

Reviews and syntheses: Heterotrophic fixation of inorganic carbon – significant but invisible flux in environmental carbon cycling

Alexander Braun¹, Marina Spona-Friedl¹, Maria Avramov¹, Martin Elsner^{1,2}, Federico Baltar³, Thomas Reinthaler³, Gerhard J. Herndl^{3,4} & Christian Griebler^{1,3*}

¹ Helmholtz Zentrum München, Institute of Groundwater Ecology, Ingolstaedter Landstrasse 1, D-85764 Neuherberg, Germany

² Technical University of Munich, Department of Analytical Chemistry and Water Chemistry, Munich, Germany

³ University of Vienna, Department of Functional and Evolutionary Ecology, Althanstrasse 14, 1090 Vienna, Austria

⁴ Department of Marine Microbiology and Biogeochemistry, Royal Netherlands Institute for Sea Research, Utrecht University, PO Box 59, 1790 AB Den Burg, The Netherlands

* Author for correspondence: christian.griebler@univie.ac.at

Abstract

Heterotrophic CO₂ fixation is a significant, yet underappreciated CO₂ flux in environmental carbon cycling. In contrast to photosynthesis and chemolithoautotrophy – the main recognized autotrophic CO₂ fixation pathways – the importance of heterotrophic CO₂ fixation remains enigmatic. All heterotrophs – from microorganisms to humans – take up CO₂ and incorporate it into their biomass. Depending on the availability and quality of growth substrates, and drivers such as the CO₂ partial pressure, heterotrophic CO₂ fixation contributes at least 1-5% and in the case of methanotrophs up to 50% of the carbon biomass. Assuming a standing stock of global heterotrophic biomass of 47-85 Pg C, we roughly estimate that up to 5 Pg C might be derived from heterotrophic CO₂ fixation and up to 12 Pg C yr⁻¹ originating from heterotrophic CO₂ fixation are funneled into the global annual heterotrophic production of 34-245 Pg C yr⁻¹. These first estimates on the importance of heterotrophic fixation of inorganic carbon indicate that this pathway should be incorporated in present and future carbon cycling budgets.

Key words: CO₂ fixation, heterotrophs, anaplerosis, carbon cycling

1. Introduction

Fixation of CO₂ is a fundamental biosynthetic process in nature (Beer et al. 2010, Berg et al. 2007) providing the main source of metabolic energy on Earth (Giovannoni and Stirling 2005). At the same time, it acts as a sink for atmospheric CO₂, the most important greenhouse gas, which is responsible for more than 60% of the 'enhanced greenhouse effect' resulting in global warming (Beer et al. 2010, Berg 2011, Houghton 2007, Le Quéré et al. 2016).

While photosynthesis and chemosynthesis are the most important processes of carbon fixation, non-autotrophic carbon fixation, i.e., the carbon fixation mediated by heterotrophic organisms might also be relevant albeit uncommonly quantified. While heterotrophs are, per definition, organisms that respire organic compounds to gain energy and build up biomass, CO₂ fixation plays also an essential role in heterotrophic carbon metabolism. The diversity of carboxylating enzymes in nature reaches far beyond autotrophy and virtually all heterotrophs harbor numerous enzymes fixing dissolved inorganic carbon. Even though the first carboxylase in heterotrophs was discovered already more than 80 years ago (Wood and Werkman 1936), the role of heterotrophs in carbon cycling has so far largely focused on the oxidation of organic substrates using oxygen or alternative electron acceptors (e.g. nitrate, ferric iron, sulfate) and the production of CO₂. Similar to the CO₂ fixation by autotrophs, "heterotrophic CO₂ fixation" might, however, constitute a significant carbon flux in specific habitats. The relevance of this process has hardly been quantified due to the lack of reliable estimates of heterotrophic CO₂ fixation for most organisms and habitats, and the presumption that CO₂ fixation in natural environments is restricted to autotrophic organisms.

To fill this gap, we review the current knowledge on (i) the role of heterotrophic CO₂ fixation for cellular metabolism, (ii) respiration and non-autotrophic CO₂ fixation, (iii) CO₂ fixation in habitats dominated by heterotrophs, and provide (iv) quantitative estimates of heterotrophic CO₂ fixation in different environments.

2. Role of heterotrophic CO₂ fixation for cellular metabolism

The non-autotrophic uptake of inorganic carbon has been reported for a wide range of organisms from prokaryotes and fungi to vertebrates (Woods & Werkman 1938, Kleiber et al. 1952, Cochrane 1958, Hartman et al. 1972, Perez & Matin 1982, Schinner et al. 1982, Parkinson et al. 1990, Roslev et al. 2004, Hesselsoe et al. 2005, Feisthauer et al. 2008, Spona-Friedl et al. 2020) and plants (Melzer and O'leary 1987). Currently, more than twenty carboxylases are known forming an integral part of the central and peripheral metabolic pathways of heterotrophic metabolism (Fig. 1), e.g., in gluconeogenesis, the synthesis of fatty acids, amino acids, vitamins and nucleotides, the assimilation of leucine, and in anaplerosis (Evans and Slotin 1940, Krebs 1941, Wood and Werkman 1941, Werkman and

Wood 1942, Kornberg and Krebs 1957, Wood and Stjernholm 1962, Kornberg 1965, Scrutton 1971, Hartman et al. 1973, Dijkhuizen and Harder 1985, Parkinson et al. 1991, Attwood 1995, Han et al 2000, Sauer and Eikmanns 2005, Erb et al. 2009, Schink 2009, Erb 2011, Bar-Even et al. 2012). Carboxylation in heterotrophs not just compensates for the dependence on organic matter, rather CO₂ fulfills the role of a “co-substrate” providing an effective and simple way to extend an existing organic carbon substrate by a single C1 unit as part of the secondary production (Erb 2011).

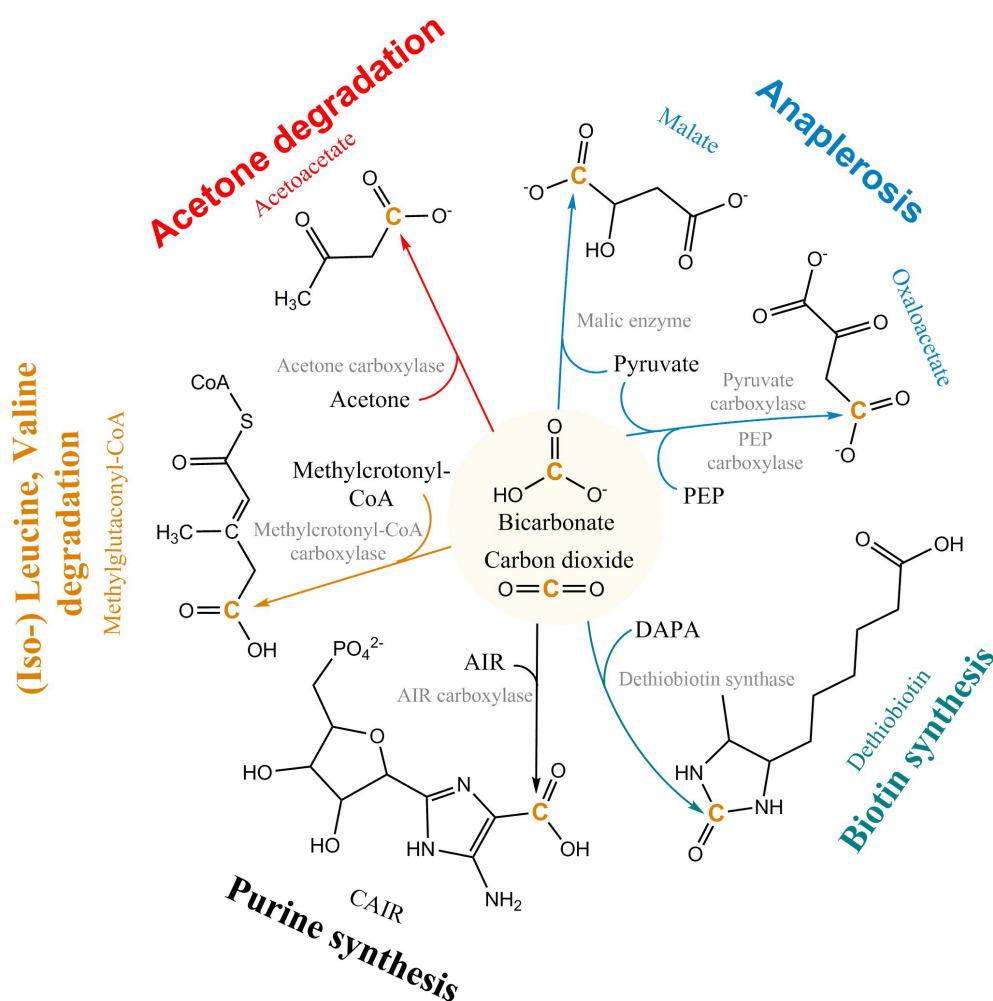
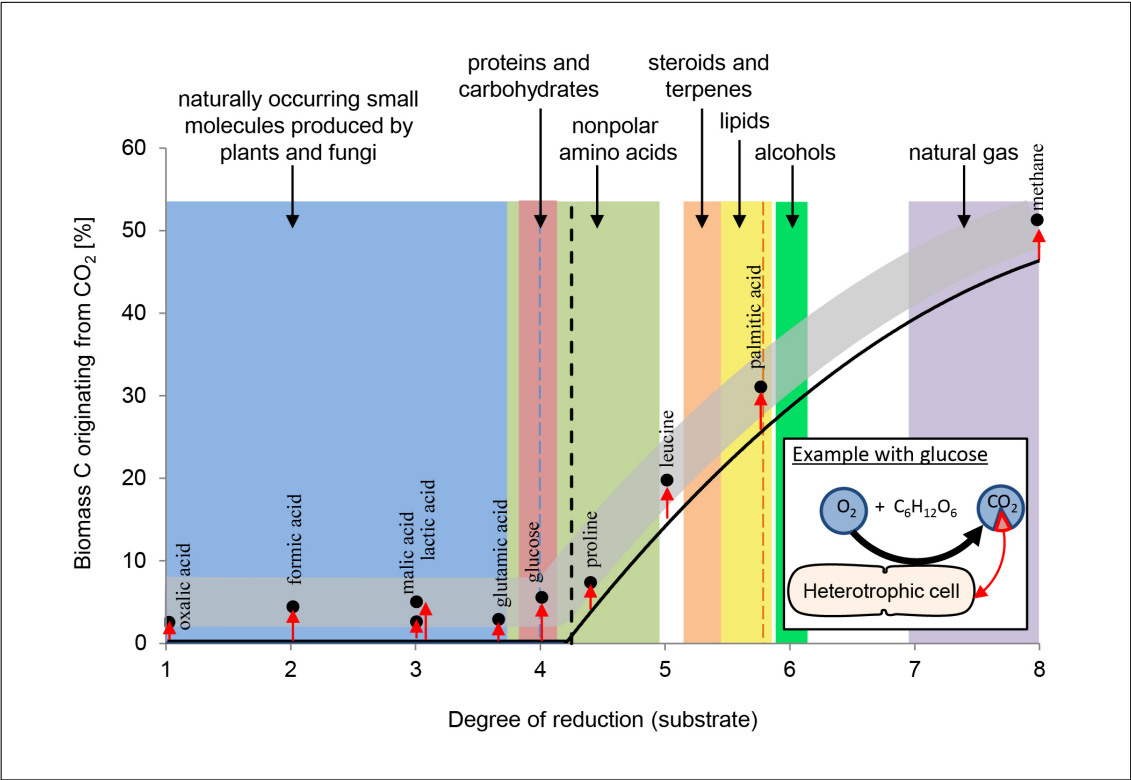


