled the prompt consumption of lactate, a substrate that is otherwise not readily used by *R. palustris*. Additionally, we found that *R. palustris* utilizes acetate and glycerol sequentially, revealing that this species has the capacity to use some substrates in a preferential order. These results provide insights into *R. palustris* physiology that will aid the use of *R. palustris* for industrial and commercial applications.

INTRODUCTION

Many bacteria in natural environments likely consume multiple carbon sources simultaneously (1, 2). However, bacterial substrate utilization is most often studied using bacteria in isolation with single carbon sources (3). Data on mixed-substrate utilization in diverse bacteria is crucial for both the understanding of nutrient acquisition, metabolism, and community dynamics within microbial ecosystems (1, 4) and the rational application of bacteria as biotechnology (5-8).

When encountering multiple carbon sources (i.e., substrates), a bacterium will utilize the substrates either simultaneously (i.e., co-utilization) or sequentially depending on the identity of the substrates. Sequential utilization typically results in a diauxic growth pattern characterized by two or more exponential phases that are each separated by a lag phase (2, 9); however, sequential utilization can also occur without an intervening lag phase (2, 4, 10), a pattern often referred to as “biphasic growth.” It generally holds true that, during sequential utilization, bacteria preferentially use the carbon source that supports the highest growth rate during the first phase of growth while utilization of the other carbon source(s) is limited until the preferred carbon source is no longer available (2); this process is commonly referred to as carbon catabolite repression (CCR) (11). A classic example of CCR involves the *lac* operon in *Escherichia coli*, which directs the preferential consumption of glucose prior to lactose when cells are provided a mix of the two sugars (11). Although sequential carbon utilization was once thought to predominate among bacteria, particularly at high substrate concentrations (2-4), it

has become clear that simultaneous utilization of carbon substrates is common (1). For example, co-utilization occurs in *Pseudomonas putida* grown with glucose and aromatic compounds (12), in *E. coli* grown with various organic acid pairs (13), and in *Lactococcus brevis* grown with glucose paired with other sugars (14).

The phototrophic purple nonsulfur bacterium *Rhodopseudomonas palustris* is a model organism for investigating metabolic flexibility in response to environmental conditions (15, 16) and is of interest for various commercial applications, including the production of hydrogen gas (17-19) and the biodegradation of aromatic inhibitors in biofuel feedstocks (20). However, there is limited information regarding carbon preference, mixed-substrate utilization, or carbon catabolite control in this species. We recently showed that *R. palustris* utilizes multiple products of *E. coli* mixed-acid fermentation when grown in a synthetic coculture (21). One of these products, lactate, is not readily utilized as a sole carbon source by phototrophically grown *R. palustris* (18). Taken together, these observations raised the question, how does coculturing enable *R. palustris* lactate consumption? Using mixtures of different carbon substrates, here we show that *R. palustris* simultaneously consumes lactate with several other organic acids and glycerol. Importantly, this co-utilization endowed *R. palustris* with the ability to promptly utilize lactate, in spite of the fact that lactate alone did not support growth in the same time frame. Hence, in coculture, the presence of mixed-acid fermentation products enables the co-utilization of lactate by *R. palustris*. Separately, experiments with additional carbon pairings revealed acetate-mediated inhibition of glycerol catabolism, establishing that *R. palustris* can exert hierarchical regulation of carbon utilization.

RESULTS

Acetate and succinate prompt the expedited and simultaneous utilization of lactate.

We previously observed that, when grown anaerobically in a mutualistic coculture with *E. coli*, *R. palustris* simultaneously consumed the acetate, succinate, and lactate excreted as

fermentation products by *E. coli* during the 200 h culturing period (21). Whereas acetate and succinate are both known to be readily utilized by *R. palustris* (19), *R. palustris* requires long-term incubation with lactate before it will utilize lactate as a sole carbon source (18). Indeed, we observed no growth of *R. palustris* with lactate alone within the time frames sufficient for growth on other organic acids (≤ 300 h; see below). As the utilization of a given carbon source can be influenced by the presence of an additional carbon source (1, 12, 13), we hypothesized that the presence of mixed fermentation products facilitated lactate utilization by *R. palustris* in coculture. We reasoned that the most likely mediators of this effect would be succinate and/or acetate given that they were the other carbon substrates consumed by *R. palustris* in coculture. To test if the presence of acetate and succinate could stimulate lactate consumption, we grew *R. palustris* Nx, the strain we had used in coculture with *E. coli*, in monoculture with a mixture of succinate, acetate, and lactate (5 mM each). Cultures grown with this mix exhibited a single exponential phase with a specific growth rate of $0.074 \pm 0.001 \text{ h}^{-1}$ (\pm SD), and growth plateaued within 150 h (**Fig. 1A**). To determine which substrate(s) in the mix had been consumed, we analyzed culture supernatants using high-performance liquid chromatography (HPLC). The data showed that all three compounds had been partially consumed by mid-log-phase and were fully consumed by stationary phase (**Fig. 1B**). Thus, acetate and succinate were sufficient to expedite lactate consumption by *R. palustris*.

To ascertain if lactate consumption was stimulated by both, one, or either succinate or acetate, we also examined growth and carbon utilization in cultures pairing lactate with either acetate or succinate. Cultures reached stationary phase with each of these mixtures within 150-200 h (**Fig. 2A**). For comparison, no growth was detected during the same time period when 10 mM lactate was provided as the sole carbon source (**Fig. 2A**). All cultures consumed all substrates provided (**Fig. 2B**), indicating that lactate consumption was stimulated by either co-substrate. These data also indicated that the co-utilization of lactate did not diminish consumption of succinate or acetate. As the disappearance of substrate from culture

supernatants could be due to either substrate transformation (e.g., degradation) or assimilation, we also compared the growth yields of the cultures. To account for the different amounts of total carbon provided to the cultures (see Fig. 2 legend), we calculated growth yields as the net cell carbon (derived from the final cell optical density less that of the inoculum) per mole carbon consumed. Final biomass yields were all near the theoretical maximum of 1 mole of biomass per mole of carbon consumed (**Fig. 2C**), signifying that all substrates were being assimilated in these cultures. Such high growth yields are consistent with those previously observed for *R. palustris* (16, 19).

