

# Carbon Isotope Fractionation by Autotrophic Bacteria with Three Different CO<sub>2</sub> Fixation Pathways

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*Dedicated to Professor Achim Trebst on the occasion of his 60th birthday*

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Carbon isotope fractionation during autotrophic growth of different bacteria which possess different autotrophic CO<sub>2</sub> fixation pathways has been studied. <sup>13</sup>C/<sup>12</sup>C-Ratios in the cell carbon of the following bacteria were determined (CO<sub>2</sub> fixation pathway suggested or proven in parentheses): *Alkaligenes eutrophus* (reductive pentose phosphate cycle), *Desulfobacterium autotrophicum* and *Acetobacterium woodii* (reductive acetyl-CoA pathway), *Desulfobacter hydrogenophilus* and *Thermoproteus neutrophilus* (reductive citric acid cycle). The Δδ<sup>13</sup>C values, which indicate the *per mille* deviation of the <sup>13</sup>C content of cell carbon from that of the CO<sub>2</sub> used as the sole carbon source, range from –10‰ (reductive citric acid cycle) over –26‰ (reductive pentose phosphate cycle) to –36‰ (reductive acetyl-CoA pathway). Acetate formed *via* the acetyl-CoA pathway by the acetogenic *Acetobacterium woodii* showed a Δδ<sup>13</sup>C = –40‰. These data are discussed in view of the different CO<sub>2</sub> fixation reactions used by the bacteria and especially with regard to the isotopic composition of sedimentary carbon through time.

## Introduction

Enzyme-catalyzed carboxylation reactions and subsequent reactions entail carbon isotope effects as a result of both thermodynamic and kinetic fractionations [1]. An outstanding example is the bias in favor of <sup>12</sup>C exercised by the ribulose 1,5-bisphosphate carboxylase reaction of Calvin cycle (C<sub>3</sub>) autotrophy, that discriminates against “heavy” carbon (<sup>13</sup>C) by about –20 to –35‰ (the negative sign indicates the <sup>13</sup>C content in cell carbon is 20 to 35‰ less than in the substrate CO<sub>2</sub>). Chemical reactions show much less pronounced carbon isotope discrimination. The isotopic composition of sedimentary carbon, which in the average deviates from inorganic carbonates by approximately –28‰ (or from CO<sub>2</sub> in the air by –21‰), is being taken as indication for biological CO<sub>2</sub> fixation by autotrophs. According to these assumptions and data autotrophic ecosystems became a dominant component of the terrestrial car-

bon cycle as early as  $3.5 \times 10^9$ , if not  $3.8 \times 10^9$  years ago.

In the past decade novel autotrophic CO<sub>2</sub> fixation pathways have been unravelled occurring mostly in strictly anaerobic prokaryotes; so far, the reductive pentose phosphate cycle (Calvin cycle) could not be found in this group ([2], for possible exception see [3]). Since the first autotrophic organisms on earth most likely were strict anaerobes, the knowledge of the magnitude of carbon isotope fractionation during autotrophic cell material synthesis from CO<sub>2</sub> in these bacteria might be of some value for the interpretation of geological isotopic composition data. On the other hand, if different CO<sub>2</sub> fixation pathways result in different carbon isotope fractionation, isotopic composition data may be taken as indication for the kind of CO<sub>2</sub> fixation reactions used.

Here we report on the carbon isotope fractionation during autotrophic growth by different bacteria which possess three different CO<sub>2</sub> fixation pathways. Special emphasis was put on comparable growth conditions and excess supply of CO<sub>2</sub>, such that the data reflect differences in the main CO<sub>2</sub> fixation reac-

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tions, rather than differences in temperature, residual CO<sub>2</sub> concentration, pH, or salinity of the medium.

## Experimental

### Organisms

*Desulfobacter hydrogenophilus* strain AcRS 1 (DSM 3380) (*Dbr. hydrogenophilus*) [4], *Desulfobacterium autotrophicum* strain HRM2 (DSM 3382) (*Dbm. autotrophicum*) [5], and *Acetobacterium woodii* (DSM 1030) (*Ac. woodii*) [6] were a kind gift of Dr. F. Widdel, University of Marburg (F.R.G.). *Alkaligenes eutrophus* strain H16 (DSM 428) [7] was a kind gift of Prof. B. Friedrich, Free University of Berlin (F.R.G.). *Thermoproteus neutrophilus* (DSM 2338) [8] was a kind gift of Prof. K. O. Stetter, University of Regensburg (F.R.G.).