Figure 1: Selected heterotrophic CO₂ fixation reactions and pathways. PEP: phosphoenolpyruvate, DAPA: 7,8-diaminononanoate, AIR: 1-(5'-phosphoribosyl)-5-aminoimidazole, CAIR: 1-(5-phospho-D-ribosyl)-5-amino-4-imidazolecarboxylate, CoA: Coenzyme-A.

The most important CO₂ fixation pathway in all organisms is anaplerosis. Anaplerosis replenishes intermediates in the tricarboxylic acid (TCA) cycle, which have been released for biosynthesis. TCA metabolites are used as building blocks for macromolecular compounds, e.g. almost half of all amino acids in prokaryotes are directly synthesized from oxaloacetate and α-ketoglutarate (Fuchs 1999). For this purpose, heterotrophs use the enzymes pyruvate carboxylase present in a large variety of organisms, including prokaryotes, archaea, yeasts,

fungi and higher organisms (e.g. mammals), and phosphoenol pyruvate (PEP) carboxylase, widely distributed in bacteria (Attwood 1995; Jitrapakdee and Wallace 1999; Sauer and Eikmanns 2005; Jitrapakdee et al. 2008) (Fig. 1). The replenishment of metabolites continuously withdrawn from the TCA cycle via the anaplerotic reaction of PEP carboxylase entails an assimilation of CO₂ corresponding to 25% of the initial substrate's carbon content. In a systematic stable isotope labelling experiments with *Bacillus subtilis*, a gram-positive heterotrophic bacterium widespread in the environment, the interdependency of pathways and rates of CO₂-fixation on the concurrent utilization of organic substrate(s) was explored (Spona-Friedl et al. 2020). Over the course of the experiments *B. subtilis* assimilated 6% and 5% of carbon biomass from the external H¹³CO₃ pool when growing on glucose and lactate, respectively (Spona-Friedl et al. 2020). Growth on malate, an intermediate of the TCA cycle, expected to serve directly to refill the oxaloacetate pool of the TCA cycle, still revealed a contribution to biomass production from inorganic carbon of 3% (Spona-Friedl et al. 2020). PEP carboxylase was still actively transforming pyruvate to oxaloacetate. Heterotrophic CO₂-fixation continued to a lower extent even in the absence of cell growth during the stationary phase (Spona-Friedl et al. 2020), indicating that anaplerotic reactions are important in low-productivity habitats (see below).

Overall, heterotrophic CO₂ fixation via anaplerosis in microorganisms contributes around 1 to 8% to the carbon biomass (Romanenko 1964, Perez and Martin 1982, Doronina and Trotsenko 1984, Miltner et al. 2004, Roslev et al. 2004, Hesselsoe et al. 2005, Sandruckova et al. 2005, Feisthauer et al. 2008, Akyniede et al. 2020, Spona-Friedl et al. 2020). Under particular environmental conditions even higher contributions were reported (Perez and Martin 1982). The advantage that CO₂ is readily available to the cell either as atmospheric gas or, more commonly, in its hydrated form HCO₃⁻, obviously outcompetes the disadvantage that carboxylation is generally an endergonic reaction (Faber et al. 2015). This thermodynamic obstacle may be less important when carboxylation supports the assimilation of organic substrates more reduced than the organism's biomass, resulting in carbon-limited but excess-energy conditions (Heijnen and Roels, 1981, Ensign et al. 1998, von Stockar et al. 2006, Battley 2013). In this case, in addition to anaplerosis further carboxylation reactions are induced (Fig. 1) to add oxidized C (from CO₂) to the reduced organic substrate for adjusting the degree of reduction to that of the biomass (Fig. 2). For example, the assimilation of leucine and propionate into biomass entails carboxylation of the initial C-6 and C-3 carbon bodies, respectively and thus, triggers an assimilation of dissolved inorganic carbon (DIC) that corresponds to 17% and 33% of the initial substrate's carbon content, respectively (Erb 2011). In aerobic methane oxidation, the full oxidation potential of one molecule of CO₂ is needed to adjust the high degree of reduction of methane to that of biomass during its assimilation. Consequently, methanotrophs derive up to 50% of their carbon biomass from CO₂ (Strong, et al. 2015, Battley 2013).



127

128 **Figure 2:** Anaplerotic CO₂ fixation contributes 1-8% of carbon biomass (indicated by the grey band) in
129 heterotrophic cells. Dependent to the organism and in relation to the uptake of the individual organic
130 compounds and their entry into the TCA cycle and central metabolic pathways the relative amount of
131 inorganic carbon assimilated varies, as highlighted by the red arrows. See examples for malic and lactic acid.
132 With organic carbon sources more reduced than the organism's biomass (right to the dashed line) further
133 carboxylation reactions are induced, increasing the overall carbon contribution from CO₂ beyond anaplerosis
134 (grey band). In methanotrophs, 50% of the cell's carbon may originate from CO₂ fixation. For further
135 explanations see text.

136

137 Besides the degree of reduction of organic carbon sources, the partial pressure of CO₂ plays
138 a role. Carboxylases may catalyze carboxylation as well as decarboxylation of organic
139 compounds and the equilibrium of the reaction depends on the concentrations of all
140 compounds involved. An increase in the CO₂ concentration may move the equilibrium of the
141 reaction toward the product of the carboxylation, and thus leading to an increase in CO₂
142 fixation.

143 In a physiological context, the amount of inorganic carbon fixed by heterotrophs, either
144 from an endogenous or exogenous source, may be dependent on the metabolic state of the
145 organisms and the specific environmental conditions. In their early work, Romanenko et al.
146 (1972) suggested that the rate of heterotrophic anaplerotic fixation of DIC is strictly
147 proportional to the heterotrophic bacterial carbon production. Since then, a number of

factors have been identified potentially influencing the relative contribution of anaplerotic and other non-autotrophic CO₂ fixation reactions on biomass production. In laboratory experiments with the bacterial strain *Thiobacillus novellus*, for example, a higher amount of CO₂ was fixed under nutrient limited conditions (Perez and Matin 1982). Moreover, mixotrophic bacterial strains fixed more DIC compared to those grown autotrophically (Perez and Matin 1982). Fungi fixed relatively more CO₂ at lower organic carbon (glucose and maltose) concentrations (Schinner et al. 1982). The degree of heterotrophic CO₂ fixation highly depended on the availability of easy degradable organic carbon sources (Schinner et al. 1982).

Studies on the possible relationship between heterotrophic DIC fixation and the activity of prokaryotic cells revealed contradicting results. While Roslev et al. (2004) mentioned actively growing cells fix more DIC than resting cells, Merlin et al. (2003) report enhanced uptake of DIC by heterotrophic bacteria during slow growth and starvation. A relationship between DIC and heterotrophic bacterial production has been reported frequently as exemplified below.