To assess the sensitivity of *R. palustris* lactate utilization to the presence of co-substrate, we next examined growth in cultures containing 5mM lactate with varying amounts of acetate (0.5mM – 5mM). Although the growth rates varied slightly with acetate concentration, stimulation of lactate utilization occurred with all acetate concentrations tested (**Fig. 2D**). Thus, lactate utilization can be stimulated by a range of co-substrate concentrations and different lactate:co-substrate ratios. Notably, cells from cultures grown with acetate plus lactate, which presumably contained the necessary enzymes for lactate catabolism, failed to grow when transferred to fresh medium with lactate as the sole carbon source (**Fig. 2E**); this suggests that co-utilization itself was necessary for expedited lactate utilization and that co-substrates are inadequate to prime cell physiology for growth on lactate as the sole carbon source. Based on these data, we conclude that utilization of acetate or succinate is sufficient to stimulate the expedited and simultaneous co-utilization of lactate by *R. palustris*.

Mixed-substrate utilization stimulates lactate consumption in diverse *R. palustris* strains.

The above experiments examined lactate utilization under conditions that mimicked coculture conditions, wherein lactate co-utilization was first observed, in the following two regards. First, we used the engineered 'Nx' strain of *R. palustris*, which harbors a mutation in *nifA* resulting in constitutive N₂ fixation, deletion of *hupS* to prevent H₂ oxidation, and deletion of *hfsE* to prevent

cell aggregation (21, 22). Second, the cultures were grown in a minimal medium (MDC) with N₂ as the sole nitrogen source (21). To assess if the engineered mutations and/or N₂-fixing conditions contributed to the co-utilization of lactate with acetate and succinate, we examined carbon utilization in CGA009, the wild-type parent strain of *R. palustris* Nx, grown with the acetate, succinate, and lactate mixture or with lactate alone in either MDC or in an NH₄⁺-containing minimal medium, PM. The presence of NH₄⁺ in PM represses N₂-fixation in CGA009 (23). Lactate utilization patterns were similar to those in *R. palustris* Nx, regardless of the media: CGA009 consumed all three compounds when provided as a mixture within 150 h and failed to grow with lactate alone in the same time frame (**Fig. 3**). Thus, the observed lactate consumption patterns were not due to either the engineered mutations in the Nx strain or the N₂-fixing conditions.

We also investigated if acetate and succinate stimulated lactate utilization in environmental *R. palustris* strains. Environmental isolates of *R. palustris* have large genetic differences and exhibit unique metabolic characteristics that are thought to aid in nutrient acquisition, anaerobic fermentation, and/or light-harvesting (24-26). Thus, it was conceivable that other *R. palustris* strains behave differently with regard to lactate utilization, either readily using lactate as a sole carbon source or failing to use lactate even in the presence of additional organic acids. However, these potential alternatives were refuted for two environmental isolates, namely, BisB5 and DX-1. When BisB5 and DX-1 were grown with acetate, succinate, and lactate in PM (**Fig. 4A**), all three compounds were consumed within 120 hours (**Fig. 4B**). In contrast, little or no growth was observed with lactate as the sole carbon source within the same time frame (**Fig. 4A**). These results indicate that stimulation of lactate catabolism via mixed-substrate utilization is conserved among diverse *R. palustris* strains.

***R. palustris* Nx lactate utilization is stimulated by diverse carbon co-substrates.** The carbon substrates available to *R. palustris* in natural environments are presumably more diverse

than *E. coli* fermentation products. Therefore, we investigated if lactate utilization by *R. palustris* was stimulated by co-consumption of carbon substrates other than acetate and succinate. Specifically, we grew *R. palustris* with malate, butyrate, and glycerol, as either the sole carbon source or paired with lactate. Glucose was not tested because *R. palustris* cannot consume sugars (27). It warrants mentioning that *R. palustris* cannot grow phototrophically on butyrate alone unless it can dispose of the excess electrons associated with this substrate; CO₂-fixation is perhaps the best-known mechanism by which *R. palustris* will dispose of excess electrons, but N₂ fixation can also fill this role (16). Because the *R. palustris* Nx strain harbors a NifA* mutation resulting in constitutive nitrogenase activity (16), and the MDC medium used in these cultures necessitates N₂ fixation for growth (21), *R. palustris* Nx is readily able to achieve electron balance and grow with butyrate alone. Similar to the results with acetate and succinate, when we grew *R. palustris* with lactate paired with malate, butyrate, or glycerol, we observed that lactate was utilized simultaneously with each of the three substrates (**Fig. 5A-C**). These data demonstrate that lactate utilization can be stimulated by diverse co-substrates.

Differential effects of mixed-substrate utilization on *R. palustris* Nx growth. In working with different substrate mixtures containing lactate, we noticed that co-utilization sometimes resulted in different growth rates compared to the growth rates on single carbon sources. Specific growth rates (μ) during co-utilization could be categorized in comparison to the growth rates on the constituent substrates alone, as follows: (i) the mixed-substrate μ was faster than that on either substrate alone (i.e., enhanced μ); (ii) the mixed-substrate μ approximated that when grown on the individual substrate allowing the fastest growth (i.e., equivalent μ); or (iii) the mixed-substrate μ was between the μ 's on the individual substrates (i.e., intermediate μ). We considered cultures with lactate as the sole carbon source to have a growth rate of 0 h⁻¹, as no growth was observed in these cultures within experimental time frames (\leq 300 h). We observed enhanced μ in cultures pairing lactate with glycerol, equivalent μ in cultures pairing lactate with

succinate or malate, and intermediate μ in cultures pairing lactate with acetate or butyrate (**Fig. 5D**). Akin to growth patterns in other species (13), there was no evident correlation between the effect of mixed-substrate utilization on growth rate and either the metabolic entry point of the co-substrate or the growth rate on the co-substrate alone.