### Growth of organisms

In order to compare <sup>13</sup>C/<sup>12</sup>C carbon isotope fractionation by different autotrophic microorganisms, the bacterial strains were grown under comparable conditions, with a large excess of CO<sub>2</sub>. The conditions were: Growth temperature 24 °C, except for *T. neutrophilus* (85 °C); culture volume 1 l; pH of medium between 6.8–7.2; mineral salt medium with CO<sub>2</sub> as the sole carbon source except for *Ac. woodii* which was grown with additional 0.02% yeast extract; continuous gassing with a 80% H<sub>2</sub>/20% CO<sub>2</sub>/0.1% H<sub>2</sub>S gas mixture at a minimal rate of 50 ml · min<sup>-1</sup> · l<sup>-1</sup> culture, except for *A. eutrophus* which was grown with 50% air/40% H<sub>2</sub>/10% CO<sub>2</sub>; stirring rate (magnetic stirring bar) was 100–150 rpm; the same type of all-glass 1 l fermenter with a filter candle, porosity 3 (Schott, Mainz, F.R.G.), for gassing was used. Except for *A. eutrophus* all organisms were grown under strictly anaerobic conditions. The cell yields were between 0.15–0.3 g dry cell matter · l culture<sup>-1</sup>, except for *A. eutrophus* which reached 2 g dry cell matter · l culture<sup>-1</sup>. The medium used for growth of *A. eutrophus* was described by [7], for *T. neutrophilus* by [9]; *Ac. woodii*, *Dbr. hydrogenophilus*, and *Dbm. autotrophicum* were grown in a salt water medium [10] as modified by [11]. Growth was determined by measuring the optical density at 578 nm (*d* = 1 cm), except for *T. neutrophilus* where the cell number was determined. Cells were harvested by centrifugation and lyophilized. Parallel to each experiment an aliquot of the gas mixture (Linde AG, Höllriegelskreuth) was continuously passed through two con-

secutive alkaline CO<sub>2</sub> traps containing 4 N KOH. Carbonate was precipitated as BaCO<sub>3</sub> [12]. From the culture supernatant of *Ac. woodii* acetate was obtained by diethylether extraction in a Kutscher-Steudel apparatus of the concentrated medium supernatant followed by vacuum distillation. Acetate from vinegar prepared from wine (Frings, Bonn, F.R.G.) was obtained after vapor distillation, diethylether extraction and crystallization.

### Determination of δ<sup>13</sup>C values

Cell material and Na-acetate were oxidized quantitatively to CO<sub>2</sub> with O<sub>2</sub> [13, 14] and precipitated as BaCO<sub>3</sub>. The <sup>13</sup>C analyses of BaCO<sub>3</sub> were performed in CO<sub>2</sub> gas extracted from the samples after acidification with 95% *ortho*-phosphoric acid. The δ<sup>13</sup>C value of CO<sub>2</sub> was determined on a specially equipped mass spectrometer MAT 230 (Varian). Corrections for the small isotope contribution of <sup>17</sup>O to the mass 45 peak were made. The δ<sup>13</sup>C values are presented as *per mille* deviation from the PDB standard (CO<sub>2</sub> obtained from Peedee Belemnite from the Peedee Formation, Upper-Cretaceous, South Carolina), as defined by [15]:

$$\delta^{13}\text{C} = \left[ \frac{^{13}\text{C}/^{12}\text{C sample}}{^{13}\text{C}/^{12}\text{C standard}} - 1 \right] \times 10^3 [\text{‰}].$$

Errors in determination of δ<sup>13</sup>C were about 0.3‰. The δ<sup>13</sup>C values of CO<sub>2</sub> in the gas mixture used in the different series of experiments were -49.8‰, -49.5‰, and -47.6‰, respectively. The Δδ<sup>13</sup>C cell value refers to the deviation of <sup>13</sup>C content of cell carbon from that of the CO<sub>2</sub> used as carbon source.