2. Respiration and non-autotrophic CO₂ fixation

The production of CO₂ via respiration and the parallel fixation of CO₂ in heterotrophs take place simultaneously. The heterotrophic fixation of CO₂ is thus generally considered a back-reaction, i.e., part of the originally produced CO₂ from respiration is re-assimilated. Following this line of arguments, the more reduced an organic substrate is the less CO₂ is released (Fig. 2). Heterotrophic fixation of DIC does not necessarily lead to a net carbon biomass production, however, if microbes oxidize geogenic methane, this would result in a net carbon biomass production. Experimentally it is difficult to differentiate respiratory CO₂ flux from concurrent anaplerotic CO₂ fixation. As a consequence, there are numerous experiments and field studies determining dark CO₂ fixation, but only a few studies quantified the assimilation of DIC by non-autotrophs.

Respiration in aquatic systems is frequently determined via the consumption of dissolved oxygen (Robinson and Williams 2005) potentially underestimating the carbon use efficiency of heterotrophs. Depending on the substrate, the respiration quotient ($\Delta\text{CO}_2/\Delta\text{O}_2$) varies between 0.7 – 1.3 (Robinson 2019) leading to an error between 20 and 40% with regard to CO₂ production from respiration. Moreover, the respiration quotient also varies because other oxygen consuming processes are potentially taking place simultaneously (e.g. nitrification) (Robinson 2019). For instance, it is 138 O₂ for 106 CO₂ for ideal Redfield type organic matter, and 150 O₂ for 106 CO₂ for more realistic marine organic matter (Fraga et al. 1998; Paulmier et al. 2009). Calculations based on a study on temperate forest soils revealed a reduction of overall CO₂ emissions due to dark CO₂ fixation by mainly

heterotrophic microbes (Akinyede et al. 2020). Collectively, with respect to C cycling, heterotrophic CO₂ fixation and the carbon flux from the inorganic pool into heterotrophic biomass can be regarded as a process more important than hitherto assumed.

3. CO₂ fixation in habitats dominated by heterotrophs

In contrast to sunlit habitats, where **photo**autotrophs make up a significant portion of the total biomass and photosynthesis is of major importance in carbon cycling, heterotrophs and chemolithoautotrophs represent the **prevailing** biota in the “dark habitats”, i.e., soils, subsurface environments and the deep sea. These dark environments exceed their photic counterparts in both, volume and biomass. In the oceans, the deep sea (below 200 m) exceeds the sunlit surface layer by a factor of 18 in volume and, remarkably, by a factor of two in biomass (Aristegui et al. 2009). Therefore, the so-called “dark CO₂ fixation” does not only occur in specific 'hot spots' on the seafloor (hydrothermal vents, cold seeps and mud volcanoes), or in anoxic waters, but throughout the entire oxygenated 'dark' water column (Reinthal et al., 2010, Yakimov et al., 2014). **In limnic environments, the dark groundwater ecosystems outnumber surface waters 100-fold in terms of water volume (Danielopol et al. 2003), and similarly, also soils are with the exception of their surface exclusively dark habitats.**

Yet, heterotrophic CO₂ fixation does not occur only in the dark environments since heterotrophs are also found in the photic zone. This is particularly relevant in the ocean because the photic zone is where the highest biomass concentrations are found. Recently, it has been estimated that the inclusion of dark CO₂ fixation (integrated over the euphotic layer, 0-150 m depth) would increase oceanic primary production estimates by 2.5–22 % (Baltar et al., 2019). **A similar situation might be assumed for surface inland waters, however, global estimations are missing so far.**

Dark DIC fixation has been reported for all types of ecosystems, including marine habitats (Wuchter et al. 2003, Middelburg 2011, DeLorenzo et al. 2012, Molari et al. 2013, Baltar and Herndl 2019, Lengger et al. 2019, Smith et al. 2019, Vasquez-Cardenas et al. 2020), brackish and freshwater systems (Bräuer et al. 2013, Santoro et al. 2013, Noguerola et al. 2015, Signori et al. 2017, Vick-Majors and Priscu 2019, Zhao et al. 2020), cave waters and groundwater ecosystems (Pedersen & Ekendahl 1992a, 1992b; Kotelnikova & Pedersen 1998, Kellermann et al. 2012, Lazar et al. 2017), and soil habitats (Ehleringer et al. 2000, Miltner et al. 2004, 2005, Šantrůčková et al. 2005, 2018, Akinyede et al. 2020 and references therein). In the absence of solar radiation, particularly in the dark ocean, CO₂ fixation rates of up to ~125 mg C m⁻³ d⁻¹ have been measured, amounting to 30% (on a per volume basis) of the phototrophic CO₂ fixation in ocean surface waters (Zopfi et al. 2001, Detmer et al. 1993, Casamayor et al. 2001, Baltar et al. 2010). **In a eutrophic lagoon, dark DIC fixation**

accounted for 31% of total DIC fixation in the water column (Lliros et al. 2011). Recently it was shown that the ratio between dark/light CO₂ fixation in oceanic surface waters which is usually around 0.1 increases with depth reaching a ratio of 1 at 120-160 m depth (Baltar et al., 2019). In the past, however, dark DIC fixation has frequently been attributed to the activity of chemoautotrophs only. A few studies provide quantitative prove or at least striking evidence for heterotrophic CO₂ fixation (Tab. 1).

As indicated, part of the dark CO₂ fixation in oceans has been attributed to chemolithoautotrophic archaea (Wuchter et al. 2003, Ingalls et al. 2006) obtaining the energy required for the endergonic carboxylation through the oxidation of reduced inorganic compounds, such as ammonia or hydrogen sulfide (Swan et al. 2011; Zhang et al. 2020). A total annual chemolithoautotrophic CO₂ fixation rate of 0.77Pg C was calculated for the oceans (Middelburg 2011). The observed fluxes of the reduced inorganic compounds available as energy sources, however, seem largely insufficient to explain the relatively high dark CO₂ fixation rates (Overbeck 1979, Tuttle and Jannasch 1979, Baltar et al. 2010, Reinthaler et al. 2010, Herndl and Reinthaler 2013). In some cases, the supply rates of the reduced inorganic compounds used as an energy source explain less than 40% of the observed dark CO₂ fixation rates (Zopfi et al. 2001). Recently, chemoautotrophic nitrification was estimated to explain <13% of the dark CO₂ fixation (integrated over the euphotic zone) with the rest coming from either heterotrophic DIC fixation or other chemoautotrophic processes (Baltar and Herndl 2019).

The potential energy sources for the unexplained proportion of the dark CO₂ fixation remain enigmatic. Possible explanations could be either an underestimation of the supply rates of reduced inorganic compounds or the uptake of CO₂ by heterotrophic organisms (Zopfi et al. 2001, Baltar et al. 2019). In the surface ocean in particular, DIC incorporation via anaplerotic reactions might play an important role in compensating metabolic imbalances in marine bacteria under oligotrophic conditions, contributing > 30 % of the carbon incorporated into biomass (González et al. 2008; Palovaara et al., 2014). Evidence for the latter comes from experiments with Arctic seawater, which exhibited high DIC fixation rates (0.5–2.5 µg C L⁻¹ d⁻¹) correlating with heterotrophic bacterial production (Alonso-Sáez et al. 2010). Using different molecular tools, DIC uptake was attributed mainly to heterotrophic *Gamma*- and *Betaproteobacteria* rather than to typical chemoautotrophs, thus showing that chemolithoautotrophs were not the main drivers of CO₂ fixation in this habitat (Alonso-Sáez et al. 2010). Further evidence comes from the genome of *Polaribacter* sp. MED152, a representative of Bacteroidetes, which typically comprise about 10–20% of the prokaryotic abundance in seawater (González et al. 2008). A unique combination of membrane transporters and carboxylases in these organisms indicates the importance of anaplerosis besides other DIC fixation pathways (González et al. 2008). If the heterotrophic metabolism of bacteria is suddenly intensified (e.g., after an input of organic matter), dark DIC fixation rates and the expression of transcripts associated with key anaplerotic enzymes increase

proportionally (Baltar et al., 2016). As mentioned above, contradicting results were obtained on the relationship between heterotrophic CO₂ fixation and the availability of organic matter. A few studies suggest a relative increase in dark DIC fixation in oligotrophic habitats harboring slow-growing or starving bacterial populations (Perez and Matin 1982, Schinner et al. 1982, Merlin et al. 2003, Alonso-Sáez et al. 2010, Santoro et al. 2013). Considering the slow community-wide specific growth rates of heterotrophic bacteria in oligotrophic and/or cold waters, such as the marine aphotic zone, the Arctic Ocean, deep sea sediments, groundwater systems and the terrestrial subsurface, alpine limnic systems and deep-lake sediments, enhanced anaplerotic DIC uptake can be expected. However, there is also evidence for the stimulation of dark DIC fixation in response to organic matter enrichment in different types of soils (Miltner et al. 2005, Šantrůčková et al. 2018). Hence, these contradictory findings require further, more systematic research.