To investigate if the changes in mixed-substrate growth rates were contingent on the co-substrate rather than on lactate itself, we examined growth of *R. palustris* with three substrate pairs that did not contain lactate: succinate with acetate, succinate with glycerol, and acetate with glycerol. When acetate was paired with succinate, the compounds were utilized simultaneously and the growth rate matched that of cultures with acetate alone (equivalent μ) (**Fig. 5D,E**). Similar results were seen in cultures containing succinate paired with glycerol, with growth rates approximating those of succinate, the 'preferred' carbon source (equivalent μ) (**Fig. 5D,F**). However, pairing acetate with glycerol resulted in two distinct exponential growth phases, with the first and second phases having growth rates that approximated those with acetate alone and glycerol alone, respectively (**Fig. 5D,G**). This pattern suggested that acetate and glycerol were being consumed sequentially, rather than simultaneously. HPLC results confirmed that acetate consumption occurred during the first exponential phase whereas glycerol consumption did not occur until acetate had been depleted from the medium (**Fig. 5G,H**). From these data, we conclude that, whereas *R. palustris* can simultaneously consume a wide range of substrates when provided in mixtures of two and three, acetate and glycerol are consumed sequentially by *R. palustris*.

DISCUSSION

Here we revealed that lactate can be readily catabolized by *R. palustris* in the presence of various other organic acids and glycerol (**Figs. 2 and 5**), despite that lactate did not support growth as the sole carbon source in the same time frames. Mixed-substrate-mediated stimulation of lactate consumption occurred in both lab-adapted and environmental WT *R.*

palustris strains that are genetically distinct (**Figs. 3 and 4**). Thus, this phenomenon appears to be broadly conserved despite the high degree of genetic diversity that exists among isolates of this species (24-26).

It is tempting to speculate how co-utilization expedites lactate consumption. The fact that we observed a similar induction effect with diverse substrates that enter central metabolism at different points of both glycolysis/gluconeogenesis and the TCA cycle makes it difficult to predict the underlying mechanism(s). However, we believe several mechanisms can be excluded. First, there are instances where co-substrates enable anaerobic growth by acting as alternative electron acceptors and thereby contributing to cellular redox balance (28-30). The contribution of co-substrates to electron balance during lactate co-utilization is unlikely because: (i) the same pattern of lactate utilization was observed in two conditions that differentially allow N₂ fixation (**Fig. 3**), a process known to satisfy electron balance in *R. palustris* (16); and (ii) lactate utilization was stimulated equivalently by carbon substrates that were more oxidized or less oxidized than lactate (**Table 1**). Second, co-transport is unlikely to be responsible, as co-consumption was not strictly dependent on lactate:co-substrate stoichiometry (**Fig. 2D**) and induction occurred with diverse co-substrates that presumably do not all utilize the same transporter (**Figs. 2 and 5**). Finally, in some instances co-substrates can have an “auxiliary” effect by providing energy during the catabolism of energy-deficit substrates (1, 31). The need for supplemental energy generation is unlikely in the case presented herein, given that *R. palustris* was grown under phototrophic conditions where energy is derived from light. Although outside of the scope of this study, we hope that future work identifies the mechanism(s) by which mixed-substrate utilization expedites lactate consumption by *R. palustris*.

This study was initiated to investigate the potential co-utilization of lactate with other carbon substrates. However, our results also revealed acetate inhibition of glycerol catabolism in *R. palustris* (**Fig. 5G, H**). *R. palustris* is well known for its strict control of nitrogen utilization, wherein the presence of ammonium strictly inhibits expression of the nitrogenase enzyme that

catalyzes N₂ fixation (23, 32). However, we are unaware of any report of CCR in this species. There was no evident lag phase between the two exponential growth phases in *R. palustris* cultures containing acetate paired with glycerol (**Fig. 5G**). This direct transition between exponential phases could indicate that acetate-mediated inhibition of glycerol assimilation in *R. palustris* occurs at the level of protein activity (e.g., transport or catabolic enzyme activity), rather than the level of protein expression (2, 4, 33). However, as brought up by a reviewer, it is also possible that new proteins required for glycerol consumption in the second phase can be synthesized upon acetate depletion in a comparatively short time frame relative to the long *R. palustris* doubling time, such that an intervening lag phase is not observed. Future studies will be needed to determine the mechanism by which acetate represses glycerol consumption. *R. palustris* CGA009 has more than 400 genes predicted to be involved in regulation and signal transduction (27). Among these are genes encoding Crp- and Hpr-like proteins (27, 34). Crp and Hpr homologues regulate diverse biological functions that include CCR in certain species (34, 35). As such, the Hpr- and Crp-like proteins seem logical initial targets for mutagenesis in the endeavor to characterize catabolite control mechanisms in *R. palustris*. Identifying the transporters used for different carbon substrates in *R. palustris* will likely also be important for elucidating such mechanisms. As *R. palustris* encodes more than 300 different transport systems (27), and results from a large-scale study of ABC transporter proteins indicate that sequence-based homology is unreliable for predicting ligand specificity (36), this will not be a trivial task.

Although simultaneous utilization of carbon substrates is most commonly described under nutrient-limited conditions (2-4), examples are accumulating, including for *R. palustris* as shown here, wherein bacteria simultaneously utilize substrates even at high concentrations (1, 13). *R. palustris* simultaneously consumed seven of the eight substrate pairs tested in this study, and published data suggest that this behavior may extend beyond organic acids and glycerol. For example, data from a recent study indicated that *R. palustris* simultaneously

utilizes acetate and various aromatic compounds when grown in corn stover hydrolysate (20), though it was not determined which compounds were being assimilated into biomass. The same study reported simultaneous biological transformation of several aromatic compounds that are not readily utilized as sole carbon sources (15, 20), perhaps indicating that mixed-substrate utilization influences the aromatic utilization spectrum of *R. palustris* as well. It is possible that assessment of bacterial nutritional repertoires using single substrates underestimates the catabolic capabilities of some bacteria. From an ecological perspective, it would not necessarily be surprising if *R. palustris* co-utilizes a large range of carbon sources. Such a strategy could allow *R. palustris* to take full advantage of the diverse carbon sources it encounters within the numerous environments it inhabits (27, 37). It has been proposed that carbon source preference reflects the likelihood of encountering various substrates in the environment (38). Thus, to speculate further, the disparity between lactate utilization in the presence and absence of a co-substrate could indicate that lactate is rarely encountered as the sole carbon source in natural environments. In this case, the inability to readily use lactate as the sole carbon source would not be of consequence to *R. palustris*. Finally, beyond these potential ecological implications, substrate co-utilization, particularly at high substrate concentrations, is preferable for industrial and commercial applications (8, 39). Specifically, such behavior is crucial for developing bioprocesses that utilize cheap, renewable waste materials, such as industrial effluents, lignocellulosic biomass, and food waste, as feedstocks for the production of biofuels and value-added products. We believe the proclivity to co-utilize carbon substrates enhances the potential biotechnological value of *R. palustris*.