## Results and Discussion

The discrimination between the stable carbon isotopes <sup>12</sup>C and <sup>13</sup>C by five different facultatively autotrophic bacteria was studied under autotrophic growth conditions, *i.e.* CO<sub>2</sub> was supplied as the only source for cell carbon. All strains but one were strict anaerobes (*Desulfobacterium autotrophicum*, *Desulfobacter hydrogenophilus*, *Acetobacterium woodii*, *Thermoproteus neutrophilus*); as a reference strain the aerobic *Alkaligenes eutrophus* was grown under similar but aerobic conditions. The <sup>13</sup>C content in cell matter (after combustion to CO<sub>2</sub>) and in CO<sub>2</sub> gas used as carbon source was determined by mass spectroscopy. The values are related to a carbonate standard of defined <sup>13</sup>C content and are referred to as δ<sup>13</sup>C

values. The difference in  $^{13}\text{C}$  content between  $\text{CO}_2$  and cell carbon is referred to as  $\Delta\delta^{13}\text{C}$  value. The cultures were grown in 1 l fermenters under continuous gassing with gas mixtures containing  $\text{CO}_2$ . Fig. 1 summarizes the growth conditions and shows representative growth curves of the 5 strains. The generation time in the exponential phase varied between 10 h and 20 h and the maximal cell yield was 0.3 g dry weight  $\cdot \text{l}^{-1}$ . Since 50% of cell dry weight is carbon the maximal  $\text{CO}_2$  assimilation rate of the 1 l culture under those conditions ( $t_d = 10$  h,  $\mu = 1.15 \times 10^{-3} \cdot \text{min}^{-1}$ ; cell mass  $X = 0.3$  g = 150 mg C = 12.5 mmol C) follows from the growth equation  $\frac{dx}{dt} = \mu \cdot X$  as 15  $\mu\text{mol}$   $\text{CO}_2$  assimilated  $\cdot \text{min}^{-1} \cdot \text{l culture}^{-1}$ . The  $\text{CO}_2$  supply varied in the experiments from 330  $\mu\text{mol}$   $\text{CO}_2$  sup-

plied  $\cdot \text{min}^{-1} \cdot \text{l culture}^{-1}$  (*A. eutrophus*) to 1.34 mmol  $\text{CO}_2$  supplied  $\cdot \text{min}^{-1} \cdot \text{l culture}^{-1}$  (*Ac. woodii*); usually 440  $\mu\text{mol}$   $\text{CO}_2$  was supplied  $\cdot \text{min}^{-1} \cdot \text{l culture}^{-1}$ . Since  $\text{CO}_2$  is well soluble it was supplied in large excess. This is important in order to prevent that due to preferential  $^{12}\text{CO}_2$  fixation  $^{13}\text{CO}_2$  selectively accumulates and therefore the  $\delta^{13}\text{C}$  value of cell carbon becomes artificially low. The results of several independent growth experiments are given in Table I; the values represent means of at least two determinations. The table also includes the  $\text{CO}_2$  fixation pathways, which have been proven or suggested, as well as the most important  $\text{CO}_2$  fixation reactions and their estimated contribution to total fixed cell carbon. The most important  $\text{CO}_2$  fixing enzymes and reactions are summarized in Table II.