Other environmental factors that may influence dark DIC fixation include the concentrations of CO₂ and bicarbonate as inorganic carbon sources. An increase in the CO₂ concentration may shift the equilibrium of the carboxylation-decarboxylation reactions increasing CO₂ fixation. Elevated partial pressure of CO₂ might stimulate dark DIC fixation. In temperate forest soils, rates of dark microbial CO₂ fixation were positively correlated with the CO₂ concentration (Spohn et al. 2019). Similarly, with increasing CO₂ concentrations, higher dark DIC fixation was observed in wetland soils affected by subcrustal CO₂ degassing (Beuling et al. 2015). Here, besides known chemoautotrophs, CO₂ fixation via anaplerotic reactions was shown for putatively heterotrophs, i.e., subdivision 1 Acidobacteriaceae, lacking enzymatic pathways for autotrophic CO₂ fixation (Beuling et al. 2015). In experiments with two marine heterotrophic bacterial isolates, elevation of CO₂ concentration provoked an increase in CO₂ fixation along with a decrease in respiration (Teiro et al. 2012). Thus, we may assume that a rise in CO₂ concentrations and CO₂-induced geochemical changes will alter carbon turnover in affected ecosystems with dark DIC fixation and anaplerotic reactions becoming more important.

4. Quantitative estimates of heterotrophic CO₂ fixation in different environments

Heterotrophic CO₂ fixation in different habitats

Quantitative data on heterotrophic DIC fixation mainly originate from laboratory experiments using cultures and tissues. Measurements of dark DIC fixation with a proven or estimated significant contribution of heterotrophic assimilation of DIC are scarce. In Table 1, we provide a compilation of studies conducted in soils, marine and limnic ecosystems. Where possible, we compared dark DIC fixation rates with heterotrophic activity. In marine and limnic systems, heterotrophic carbon production as a widely applied activity measurement was used. In soils, we compared dark DIC fixation rates with respiration, i.e.,

CO₂ production. Dark DIC fixation rates in different marine systems range between 0.1 and 206 $\mu\text{g C L}^{-1} \text{ d}^{-1}$ with highest values found in a eutrophic lagoon and lowest values in the deep waters of the Mediterranean Sea (Tab. 1). Data from limnic systems originate from lake sediments with dark DIC fixation rates between 0.12 and 48 $\text{mg C m}^{-2} \text{ d}^{-1}$ (Tab. 1). Projecting these numbers to only the top 10 cm of sediment in the different lakes (which is a gross simplification), values of 1.2-480 $\mu\text{g C L}^{-1} \text{ sediment d}^{-1}$ are obtained. When compared to rates of bacterial carbon production, dark DIC fixation rates in these habitats accounted for a considerable fraction of total carbon assimilation, occasionally even exceeding it (Tab. 1). In soils, the dark DIC fixation rates which were attributed mainly to the activity of heterotrophs amounted to 0.04-39% of the overall respiration rate (Tab. 1). Dark DIC fixation rates range from 36 ng C to 23.6 $\mu\text{g C g}^{-1} \text{ d}^{-1}$ ranging over three orders of magnitude (Tab. 1). The contribution of heterotrophically fixed DIC to biomass carbon of microbes ranged from 0.2-1.1% in temperate forest soil (Akinyede et al. 2020), 0.2-4.6% in temperate forest and field soils (Santruckova et al. 2005), to 7% in arable soil (Miltner et al. 2004). Santruckova et al. (2005) estimated the overall heterotrophic CO₂ fixation to be even higher, i.e., 1.9-11.3% taking into account that the labile fraction of the biodegradable organic carbon resulted from metabolites released by spilling reactions of microorganisms due to a limitation in inorganic nutrients or due to the presence of highly reduced energy-rich carbon sources (e.g. Tempest et al. 1992). A contribution of heterotrophic CO₂ fixation to biomass carbon of 6.5±2.8% was found in drinking water biofilms and activated sludge (Roslev et al. 2004).

Carbon biomass stock originating from heterotrophic CO₂ fixation

While it is difficult to derive global estimations from the few studies that measured heterotrophic CO₂ fixation rates in marine, limnic and terrestrial ecosystems, we may use a conservative approach assuming that at least 1-5% of carbon biomass of all heterotrophs originates from anaplerotic DIC fixation. Earth's total living biomass is estimated to amount to about 499 – 738 Pg C, of which approx. 451 – 653 Pg C is photoautotrophic biomass (Bar-On et al. 2018). Heterotrophic biomass thus contributes 47 – 85 Pg C (Table SI-1). The uncertainties of the estimates of heterotrophic biomass of the terrestrial subsurface, however, are high (Whitman et al. 1998, McMahon and Parnell 2014, Bar-On et al. 2018). Nevertheless, following this line of evidence anaplerotic CO₂ fixation contributes between 0.5 – 5 Pg C to the living biomass.

Tab. 1.: Dissolved inorganic carbon (DIC) assimilation rates from a range of aquatic (marine and limnic) and soil environments. Dark carbon fixation (DCF) is shown as fraction of either bacterial heterotrophic production (BP) or respiration. Original data were converted to similar units whenever possible to allow comparison.

Aquatic ecosystems	Depth [m]	DIC fixation [$\mu\text{g C L}^{-1} \text{d}^{-1}$]	BP [$\mu\text{g C}^{-1} \text{d}^{-1}$]	DCF/BP [%]	Source	Remarks
Arctic	Seawater cultures	0.5-2.3	0.4-2.5	100%	Alonso-Saéz et al. 2010	Only potential for DCF
Mediterranean Sea	4900	0.096 \pm 0.02	0.048	200%	Yakimov et al. 2014	Only anaplerotic
Tropical South China Sea	200-1500	0.72-1.68	0.48- 4.8	40-105%	Zhou et al. 2017	Probably a large fraction anaplerotic
Tropical Estuary	1-18	4.8-14.4	55.2-1142	1.3-9%	Signori et al. 2018	Probably mostly anaplerotic
Eutrophic lagoon	1-5	206			Lliros et al. 2011	Probably mostly anaplerotic
Boreal lakes sediments	1-3	13.2-48 $\text{mg C m}^{-2} \text{d}^{-1}$	BP 96-216 $\text{mg C m}^{-2} \text{d}^{-1}$	8.4-37.4%	Santoro et al. 2013	Probably a large fraction anaplerotic
Tropical lakes sediments	1-3	0.12-20.4 $\text{mg C m}^{-2} \text{d}^{-1}$	BP 14.4- 583 $\text{mg C m}^{-2} \text{d}^{-1}$	0.4-80.4%	Santoro et al. 2013	Probably a large fraction anaplerotic
Deep granitic groundwater biofilms	812-1240	0. 2-2 $\mu\text{g C m}^{-2} \text{d}^{-1}$	n.d.	n.d.	Ekendahl and Pedersen 1994	Probably a large fraction anaplerotic
Terrestrial ecosystems		DIC fixation [$\mu\text{g C g}^{-1} \text{d}^{-1}$]	R [$\mu\text{g CO}_2\text{-C g}^{-1} \text{d}^{-1}$]	DCF/R [%]		
Temperate forest soil	0-0.7	0.036-0.32	0.95-19.1	1.2-3.9%	Spohn et al. 2019	^{13}C label mainly in AA, indicating anaplerosis
	0-1	0.06-0.86	n.d.	n.d.	Akinyede et al. 2020	Dominance of heterotrophs
Temperate agricultural soil	0-0.3	0.26	.63	2.7%	Miltner et al. 2004	Probably a large fraction anaplerotic
	0-0.3	0.19	9.82	1-5%	Miltner et al. 2005	DCF mainly driven by aerobic heterotrophs
Range of temperate forest & field soils	0.05-0.15	1.82-23.6*	0.65-9.16	3-39%	Šantrůčková et al. 2005	Probably a large fraction anaplerotic
	0-0.15	0.035-0.4	n.d.	n.d.	Nel and Cramer 2019	Probably mostly anaplerotic
Arctic tundra soils		0.04-0.08	0.79-10.7	0.04-16%	Šantrůčková et al. 2018	Anaplerotic enzymes comprised the majority of carboxylase genes.

*Values taken from Table 2 in Akinyede et al. 2020
n.d. not determined

Carbon flux related to heterotrophic CO_2 fixation

In terms of annual global heterotrophic production rates, oceans and the terrestrial subsurface (including soils) are the main habitats of heterotrophic CO_2 fixation (Cole et al. 2002; Magnabosco et al. 2018) (Table SI-2). Recently, Akinyede et al. (2020) estimated a

global dark CO₂ fixation rate of all temperate forest soils of 0.26 ± 0.07 Pg C yr⁻¹. We calculated a global heterotrophic C production of 34 – 245 Pg C yr⁻¹, which would translate into **0.34 – 12.3 Pg** of DIC bound by heterotrophic CO₂ fixation each year. Interestingly, these numbers are consistent with the recently calculated contribution of CO₂ fixation for the integrated epipelagic ocean of ca. 1.2– 11 Pg C yr⁻¹ (Baltar and Herndl 2019). This is a significant carbon flux amounting to **0.3-14%** of the global net amount of carbon produced annually by photoautotrophs (90 – 110 Pg C yr⁻¹; Ciais et al. 2013).

Our estimates are subject to a high uncertainty, which, on the one hand, results from the dependency of the extent of heterotrophic CO₂ fixation on the organic carbon oxidized and, on the other hand, on the predominant environmental conditions. Moreover, data on terrestrial and marine subsurface environments, although large in dimension, are **scarce**. For these environments, no detailed information on the abundance, growth (yield) and metabolic activity of microbial communities is available, particularly with increasing depth. Most of the deeper subsurface environments, even when harboring considerable living biomass, do not participate in the global carbon cycle on a short and medium time scales (years to decades), but rather in centennial to geological time scales. Nevertheless, in order to provide a first estimate and to be able to roughly evaluate the relevance of heterotrophic CO₂ fixation for all habitats of high uncertainty (e.g. the continental subsurface) we adopted a conservative approach (see also **Tables SI-1 and SI-2**).