MATERIALS AND METHODS

Chemicals, strains, and growth conditions. The *R. palustris* strains used in this study are listed in Table 2. *R. palustris* was routinely cultivated on defined mineral (PM) (40) agar

supplemented with 10 mM succinate. All cultures were grown in 27-mL anaerobic test tubes containing 10 mL of either defined M9-derived coculture medium (MDC) (21) or PM medium. MDC or PM were bubbled with 100% N₂ or Ar, respectively, and tubes were sealed with rubber stoppers and aluminum crimps prior to autoclaving.

For starter cultures, single colonies were used to inoculate MDC with limiting (3 mM) acetate. For experimental cultures, 100 µL aliquots of replicate stationary-phase starter cultures were used to inoculate MDC or PM supplemented with either 10 mM of a single carbon substrate or 5 mM each of multiple carbon substrates, unless indicated otherwise in figure legends. Carbon sources were added to desired final concentrations from 1M stock solutions of glycerol and sodium salts of L-lactate, acetate, succinate, L-malate, and butyrate. All cultures were incubated horizontally at 30°C under a 43 W A19 halogen bulb (750 lumens) with shaking at 150 rpm. At least three independent biological replicates were performed for each culture condition.

Analytical procedures. *R. palustris* growth was monitored via optical density at 660 nm (OD₆₆₀) using a Genesys 20 spectrophotometer (Thermo-Fisher, Waltham, MA, USA). Growth readings were measured in culture tubes without sampling. Specific growth rates were calculated using OD₆₆₀ values between 0.1—1.0 where cell density and OD₆₆₀ are linearly correlated. Final cell densities were measured in cuvettes with samples diluted as needed to achieve an OD₆₆₀ within the linear range. Organic acids and glycerol were quantified using a Shimadzu high-performance liquid chromatograph, as previously described (41).

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REFERENCES

1. Egli T. 1995. The Ecological and Physiological Significance of the Growth of Heterotrophic Microorganisms with Mixtures of Substrates, p 305-386. *In* Jones JG (ed), *Advances in Microbial Ecology*. Springer US, Boston, MA.
2. Harder W, Dijkhuizen L. 1982. Strategies of mixed substrate utilization in microorganisms. *Philos Trans R Soc Lond B Biol Sci* 297:459-80.
3. Gottschal JC. 1992. Principles of enrichment, isolation, cultivation, and preservation of bacteria., p 149–195. *In* Balows A, Trüper, H.G. , Dworkin, M. , Harder, W. , and Schleifer, K.H. (ed), *The Prokaryotes: a handbook on the biology of bacteria : ecophysiology, isolation, identification, applications*. Springer-Verlag, New York.
4. Yoon H, Klinzing G, Blanch HW. 1977. Competition for mixed substrates by microbial populations. *Biotechnol Bioeng* 19:1193-210.
5. Chen GQ. 2012. New challenges and opportunities for industrial biotechnology. *Microb Cell Fact* 11:111.
6. Nakashima N, Tamura T. 2012. A new carbon catabolite repression mutation of *Escherichia coli*, *mlc* *, and its use for producing isobutanol. *J Biosci Bioeng* 114:38-44.
7. Gronenberg LS, Marcheschi RJ, Liao JC. 2013. Next generation biofuel engineering in prokaryotes. *Curr Opin Chem Biol* 17:462-71.
8. Wu Y, Shen X, Yuan Q, Yan Y. 2016. Metabolic Engineering Strategies for Co-Utilization of Carbon Sources in Microbes. *Bioengineering (Basel)* 3.
9. Monod J. 1949. The Growth of Bacterial Cultures. *Annual Review of Microbiology* 3:371-394.

- 364 10. Ghosh S, Pohland FG, Gates WE. 1972. Phasic utilization of substrates by aerobic
365 cultures. *J Water Pollut Control Fed* 44:376-400.
- 366 11. Gorke B, Stulke J. 2008. Carbon catabolite repression in bacteria: many ways to make
367 the most out of nutrients. *Nat Rev Microbiol* 6:613-24.
- 368 12. Reber HH, Kaiser P. 1981. Regulation of the Utilization of Glucose and Aromatic
369 Substrates in 4 Strains of *Pseudomonas-Putida*. *Archives of Microbiology* 130:243-247.
- 370 13. Narang A, Konopka A, Ramkrishna D. 1997. New patterns of mixed-substrate utilization
371 during batch growth of *Escherichia coli* K12. *Biotechnol Bioeng* 55:747-57.
- 372 14. Kim JH, Shoemaker SP, Mills DA. 2009. Relaxed control of sugar utilization in
373 *Lactobacillus brevis*. *Microbiology* 155:1351-9.
- 374 15. Harwood CS, Gibson J. 1988. Anaerobic and aerobic metabolism of diverse aromatic
375 compounds by the photosynthetic bacterium *Rhodopseudomonas palustris*. *Appl*
376 *Environ Microbiol* 54:712-7.
- 377 16. McKinlay JB, Harwood CS. 2010. Carbon dioxide fixation as a central redox cofactor
378 recycling mechanism in bacteria. *Proc Natl Acad Sci U S A* 107:11669-75.
- 379 17. Huang JJ, Heiniger EK, McKinlay JB, Harwood CS. 2010. Production of hydrogen gas
380 from light and the inorganic electron donor thiosulfate by *Rhodopseudomonas palustris*.
381 *Appl Environ Microbiol* 76:7717-22.
- 382 18. Adessi A, McKinlay JB, Harwood CS, De Philippis R. 2012. A *Rhodopseudomonas*
383 *palustris* nifA* mutant produces H₂ from NH₄⁺-containing vegetable wastes.
384 *International Journal of Hydrogen Energy* 37:15893-15900.
- 385 19. McKinlay JB, Harwood CS. 2011. Calvin cycle flux, pathway constraints, and substrate
386 oxidation state together determine the H₂ biofuel yield in photoheterotrophic bacteria.
387 *MBio* 2.
- 388 20. Austin S, Kontur WS, Ulbrich A, Oshlag JZ, Zhang W, Higbee A, Zhang Y, Coon JJ,
389 Hodge DB, Donohue TJ, Noguera DR. 2015. Metabolism of Multiple Aromatic

Compounds in Corn Stover Hydrolysate by *Rhodopseudomonas palustris*. *Environ Sci Technol* 49:8914-22.

21. LaSarre B, McCully AL, Lennon JT, McKinlay JB. 2017. Microbial mutualism dynamics governed by dose-dependent toxicity of cross-fed nutrients. *ISME J* 11:337-348.
22. Fritts RK, LaSarre B, Stoner AM, Posto AL, McKinlay JB. 2017. A Rhizobiales-Specific Unipolar Polysaccharide Adhesin Contributes to *Rhodopseudomonas palustris* Biofilm Formation across Diverse Photoheterotrophic Conditions. *Appl Environ Microbiol* 83.
23. Rey FE, Heiniger EK, Harwood CS. 2007. Redirection of metabolism for biological hydrogen production. *Appl Environ Microbiol* 73:1665-71.
24. Oda Y, Larimer FW, Chain PS, Malfatti S, Shin MV, Vergez LM, Hauser L, Land ML, Braatsch S, Beatty JT, Pelletier DA, Schaefer AL, Harwood CS. 2008. Multiple genome sequences reveal adaptations of a phototrophic bacterium to sediment microenvironments. *Proc Natl Acad Sci U S A* 105:18543-8.
25. Oda Y, Wanders W, Huisman LA, Meijer WG, Gottschal JC, Forney LJ. 2002. Genotypic and phenotypic diversity within species of purple nonsulfur bacteria isolated from aquatic sediments. *Appl Environ Microbiol* 68:3467-77.
26. Oda Y, Star B, Huisman LA, Gottschal JC, Forney LJ. 2003. Biogeography of the purple nonsulfur bacterium *Rhodopseudomonas palustris*. *Appl Environ Microbiol* 69:5186-91.
27. Larimer FW, Chain P, Hauser L, Lamerdin J, Malfatti S, Do L, Land ML, Pelletier DA, Beatty JT, Lang AS, Tabita FR, Gibson JL, Hanson TE, Bobst C, Torres JL, Peres C, Harrison FH, Gibson J, Harwood CS. 2004. Complete genome sequence of the metabolically versatile photosynthetic bacterium *Rhodopseudomonas palustris*. *Nat Biotechnol* 22:55-61.
28. De Meur Q, Deutschbauer A, Koch M, Wattiez R, Leroy B. 2018. Genetic Plasticity and Ethylmalonyl Coenzyme A Pathway during Acetate Assimilation in *Rhodospirillum rubrum* S1H under Photoheterotrophic Conditions. *Appl Environ Microbiol* 84.

29. Letoffe S, Chalabaev S, Dugay J, Stressmann F, Audrain B, Portais JC, Letisse F, Ghigo JM. 2017. Biofilm microenvironment induces a widespread adaptive amino-acid fermentation pathway conferring strong fitness advantage in *Escherichia coli*. *PLoS Genet* 13:e1006800.
30. Guest JR. 1979. Anaerobic growth of *Escherichia coli* K12 with fumarate as terminal electron acceptor. Genetic studies with menaquinone and fluoroacetate-resistant mutants. *J Gen Microbiol* 115:259-71.
31. Babel W, Muller RH. 1985. Mixed Substrate Utilization in Microorganisms - Biochemical Aspects and Energetics. *Journal of General Microbiology* 131:39-45.
32. Zumft WG, Castillo F. 1978. Regulatory Properties of Nitrogenase from *Rhodopseudomonas-Palustris*. *Archives of Microbiology* 117:53-60.
33. Kompala DS, Ramkrishna D, Jansen NB, Tsao GT. 1986. Investigation of bacterial growth on mixed substrates: Experimental evaluation of cybernetic models. *Biotechnology and Bioengineering* 28:1044-1055.
34. Pinedo CA, Gage DJ. 2009. HPrK regulates succinate-mediated catabolite repression in the gram-negative symbiont *Sinorhizobium meliloti*. *J Bacteriol* 191:298-309.
35. Soberon-Chavez G, Alcaraz LD, Morales E, Ponce-Soto GY, Servin-Gonzalez L. 2017. The Transcriptional Regulators of the CRP Family Regulate Different Essential Bacterial Functions and Can Be Inherited Vertically and Horizontally. *Front Microbiol* 8:959.
36. Giuliani SE, Frank AM, Corgliano DM, Seifert C, Hauser L, Collart FR. 2011. Environment sensing and response mediated by ABC transporters. *BMC Genomics* 12 Suppl 1:S8.
37. Madigan MT, Jung DO. 2009. An Overview of Purple Bacteria: Systematics, Physiology, and Habitats, p 1-15. *In* Hunter CN, Daldal F, Thurnauer MC, Beatty JT (ed), *The Purple Phototrophic Bacteria*. Springer Netherlands, Dordrecht.