5. Conclusions

Current models of carbon cycling and carbon sequestration do not account for heterotrophic CO₂ fixation (Gruber et al. 2004, Le Quéré et al. 2009). Despite the uncertainties in the data on heterotrophic biomass and production rates for some habitats (e.g. the terrestrial subsurface), the numbers presented here represent the first attempt to quantify the global contribution and relevance of heterotrophic CO₂ fixation to carbon cycling. Our results indicate that heterotrophs significantly contribute to global CO₂ fixation – especially (although not restricted to) **in habitats experiencing elevated CO₂ concentrations and/or lacking a sufficient supply of degradable organic carbon**. In specific environments, this may explain the mismatch between autotrophic C input, consumption, and sequestration that **has been** observed in marine systems (Baltar et al. 2009, Burd et al. 2010, Reinthaler et al. 2010, **Morán et al. 2007**, Hoppe et al. 2002, Tait and Schiel 2013). Particularly in aphotic habitats (which outnumber the photic habitats in both size and volume) such as the dark ocean, seafloor sediments, soils, as well as the sediments and rocks of the terrestrial subsurface (Miltner et al. 2004, Miltner et al. 2005, Yakimov et al. 2014, Wegener et al. 2012), carbon cycling needs to be re-evaluated taking into account anaplerotic CO₂ fixation and other inorganic carbon uptake pathways in heterotrophs. In seafloor sediments, wetlands and marshes, as well as in other habitats where methane

oxidation is a key process, a large fraction (10-50%) of heterotrophic biomass potentially originates from heterotrophic DIC fixation. Recently, a time-series study showed a tendency towards higher ratios of dark to light DIC fixation in the top half of the euphotic layer (0– 65 m) in the years 2012-2019 than in the preceding years (data started in 1989), which was linked to oceanographic changes (i.e., a deepening of the mixed zone) (Baltar et al., 2019). Moreover, the metabolic theory of ecology posits that heterotrophic metabolism increases more than gross primary production in the ocean in response to warming (see Baltar et al., 2019 and reference therein), which might also make heterotrophic DIC fixation relatively more important in a warmer ocean. In the light of global warming leading to an extensive thawing of permafrost soils and providing new habitats for methanotrophs, these processes are expected to become more important in the future. Hence, the potential contribution of heterotrophic CO₂ fixation under climate change conditions clearly deserves further investigations.

Author contributions

A.B., M.E. and C.G. conceived the idea for the manuscript. A.B., G.J.H. and C.G. wrote the manuscript. M.S.F., M.E., M.A. F.B. and T.R. substantially commented on and edited the manuscript. M.A., M.S.F. and C.G. did the literature search on available global carbon data. C.G. and M.A. performed the estimation of heterotrophic CO₂ fixation on a global scale.

Acknowledgments

We acknowledge B.B. Jørgensen for commenting on an earlier draft of the manuscript. We thank R. Thauer and W. Eisenreich for fruitful discussions on heterotrophic CO₂ fixation. Financial support was provided by the Wittgenstein Prize (Austrian Science Fund, project number Z194-B17), by the European Research Council under the European Community's Seventh Framework Program (FP7/2007-2013) / ERC grant agreement No. 268595 (MEDEA project) and the Austrian Science Fund (P 28781-B21) to G.J.H. Financial support was further provided by the Helmholtz Center Munich to A.B., M.E., M.S.F. and C.G.

References:

- Akyniede, R., Taubert, M., Schrumpf, M., Trumbore, S. & Küsel, K. Rates of dark CO₂ fixation are driven by microbial biomass in a temperate forest soil. *Soil Biol. Biochem.* 150, 107950, 2020.
- Alonso-Sáez, L., Galand, P. E., Casamayor, E. O., Pedrós-Alió, C., and Bertilsson, S.: High bicarbonate assimilation in the dark by Arctic bacteria, *ISME J.*, 4, 1581–1590, 2010.

418 Arístegui, J., Gasol, J. M., Duarte, C. M., and Herndl, G. J. Microbial oceanography of the dark ocean's
419 pelagic realm. *Limnol. Oceanogr.*, 54, 1501-1529, 2009.

420 Attwood, P. V. The structure and the mechanism of action of pyruvate-carboxylase. *Int. J. Biochem.*
421 *Cell B* 27, 231–249, 1995.

422 Baltar, F., and Herndl, G. J. Ideas and perspectives: Is dark carbon fixation relevant for oceanic
423 primary production estimates? *Biogeosci.*, 16, 3793-3799, 2019.

424 Baltar, F., Bayer, B., Bednarsek, N., Deppeler, S., Escribano, R., Gonzalez, C. E., ... , and Robinson, C.
425 Towards integrating evolution, metabolism, and climate change studies of marine ecosystems.
426 *Trends Ecol. Evol.*, 34, 1022-1033, 2019.

427 Baltar, F., Arístegui, J., Sintes, E., Gasol, J. M., Reinthaler, T., and Herndl, G. J. Significance of non-
428 sinking particulate organic carbon and dark CO₂ fixation to heterotrophic carbon demand in the
429 mesopelagic northeast Atlantic. *Geophys. Res. Lett.*, 37, 1-6, 2010.

430 Baltar, F., Lundin, D., Palovaara, J., Lekunberri, I., Reinthaler, T., Herndl, G. J., and Pinhassi, J.
431 Prokaryotic responses to ammonium and organic carbon reveal alternative CO₂ fixation pathways
432 and importance of alkaline phosphatase in the mesopelagic North Atlantic. *Front. Microbiol.*, 7,
433 1670, 2016.

434 Bar-Even, A., Noor, E., and Milo, R. A survey of carbon fixation pathways through a quantitative lens.
435 *J. Exp. Bot.* 63, 2325–2342, 2012.

436 Bar-On, Y. M., Phillips, R., and Milo, R. The biomass distribution on Earth. *PNAS*, 115, 6506-6511,
437 2018.

438 Battley, E. H. A theoretical study of the thermodynamics of microbial growth using *Saccharomyces*
439 *cerevisiae* and a different free energy equation. *Quart. Rev. Biol.*, 88, 69-96, 2013.

440 Beer, C., Reichstein, M., Tomelleri, E., Ciais, P., Jung, M., Carvalhais, N., Rödenbeck, C., Arain, M. A.,
441 Baldocchi, D., Bonan, G. B., Bondeau, A., Cescatti, A., Lasslop, G., Lindroth, A., Lomas, M., Luyssaert,
442 S., Margolis, H., Oleson, K. W., Rouspard, O., Veenendaal, E., Viovy, N., Williams, C., Woodward, F. I.,
443 and Papale, D. Terrestrial gross carbon dioxide uptake: global distribution and covariation with
444 climate. *Science*, 329, 834-838, 2010.

445 Berg, I. A. Ecological aspects of the distribution of different autotrophic CO₂ fixation pathways. *Appl.*
446 *Environ. Microbiol.*, 77, 1925-1936, 2011.

447 Berg, I. A., Kockelkorn, D., Buckel, W., and Fuchs, G. A 3-hydroxypropionate/4-hydroxybutyrate
448 autotrophic carbon dioxide assimilation pathway in Archaea. *Science*, 318, 1782-1786, 2007.

449 Beulig, F., Heuer, V.B., Akob, D.M., Viehweger, B., Elvert, M., Herrmann, M., Hinrichs, K.-U., and
450 Küsel, K. Carbon flow from volcanic CO₂ into soil microbial communities of a wetland mofette. *ISME*
451 *J.* 9, 746–759, 2015.

452 Bräuer, S.L., Kranzler, K., Goodson, N., Murphy, D., Simon, H.M., Baptista, A.M., and Tebo, B.M. Dark
453 carbon fixation in the Columbia River's estuarine turbidity maxima: molecular characterization of
454 red-type *cbbI* genes and measurement of DIC uptake rates in response to added electron donors.
455 *Estuarine, Coast. Shelf Sci.* 36, 1073-1083, 2013.

456 Burd, A. B., Hansell, D. A., Steinberg, D. K., Anderson, T. R., Arístegui, J., Baltar, F., Beupre, S. R.,
457 Buesseler, K. O., De- Hairs, F., Jackson, G. A., Kadko, D. C., Koppelman, R., Lampitt, R. S., Nagata, T.,
458 Reinthaler, T., Robinson, C., Robison, B. H., Tamburini, C., and Tanaka, T.: Assessing the apparent
459 imbalance between geochemical and biochemical indicators of meso-and bathypelagic biological
460 activity: What the @ \$?! Is wrong with present calculations of carbon budgets?, *Deep-Sea Res. Pt. II*,
461 57, 1557–1571, 2010.

462 Casamayor, E. O., García-Cantizano, J., Mas, J., and Pedrós-Alió, C. Primary production in estuarine
 463 oxic/anoxic interfaces: contribution of microbial dark CO₂ fixation in the Ebro River Salt Wedge
 464 Estuary. *Mar. Ecol. Prog. Ser.*, 215, 49-56, 2001.