38. Lahme S, Trautwein K, Strijkstra A, Dorries M, Wohlbrand L, Rabus R. 2014. Benzoate mediates the simultaneous repression of anaerobic 4-methylbenzoate and succinate utilization in *Magnetospirillum* sp. strain pMbN1. *BMC Microbiol* 14:269.
39. Vinuselvi P, Kim MK, Lee SK, Ghim CM. 2012. Rewiring carbon catabolite repression for microbial cell factory. *BMB Rep* 45:59-70.
40. Kim MK, Harwood CS. 1991. Regulation of Benzoate-CoA Ligase in *Rhodopseudomonas-Palustris*. *Fems Microbiology Letters* 83:199-203.
41. McKinlay JB, Zeikus JG, Vieille C. 2005. Insights into *Actinobacillus succinogenes* fermentative metabolism in a chemically defined growth medium. *Appl Environ Microbiol* 71:6651-6.
42. Xing D, Zuo Y, Cheng S, Regan JM, Logan BE. 2008. Electricity generation by *Rhodopseudomonas palustris* DX-1. *Environ Sci Technol* 42:4146-51.
43. McKinlay JB, Oda Y, Ruhl M, Posto AL, Sauer U, Harwood CS. 2014. Non-growing *Rhodopseudomonas palustris* increases the hydrogen gas yield from acetate by shifting from the glyoxylate shunt to the tricarboxylic acid cycle. *J Biol Chem* 289:1960-70.
44. Carlozzi P, Sacchi A. 2001. Biomass production and studies on *Rhodopseudomonas palustris* grown in an outdoor, temperature controlled, underwater tubular photobioreactor. *J Biotechnol* 88:239-49.

TABLES

Table 1. Oxidation states of tested *R. palustris* growth substrates

Substrate	Formula	Oxidation state ^a
Malate	C ₄ H ₆ O ₅	+1
Succinate	C ₄ H ₆ O ₄	+0.5
Lactate	C ₃ H ₆ O ₃	0

Acetate	C ₂ H ₃ O ₂	0
Glycerol	C ₃ H ₈ O ₃	-0.67
Butyrate	C ₄ H ₈ O ₂	-1

^aOxidation states were calculated as previously described (19)

Table 2. *R. palustris* strains used in this study.

Strain	Description or Sequence (5'-3'); Paper designation	Reference
CGA009	Wild-type strain; spontaneous Cm ^R derivative of CGA001	(40)
CGA4005	CGA009 <i>nifA</i> * Δ <i>hupS</i> Δ <i>rpa2750</i> ; Nx	(21)
BisB5	Environmental isolate	(24)
DX-1	Environmental isolate	(42)

FIGURE LEGENDS

Fig. 1. *R. palustris* Nx simultaneously utilizes succinate, acetate, and lactate when provided a mix of the three substrates. (A) Representative growth curve of *R. palustris* Nx grow in MDC with 5 mM each succinate, acetate, and lactate (45 mM carbon total). Similar trends were observed for three other biological replicates (B) Amount (%) of succinate (circles), acetate (triangles), and lactate (squares), remaining in culture supernatants at the indicated cell densities. Each of the four shades of gray indicates an independent biological replicate.

Fig. 2. Both succinate and acetate individually stimulate expedited co-utilization of lactate by *R. palustris* Nx. (A) Representative growth curves of *R. palustris* Nx in MDC with 10 mM individual substrates (acetate [ace; 20 mM carbon], lactate [lac; 30 mM carbon], or succinate [succ; 40 mM carbon]) or 5 mM each of paired substrates (ace/lac [25 mM carbon total]; succ/lac [35 mM carbon total]). (B, C) Amount (%) of each substrate consumed at stationary phase (B) and growth yields (C) of cultures provided 10 mM acetate (20 mM carbon),

5 mM each acetate and lactate (25 mM carbon total), or 5 mM each succinate, acetate, and lactate (45 mM carbon total). Error bars, SD; $n \geq 3$. (C) Growth yields were derived from the change in OD₆₆₀ divided by the change in substrate carbon concentration between the initial and final time point. OD₆₆₀ values were converted into molar carbon using conversion factors of 625 mg dry cell weight / L / OD₆₆₀ (43) and the molecular weight for biomass of 22.426 g/mol based on the elemental composition of *R. palustris* 42OL, CH_{1.8}N_{0.18}O_{0.38} (44). The theoretical maximum growth yield is 1 mole biomass C per mole C consumed. (D) Representative growth curves of *R. palustris* Nx in MDC with 5 mM acetate alone, 5 mM lactate alone, or 5 mM lactate supplemented with the indicated concentrations of acetate. (A, D) Similar trends were observed for three other biological replicates for each condition. (E) Growth curve of *R. palustris* Nx that was subcultured from stationary-phase ace/lac cultures (orange diamonds in panel A) into MDC with 10 mM lactate alone. Error bars, SD; $n=4$.

Fig. 3. Stimulation of lactate consumption via co-utilization of acetate and succinate also occurs in wild-type *R. palustris* CGA009 and is independent of N₂ fixation. (A)

Representative growth curves of *R. palustris* CGA009 in MDC or PM with either 5 mM each succinate, acetate, and lactate (ace/succ/lac; 45 mM carbon total) or 10 mM lactate alone (30 mM carbon). Similar trends were observed for three other biological replicates in each condition. (B) Amount (%) of each substrate consumed at stationary phase in cultures of *R. palustris* CGA009 in MDC or PM with 5 mM each succinate, acetate, and lactate (45 mM carbon total). Error bars, SD; $n=4$.

Fig. 4. Stimulation of lactate consumption via co-utilization of acetate and succinate also occurs in environmental *R. palustris* isolates. (A)

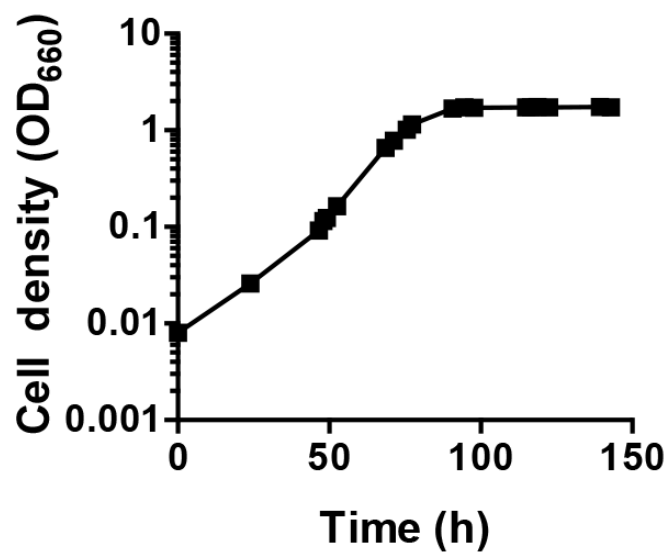
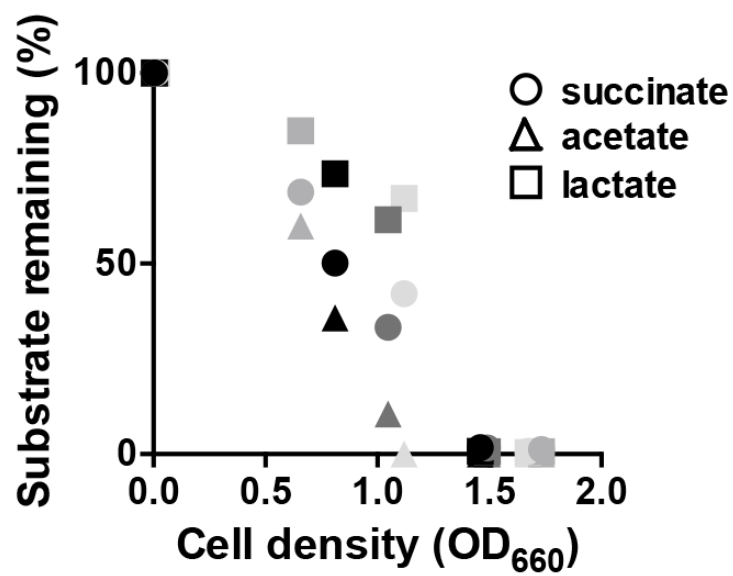
Representative growth curves for *R. palustris* strains BisB5 or DX-1 in PM with either 5 mM each succinate, acetate, and lactate (45 mM carbon total) or 10 mM lactate alone (30 mM carbon). Similar trends were observed for

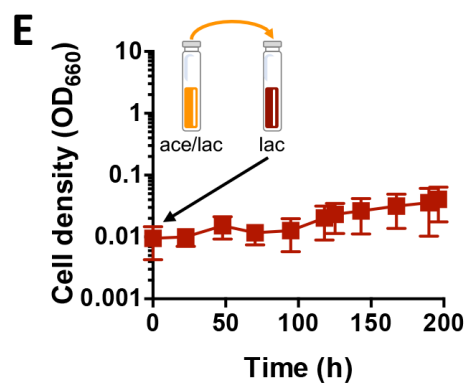
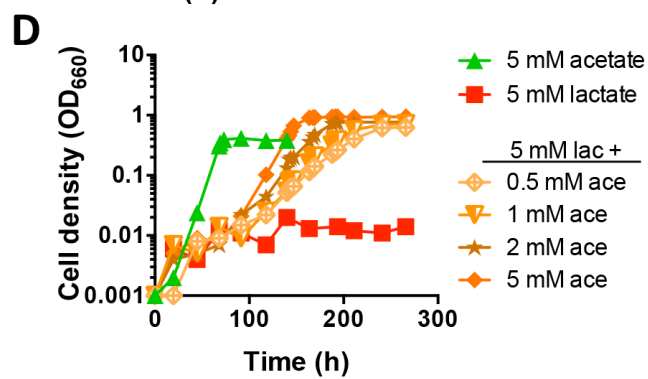
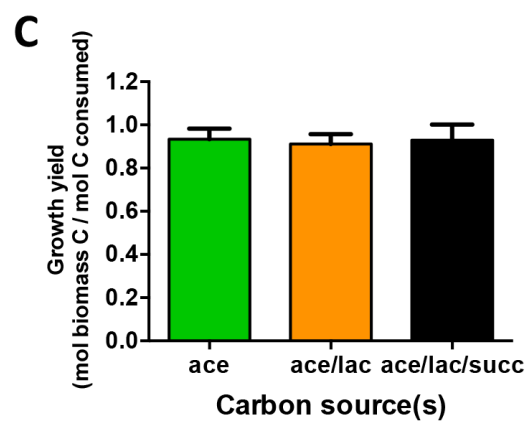
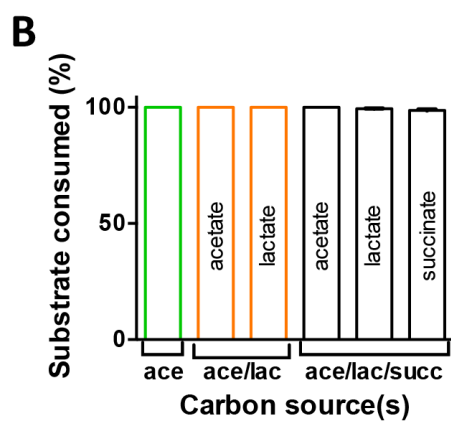
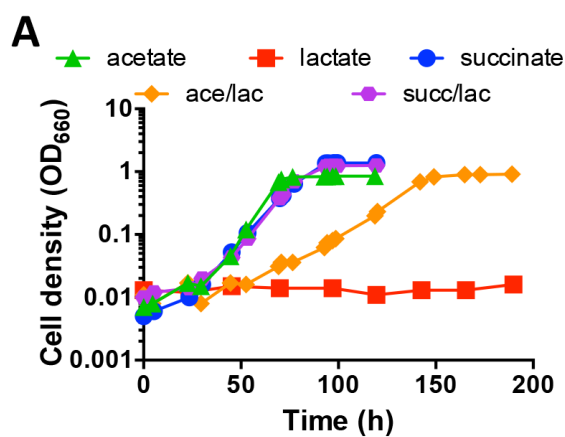
three other biological replicates for each strain in each condition. **(B)** Amount (%) of each substrate consumed at stationary phase in cultures of *R. palustris* BisB5 or DX-1 in PM with 5 mM each succinate, acetate, and lactate (45 mM carbon total). Error bars, SD; n=4.