465 Ciais, P., C. Sabine, G. Bala, L. Bopp, V. Brovkin, J. Canadell, A. Chhabra, R. DeFries, J. Galloway, M.
 466 Heimann, C. Jones, C. Le Quéré, R. B. Myneni, S. Piao, and P. Thornton. Carbon and other
 467 biogeochemical cycles. In *Climate change 2013: The physical science basis. Contribution of working*
 468 *group I to the fifth assessment report of the Intergovernmental Panel on Climate Change*, eds. T. F.
 469 Stocker, D. Qin, G.-K. Plattner, M. Tignor, S. K. Allen, J. Boschung, A. Nauels, Y. Xia, V. Bex & P. M.
 470 Midgley, 465-570. Cambridge, United Kingdom and New York, NY, USA: Cambridge University Press,
 471 2013.

472 **Cochrane, V.W. Physiology of fungi. John Wiley, New York, 1958.**

473 Cole, J. J., Findlay, S. E. G., and Pace, M. L. Bacterial production in fresh and saltwater ecosystems : a
 474 cross-system overview. *Mar. Ecol. Prog. Ser.*, 43, 1-10, 1988.

475 **Danielopol, D. L., Griebler, C., Gunatilaka, A., and Notenboom, J. Present state and future prospects**
 476 **for groundwater ecosystems. *Environ. Conserv.* 30, 104-130, 2003.**

477 **DeLorenzo, S., Bräuer, S. L., Edgmont, C. A., Herfort, L., Tebo, B.M., and Zuber, P. Ubiquitous**
 478 **dissolved inorganic carbon assimilation by marine bacteria in the Pacific Northwest Coastal Ocean as**
 479 **determined by stable isotope probing. *PLoS ONE* 7, e46695, 2012.**

480 Detmer, A. E., Giesenhausen, H. C., Trenkel, V. M., Auf dem Venne, H., and Jochem, F. J. Phototrophic
 481 and heterotrophic pico- and nanoplankton in anoxic depths of the central Baltic Sea. *Mar. Ecol.*
 482 *Progr. Ser.*, 99, 197-203, 1993.

483 Dijkhuizen, L., and Harder, W. Current views on the regulation of autotrophic carbon dioxide fixation
 484 via the Calvin cycle in bacteria. *Antonie van Leeuwenhoek*, 50, 473-87, 1984.

485 Doronina, N. V., and Trotsenko, Y. A. The levels of carbon dioxide assimilation in bacteria with
 486 different pathways of 1-carbon metabolism. *Mikrobiologiya*, 53, 885-889, 1984.

487 **Ensign, S. A., Small, F. J., Allen, J. R., Sluis, M. K. New roles for CO₂ in the metabolism of aliphatic**
 488 **epoxides and ketones. *Arch. Microbiol.* 169, 179–187, 1998.**

489 Erb, T. J., Brecht, V., Fuchs, G., Muller, M., Alber, B. E. Carboxylation mechanism and stereochemistry
 490 of crotonyl-CoA carboxylase/reductase, a carboxylating enoyl-thioester reductase. *PNAS* 106, 8871–
 491 8876, 2009.

492 Erb, T. J. Carboxylases in natural and synthetic microbial pathways. *Appl. Environ. Microbiol.*, 77,
 493 8466-8477, 2011.

494 **Evans, E. A., Jr., and Slotin, L. The utilization of carbon dioxide in the synthesis of α-ketoglutaric acid.**
 495 ***J. Biol. Chem.*, 136, 301, 1940.**

496 Faber, K., Fessner, W. D., and Turner, N. J. *Science of synthesis: biocatalysis in organic synthesis Vol.*
 497 *2.* 672. Thieme Chemistry, 2015.

498 **Feisthauer, S., Wick, L. Y., Kastner, M., Kaschabek, S. R., Schlomann, M., Richnow, H. H., Differences**
 499 **of heterotrophic ¹³CO₂ assimilation by *Pseudomonas knackmussii* strain B13 and *Rhodococcus opacus***
 500 **1CP and potential impact on biomarker stable isotope probing. *Environ. Microbiol.* 10, 1641–1651,**
 501 **2008.**

502 Fraga, F., Rios, A., Perez, F., and Figueras, F. Theoretical limits of oxygen:carbon and oxygen:nitrogen
 503 ratios during photosynthesis and mineralisation of organic matter in the sea. *Mar. Chem.*, 62, 161–
 504 168, 1998.

505 Fuchs, G. Biosynthesis of building blocks. In Biology of the prokaryotes, eds. Lengeler, J. W., Drews,
506 G., and Schlegel, H. G., 110-160, Stuttgart, New York: Thieme, 1999.

507 González, J. M., Fernández-Gómez, B., Fernández-Guerra, A., Gómez-Consarnau, L., Sánchez, O., Coll-
508 Lladó, M., del Campo, J., Escudero, L., Rodríguez-Martínez, R., Alonso-Sáez, L., Latasa, M., Paulsen, I.,
509 Nedashkovskaya, O., Lekumberri, I., Pinhassi, J., and Pedrós-Alió, C.: Genome analysis of the
510 proteorhodopsin-containing marine bacterium *Polaribacter* sp. MED152 (Flavobacteria), P. Natl.
511 Acad. Sci. USA, 105, 8724– 8729, 2008.

512 Gruber, N., Friedlingstein, P., Field, C., Valentini, R., Heimann, M., Richey, J. E., Romero-Lankao, P.,
513 Schulze, E. D. & Chen, C.-T. A. The vulnerability of the carbon cycle in the 21st century: an assessment
514 of carbon-climate-human interactions. In: The global carbon cycle: integrating humans, climate, and
515 the natural world, eds. Field, C. B., and Raupach, M. R., 45-76. Washington D.C., London: Island
516 Press, 2004.

517 Han, L., Yang, K., Kulowski, K., Wendt-Plienkowski, E., Hutchinson, C. R., and Vining, L. C. An acyl-
518 coenzyme A carboxylase encoding gene associated with jadomycin biosynthesis in *Streptomyces*
519 *venezuelae* ISP5230. Microbiol. UK 146, 903–910, 2000.

520 Hartman, R. E., and Keen, N. T. Enzymes catalysing anaplerotic carbon dioxide fixation in *Verticillium*
521 *albo-atrum*. Phytopathol. 63, 947-953, 1973.

522 Hartman, R. E., Keen, N. T., and Long, M. Carbon dioxide fixation by *Verticillium albo-atrum*. J. Gen.
523 Microbiol. 73, 29-34, 1972.

524 Heijnen, J. J., and Roels, J. A. A macroscopic model describing yield and maintenance relationship in
525 aerobic fermentation processes. Biotechnol. Bioeng. 23, 739–763, 1981.

526 Herndl, G. J., and Reinthaler, T. Microbial control of the dark end of the biological pump. Nat. Geosc.,
527 6, 718-724, 2013.

528 Hesselsoe, M., Nielsen, J. L., Roslev, P., and Nielsen, P. H. Isotope labeling and microautoradiography
529 of active heterotrophic bacteria on the basis of assimilation of ¹⁴CO₂. Appl. Environ. Microbiol., 71,
530 646-655, 2005.

531 Hoppe, H. G., Gocke, K., Koppe, R., and Begler, C. Bacterial growth and primary production along a
532 north-south transect of the Atlantic Ocean. Nature, 416, 168-171, 2002.

533 Houghton, R. A. Balancing the global carbon budget. Ann. Rev. Earth Planet. Sci., 35, 313-347, 2007.

534 Ingalls, A. E., Shah, S. R., Hansman, R. L., Aluwihare, L. I., Santos, G. M., Druffel, E. R., and Pearson, A.
535 Quantifying archaeal community autotrophy in the mesopelagic ocean using natural radiocarbon.
536 PNAS, 103, 6442-6447, 2006.

537 Jitrapakdee, S., and Wallace, J.C. Structure, function and regulation of pyruvate carboxylase.
538 Biochem. J. 340, 1–16, 1999.

539 Jitrapakdee, S., St. Maurice, M., Rayment, I., Cleland, W.W., Wallace, J.C., and Attwood, P.V.
540 Structure, mechanism and regulation of pyruvate carboxylase. Biochem. J. 413, 369–387, 2008.

541 Kellermann, C., Selesi, D., Lee, N., Hügler, M., Esperschütz, J., Hartmann, A. & Griebler, C. Microbial
542 CO₂ fixation potential in a tar-oil-contaminated porous aquifer. FEMS Microbiol. Ecol. 81, 172-187,
543 2012.

544 Kleiber, M., Smith, A. H., and Black, A. L. Carbonate as precursor of milk constituents in the intact
545 dairy cow. J. Biol. Chem., 195, 707-714, 1952.

546 Kornberg, H. L., and Krebs, E. H. Synthesis of cell constituents from C₂-units by a modified
547 tricarboxylic acid cycle. Nature 179, 988–991, 1957.

548 Kornberg, H.L. Anaplerotic sequences in microbial metabolism. *Angew. Chem. internat. Edit.* 4, 558-
549 565, 1965.

550 Kotelnikova, S., and Pedersen, K. Distribution and activity of methanogens and homoacetogens in
551 deep granitic aquifers at Äspö Hard Rock Laboratory, Sweden. *FEMS Microbiol. Ecol.*, 26, 121-134,
552 1998.