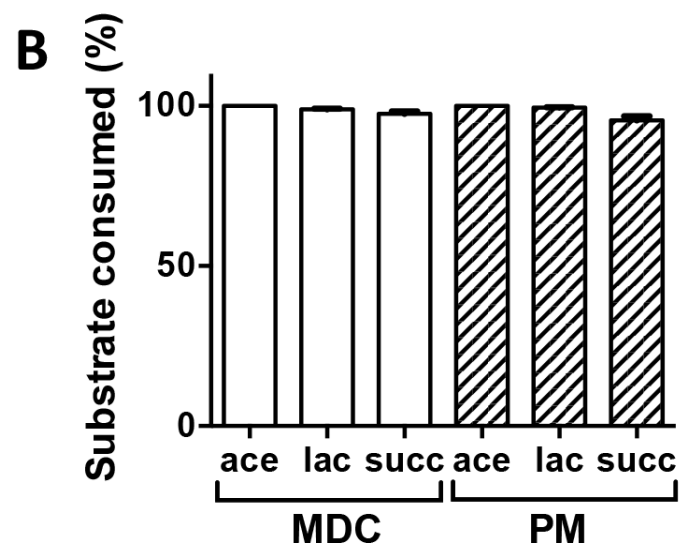
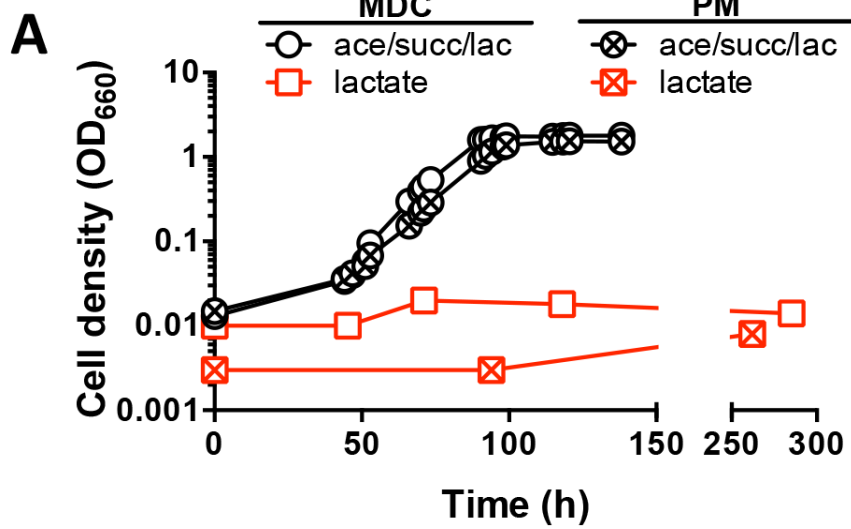
Fig. 5. *R. palustris* co-utilizes many, but not all, carbon substrate pairs. (A-C)

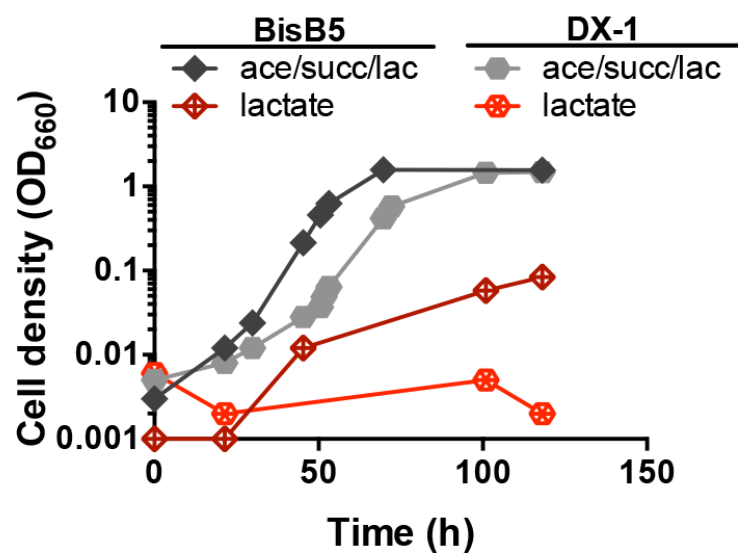
Representative growth curves and amount (%) of substrates remaining at early-log phase (filled symbols) for *R. palustris* Nx in MDC with 5 mM each of paired substrates, as follows: malate + lactate (mal/lac; 35 mM carbon total) **(A)**, butyrate + lactate (buty/lac; 35 mM carbon total) **(B)**, and glycerol + lactate (gly/lac; 30 mM carbon total) **(C)**. Representative growth curves (open symbols) for *R. palustris* Nx in MDC with 10 mM non-lactate substrates alone (malate [40 mM carbon], butyrate [40 mM carbon], glycerol [30 mM carbon]) are included for comparison. Similar trends were observed for two or more additional biological replicates for each condition. **(D)** Specific growth rates of *R. palustris* Nx in MDC with indicated carbon substrates. NG, no growth. Error bars, SD; n≥3. Different letters indicate statistically significant differences between groups ($P < 0.05$; one-way ANOVA with Tukey's multiple-comparison test). **(E-G)**

Representative growth curves and amount (%) of substrates remaining at log phase and stationary phase for *R. palustris* Nx in MDC with 5 mM each of the following paired substrates: succinate + acetate (succ/ace; 30 mM carbon total) **(E)**, succinate + glycerol (succ/gly; 35 mM carbon total) **(F)**, and acetate + glycerol (ace/gly; 25 mM carbon total) **(G)**. Similar trends were observed for two or more additional biological replicates. **(G)** The numbered brackets indicate the two exponential growth phases (see **D**). **(H)** Amount (%) of acetate and glycerol remaining in supernatants of ace/gly cultures at indicated cell densities. Each of the four shades of green (acetate) or purple (glycerol) indicates an independent biological replicate.

A**B**





A**B**