553 Krebs, H.A. Carbon dioxide assimilation in heterotrophic organisms. *Nature* 147, 560-563, 1941.

554 Lazar, C.S., Stoll, W., Lehmann, R., Herrmann, M., Schwab, V.F., Akob, D.M., Nawaz, A., Wubet, T.,
555 Buscot, F., Totsche, K.-U., and Küsel, K. Archaeal diversity and CO₂ fixers in carbonate-/siliciclastic-
556 rock groundwater ecosystems. *Archaea* 2136287, 1-13, 2017.

557 Le Quéré, C., M. R. Raupach, J. G. Canadell, G. Marland, L. Bopp, P. Ciais, T. J. Conway, S. C. Doney, R.
558 A. Feely, P. Foster, P. Friedlingstein, K. Gurney, R. A. Houghton, J. I. House, C. Huntingford, P. E. Levy,
559 M. R. Lomas, J. Majkut, N. Metzl, J. P. Ometto, G. P. Peters, I. C. Prentice, J. T. Randerson, S. W.
560 Running, J. L. Sarmiento, U. Schuster, S. Sitch, T. Takahashi, N. Viovy, G. R. van der Werf & F. I.
561 Woodward. Trends in the sources and sinks of carbon dioxide. *Nat. Geosci.*, 2, 831-836, 2009.

562 Le Quéré, C., R. M. Andrew, J. G. Canadell, S. Sitch, J. I. Korsbakken, G. P. Peters, A. C. Manning, T. A.
563 Boden, P. P. Tans, R. A. Houghton, R. F. Keeling, S. Alin, O. D. Andrews, P. Anthoni, L. Barbero, L.
564 Bopp, F. Chevallier, L. P. Chini, P. Ciais, K. Currie, C. Delire, S. C. Doney, P. Friedlingstein, T. Gkritzalis,
565 I. Harris, J. Hauck, V. Haverd, M. Hoppema, K. Klein Goldewijk, A. K. Jain, E. Kato, A. Körtzinger, P.
566 Landschützer, N. Lefèvre, A. Lenton, S. Lienert, D. Lombardozzi, J. R. Melton, N. Metzl, F. Millero, P.
567 M. S. Monteiro, D. R. Munro, J. E. M. S. Nabel, S. I. Nakaoka, K. O'Brien, A. Olsen, A. M. Omar, T. Ono,
568 D. Pierrot, B. Poulter, C. Rödenbeck, J. Salisbury, U. Schuster, J. Schwinger, R. Séférian, I. Skjelvan, B.
569 D. Stocker, A. J. Sutton, T. Takahashi, H. Tian, B. Tilbrook, I. T. van der Laan-Luijkx, G. R. van der Werf,
570 N. Viovy, A. P. Walker, A. J. Wiltshire & S. Zaehle. Global Carbon Budget 2016. *Earth System Science*
571 *Data*, 8, 605-649, 2016.

572 Lengger, S.K., Rush, D., Mayser, J.P., Blewett, J., Schwartz-Narbonne, R., Talbot, H.B., Middelburg,
573 J.J., Jetten, M.S.M., Schouten, S., Sinninghe Damsté, J.S., and Pancost, R.D. Dark carbon fixation in
574 the Arabian Sea oxygen minimum zone contributes to sedimentary organic carbon (SOM). *Global*
575 *Biogeochem. Cycl.* 33, 1715-1732, 2019.

576 Lliros, M., Alonso-Saéz, L., Gich, F., Plasencia, A., Auguet, O., Casamayor, E.O., and Borrego, C.M.
577 Active bacteria and archaea cells fixing bicarbonate in the dark along the water column of a stratified
578 eutrophic lagoon. *FEMS Microbiol. Ecol.* 77, 370-384, 2011.

579 Magnabosco, C., Lin, L. H., Dong, H., Bomberg, M., Ghiorse, W., Stan-Lotter, H., Pedersen, K., Kieft, T.
580 L., van Heerden, E., and Onstott, T. C. The biomass and biodiversity of the continental subsurface.
581 *Nature Geoscience*, 11, 707-717, 2018.

582 McMahon, S., and Parnell J. Weighing the deep continental biosphere. *FEMS Microbiol. Ecol.*, 87,
583 113-120, 2014.

584 Melzer, E., and O'Leary M. H. Anapleurotic CO₂ fixation by phosphoenolpyruvate carboxylase in C3
585 plants. *Plant Physiol.*, 84, 58-60, 1987.

586 Merlin, C., Masters, M., McAteer, S., and Coulson, A. Why is carbonic anhydrase essential to
587 *Escherichia coli*? *J. Bacteriol.* 185, 6415-6424, 2003.

588 Middelburg, J. J. Chemoautotrophy in the ocean. *Geophy. Res. Lett.*, 38, 1-4, 2011.

589 Miltner, A., Kopinke, F.-D., Kindler, R., Selesi, D., Hartmann, A., and Kästner, M. Non-phototrophic
590 CO₂ fixation by soil microorganisms. *Plant Soil*, 269, 193-203, 2005.

591 Miltner, A., Richnow H.-H., Kopinke F.-D., and Kästner, M. Assimilation of CO₂ by soil microorganisms
592 and transformation into soil organic matter. *Org. Geochem.*, 35, 1015-1024, 2004.

593 Molari, M., Manini, E., and Dell'Anno, A. Dark inorganic carbon fixation sustains the functioning of
594 benthic deep-sea ecosystems. *Glob. Biogeochem. Cycl.* 27, 212-221, 2013.

595 Morán, X. A. G., Pérez, V. & Fernández, E. Mismatch between community respiration and the
596 contribution of heterotrophic bacteria in the NE Atlantic open ocean: What causes high respiration
597 in oligotrophic waters? *J. Mar. Res.*, 65, 545-560, 2007.

598 Nel, J.A., and Cramer, M.D. Soil microbial anaplerotic CO₂ fixation in temperate soils. *Geoderma* 335,
599 170-178, 2019.

600 Noguerola, I., Picazo, A., Lliros, M., Camacho, A., and Borrego, C.M. Diversity of freshwater
601 Epsilonproteobacteria and dark inorganic carbon fixation in the sulphidic redoxcline of a meromictic
602 karstic lake. *FEMS Microbiol. Ecol.* 91, fiv086, 2015.

603 Overbeck, J. Dark CO₂ uptake - biochemical background and its relevance to in situ bacterial
604 production. *Arch. Hydrobiol. Beiheft*, 12, 38-47, 1979.

605 Palovaara, J., Akram, N., Baltar, F., Bunse, C., Forsberg, J., Pedrós- Alió, C., González, J. M., and
606 Pinhassi, J.: Stimulation of growth by proteorhodopsin phototrophy involves regulation of central
607 metabolic pathways in marine planktonic bacteria, *P. Natl. Acad. Sci. USA*, 111, E3650–E3658, 2014.

608 Parkinson, S. M., Jones, R., Meharg, A. A., Wainwright, M., and Killham, K. The quantity and fate of
609 carbon assimilated from ¹⁴CO₂ by *Fusarium oxysporum* grown under oligotrophic and near
610 oligotrophic conditions. *Mycol. Res.* 95, 1345–1349, 1991

611 Parkinson, S. M., Killham, K., and Wainwright, M. Assimilation of ¹⁴CO₂ by *Fusarium oxysporum*
612 grown under oligotrophic conditions. *Mycol. Res.* 94, 959–964, 1990.

613 Paulmier, A., Kriest, I. & Oschlies, A. Stoichiometries of remineralisation and denitrification in global
614 biogeochemical ocean models. *Biogeosci.* 6, 923–935, 2009.

615 Pedersen, K., and Ekendahl, S. Assimilation of CO₂ and introduced organic compounds by bacterial
616 communities in groundwater from southeastern Sweden deep crystalline bedrock. *Microb. Ecol.* 23,
617 1-14, 1992.

618 Pedersen, K., and Ekendahl, S. Incorporation of CO₂ and introduced organic compounds by bacterial
619 populations in groundwater from deep crystalline bedrock of Stripa mine. *J. Gen. Microbiol.* 138,
620 369-376, 1992.

621 Perez, R.C., and Matin, A. Carbon dioxide assimilation by *Thiobacillus novellus* under nutrient-limited
622 mixotrophic conditions. *J. Bacteriol.* 150, 46-51, 1982.

623 Reinthaler, T., Van Aken, H. M., and Herndl, G. J. Major contribution of autotrophy to microbial
624 carbon cycling in the deep North Atlantic, Åôs interior, *Deep-Sea Res. Pt. II*, 57, 1572– 1580, 2010.

625 Robinson, C., and Williams, P.J. Respiration and its measurement in surface marine waters. In:
626 Respiration in aquatic ecosystems (eds. P. A. del Giorgio and P. J. Williams) Oxford: Oxford University
627 Press, 2005.

628 Robinson, C. Microbial respiration, the engine of ocean deoxygenation. *Front. Mar. Sci.*, 5, 533, 2019.

629 Romanenko, V. I. Heterotrophic CO₂ assimilation by water bacterial flora. *Mikrobiologiya*, 33, 679-
630 683, 1964.

631 Romanenko, V. I., Overbeck, J., and Sorokin, Y. I. Estimation of production of heterotrophic bacteria
632 using ¹⁴C. In: Sorokin, Y. I., Kadota, H. (eds.) *Techniques for the assessment of microbial production*
633 *and decomposition in fresh waters. IBP Handbook No. 23*, Blackwell, Oxford, pp. 82-85, 1972.

634 Roslev, P., Larsen, M. B., Jørgensen, D. & Hesselsoe, M. Use of heterotrophic CO₂ assimilation as a
635 measure of metabolic activity in planktonic and sessile bacteria. *J. Microbiol. Meth.*, 59, 381-393,
636 2004.

637 Santoro, A.L., Bastviken, D., Gudas, C., Tranvik, L., Enrich-Prast, A. Dark carbon fixation: an
638 important process in lake sediments. *PLoS ONE* 8: e65813, 2013.

639 Šantrůčková, H., Bird, M. I., Elhottova, D., Novak, J., Pícek, T., Simek, M., and Tykva, R. Heterotrophic
640 fixation of CO₂ in soil. *Microb. Ecol.* 49, 218–225, 2005.

641 Šantrůčková, H., Kotas, P., Bárta, J., Urich, T., Čapek P., Palmtag J., Eloy Alves, R. J., Biasi, C., Diáková,
642 K., Gentsch, N., Gittel, A., Guggenberger, G., Hugelius, G., Lashchinsky, N., Martikainen, P. J.,
643 Mikutta, R., Schleper, C., Schneckner, J., Schwab, C., Shibistova, O., Wild, B., and Richter, A.
644 Significance of dark CO₂ fixation in arctic soils. *Soil Biol. Biochem.* 119, 11–21, 2018.

645 Sauer, U., and Eikmanns B. J. The PEP–pyruvate–oxaloacetate node as the switch point for carbon
646 flux distribution in bacteria. *FEMS Microbiol. Rev.*, 29, 765-794, 2005.

647 Schink, B. An alternative to the glyoxylate shunt. *Mol. Microbiol.* 73, 975–977, 2009.

648 Schinner, F., Concin, R., & Binder, H. Heterotrophic CO₂ -fixation by fungi in dependence on the
649 concentration of the carbon source. *Phyton* 22, 81-85, 1982.

650 Scrutton, M. C. Assay of enzymes of CO₂ metabolism. *Methods in Microbiology* Vol 6, Part A, 479-
651 541, 1971.

652 Signori, C. N., Valentin, J. L., Pollery, R. C. G., and Enrich-Prast, A. Temporal variability of dark carbon
653 fixation and bacterial production and their relation with environmental factors in a tropical estuarine
654 system, *Estuaries and Coasts*, 41, 1089–1101, 2018.

655 Smith, A. R., Kieft, B., Mueller, R., Fisk, M. R., Mason, O. U., Popa, R., and Colwell, F. S. Carbon
656 fixation and energy metabolisms of a subseafloor olivine biofilm. *ISME J.*, 13, 1737-1749, 2019.

657 Spohn, M., Müller, K., Höschen, C., Mueller, C.W., and Marhan, S. Dark microbial CO₂ fixation in
658 temperate forest soils increases with CO₂ concentrations. *Global Change Biology* 26, 1926-1935,
659 2019.

660 Spona-Friedl, M., Braun, A., Huber, C., Eisenreich, W., Griebler, C., Kappler, A., and Elsner M.
661 Substrate-dependent CO₂-fixation in heterotrophic bacteria revealed by stable isotope labelling.
662 *FEMS Microbiol. Ecol.*, 96, fiae080, 2020.

663 Strong, P. J., Xie, S., and Clarke, W. P. Methane as a resource: can the methanotrophs add value?
664 *Environ. Sci. Technol.*, 49, 4001-4018, 2015.

665 Swan, B. K., Martinez-Garcia M., Preston C. M., Sczyrba A., Woyke T., Lamy D., et al. Potential for
666 chemolithoautotrophy among ubiquitous bacteria lineages in the dark ocean. *Science* 333, 1296-
667 1300, 2011.

668 Tait, L. W., and Schiel D. R. Impacts of temperature on primary productivity and respiration in
669 naturally structured macroalgal assemblages. *PLoS ONE*, 8, e74413, 2013.

670 Teiro, E., Fernández, A., Álvarez-Salgado, X. A., García-Martín, E. E., Serret, P., and Sobrino, C.
671 Response of two marine bacterial isolates to high CO₂ concentration *Mar. Ecol. Prog. Ser.*, 453, 27–36,
672 2012.

673 Tempest, D. W., and Neijssel, O. M. Physiological and energetic aspects of bacterial metabolite
674 overproduction. *FEMS Microbiol. Lett.* 100, 169–176, 1992.

675 Tuttle, J. H., and Jannasch H. W. Microbial dark assimilation of CO₂ in the Cariaco Trench. *Limnol.*
676 *Oceanogr.*, 24, 746-753, 1979.

677 Vasquez-Cardenas, D., Meysman, F. J. R., & Boschker, H. T. S. A cross-system comparison of dark
678 carbon fixation in coastal sediments. *Glob. Biogeochem. Cycl.* 34, 1-14, 2020.

679 Vick-Majors, T. J., and Priscu, J. C. Inorganic carbon fixation in ice-covered lakes of the McMurdo Dry
680 Valleys. *Antarctic Sci.* 1-10, 2019.

681 von Stockar, U., Maskow, T., Liu, J., Marison, I. W., and Patiño, R. Thermodynamics of microbial
682 growth and metabolism: An analysis of the current situation. *J. Biotechnol.*, 121, 517-533, 2006.

683 Wegener, G., Bausch, M., Holler, T., Thang, N. M., Mollar, X. P., Kellermann, M. Y., Hinrichs, K. U.,
684 and Boetius, A. Assessing sub-seafloor microbial activity by combined stable isotope probing with
685 deuterated water and ^{13}C -bicarbonate. *Environ. Microbiol.*, 14, 1517-1527, 2012.

686 Werkman, C.H., and Wood, H.G. Heterotrophic assimilation of carbon dioxide. In: *Advances in*
687 *Enzymology and Related Areas of Molecular Biology* (Nord, F.F. and Werkman, C.H., eds.), 2, 135-
688 182, 1942.

689 Whitman, W. B., Coleman, D. C., and Wiebe, W. J. Prokaryotes: The unseen majority. *PNAS*, 95,
690 6578-6583, 1998.

691 Wood, H. G., and Werkman, C. H. The utilisation of CO_2 in the dissimilation of glycerol by the
692 propionic acid bacteria. *Biochem. J.*, 30, 48-53, 1936.

693 Wood, H.G., and Werkman, C.H. The utilization of CO_2 by the propionic acid bacteria. *Biochem. J.*, 32,
694 1262–1271, 1938.

695 Wood, H.G., and Werkman, C.H. The position of carbon dioxide-carbon in succinic acid synthesized
696 by heterotrophic bacteria. *Jour. Biol. Chem.*, 139, 377–381, 1941.

697 Wood, H. G., and Stjernholm, R. L. Assimilation of carbon dioxide by heterotrophic organisms. In
698 Gunsalus, IC, Stanier, RY (Eds.) *The Bacteria: A Treatise on Structure and Function*, vol 3. Biosynthesis
699 Academic Press, New York, 41–117, 1962.

700 Wuchter, C., Schouten, S., Boschker, H. T. S., and Sinninghe Damsté, J. S. Bicarbonate uptake by
701 marine Crenarchaeota. *FEMS Microbiol. Lett.*, 219, 203-207, 2003.

702 Yakimov, M. M., La Cono, V., Smedile, F., Crisafi, F., Arcadi, E., Leonardi, M., Decembrini, F.,
703 Catalfamo, M., Bargiela, R., Ferrer, M., Golyshin, P. N., and Giuliano, L. Heterotrophic bicarbonate
704 assimilation is the main process of de novo organic carbon synthesis in hadal zone of the Hellenic
705 Trench, the deepest part of Mediterranean Sea. *Environ. Microbiol. Rep.*, 6, 709–722, 2014.

706 Zhang, Y., Qin, W., Hou, L., Zakem, E.J., Wan, X., Zhao, Z., Liu, L., Hunt, K.A., Jiao, N., Kao, S.-J., Tang,
707 K., Xie, X., Shen, J., Li, Y., Chen, M., Dai, X., Liu, C., Deng, W., Dai, M., Ingalls, A.E., Stahl, D.A., and
708 Herndl, G.J. Nitrifier adaptation to low energy flux controls inventory of reduced nitrogen in the dark
709 ocean. *PNAS* 117, 4823-4830, 2020.

710 Zhao, Y., Liu, P., Rui, J., Cheng, L., Wang, Q., Liu, X., and Yuan, Q. Dark carbon fixation and
711 chemolithotrophic microbial community in surface sediments of the cascade reservoirs, Southwest
712 China. *Sci.Tot. Environ.* 698, 134316, 2020.

713 Zhou, W., Liao, J., Guo, Y., Yuan, X., Huang, H., Yuan, T., and Liu, S.: High dark carbon fixation in the
714 tropical South China Sea, *Cont. Shelf Res.*, 146, 82–88, 2017.

715 Zopfi, J., Ferdelman, T. G., Jørgensen, B. B., Teske, A., and Thamdrup, B. Influence of water column
716 dynamics on sulfide oxidation and other major biogeochemical process in the chemocline of
717 Mariager Fjord (Denmark). *Mar. Chem.*, 74, 29-51, 2001.