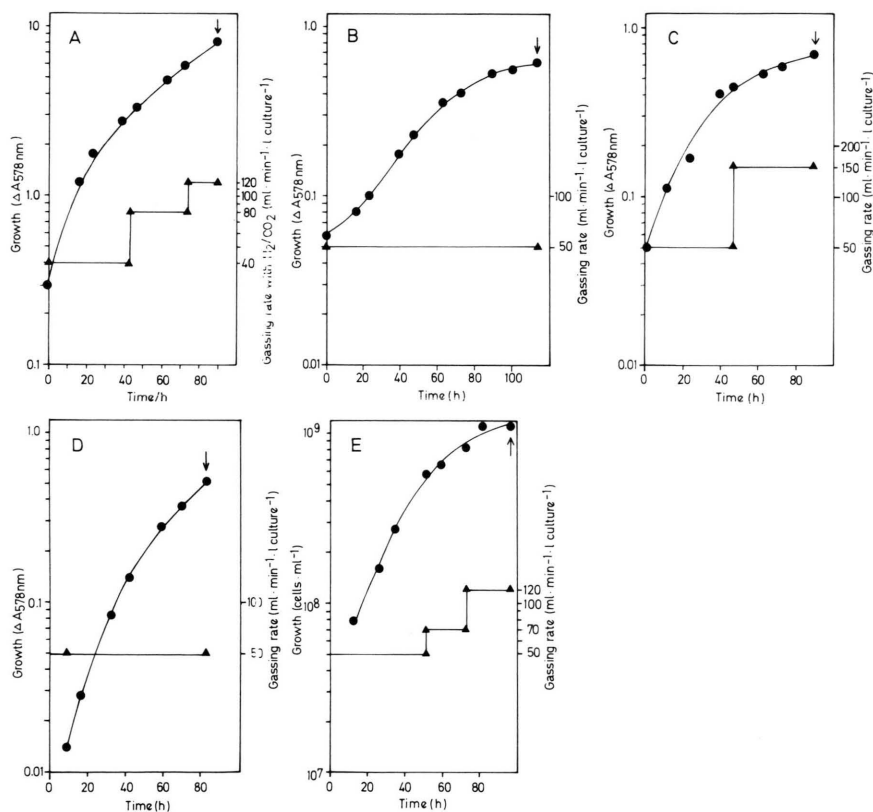


Fig. 1. Autotrophic growth of different bacteria on mineral salt medium at neutral pH and under excess gassing with  $\text{CO}_2$ .  $\text{CO}_2$  was the only carbon source added.

A. *Alkaligenes eutrophus* grown aerobically with  $\text{H}_2 + \text{O}_2$  as energy source (24 °C).

B. *Desulfobacterium autotrophicum* grown anaerobically with  $\text{H}_2$  + sulfate as energy source (24 °C).

C. *Acetobacterium woodii* grown anaerobically with  $\text{H}_2$  +  $\text{CO}_2$  as energy source (24 °C).

D. *Desulfobacter hydrogenophilus* grown anaerobically with  $\text{H}_2$  + sulfate as energy source (24 °C).

E. *Thermoproteus neutrophilus* grown anaerobically with  $\text{H}_2$  + elementary sulfur as energy source (85 °C).

The arrow indicates the time of harvest.

Table I. Fractionation of stable carbon isotopes  $^{13}\text{C}/^{12}\text{C}$  during autotrophic growth of bacterial cultures excessively gassed with 20%  $\text{CO}_2/80\%$   $\text{H}_2$  and 10%  $\text{CO}_2/40\%$   $\text{H}_2/50\%$  air (*Alkaligenes*), respectively. The main  $\text{CO}_2$  fixation reactions are presented in Table II. The estimation of the contribution of the individual  $\text{CO}_2$  fixation reaction to total carbon in the sample is based on the assumption that in the Calvin cycle 90% of cell carbon is derived from ribulose 1,5-bisphosphate carboxylase, and 10% from anaplerotic and other minor  $\text{CO}_2$  fixation reactions. In anaerobes in which acetyl CoA is central carbon fixation intermediate, approximately 2/3 of cell carbon is derived from acetyl CoA, the rest from additional carboxylations.

Organism or sample	Culture conditions	Autotrophic $\text{CO}_2$ Fixation pathway	$\delta^{13}\text{C}_{\text{cell}}$ [‰]	$\delta^{13}\text{C}_{\text{CO}_2}$ [‰]	$\Delta\delta^{13}\text{C}_{\text{cell}}$ [‰]	Main $\text{CO}_2$ fixation reactions (see Table II)	Estimated contribution of individual $\text{CO}_2$ fixation step to total cell carbon [%]
<i>Alkaligenes eutrophus</i>	28 °C pH 6.7	Reductive pentose phosphate cycle	-75.6	-49.5	-26.1	1 7	90 5
<i>Desulfobacterium autotrophicum</i>	28 °C pH 6.8–7.2	Reductive acetyl CoA pathway	-85.4 -84.8	-49.5 -49.8	-35.9 -35.0	2 3 4 7	33 33 20 10
<i>Acetobacterium woodii</i>	28 °C pH 6.8–7.0	Reductive acetyl CoA pathway	-70.4 -76.3 -62.4	-49.8 -49.5 -47.6	-20.6 -26.8 -14.8	2 3 4 8	33 33 20 10
Acetate from <i>Ac. woodii</i> culture	28 °C pH 6.8–7.0	Reductive acetyl CoA pathway	-89.5 -87.2	-49.8 -47.6	-39.7 -39.6	2 3	50 50
<i>Desulfobacter hydrogenophilus</i>	28 °C pH 6.8–7.2	Reductive citric acid cycle	-59.2 -57.9 -60.9	-49.8 -49.5 -47.6	- 9.4 - 8.4 -13.3	5 6 4 7	33 33 20 10
<i>Thermoproteus neutrophilus</i>	85 °C pH 6.8–6.9	Modified reductive citric acid cycle (?)	-55.8	-47.6	- 8.2	5 6 4 7	33 33 20 10

Table II. Main  $\text{CO}_2$  fixing enzymes and  $\text{CO}_2$  fixation reactions catalyzed by these enzymes in bacteria. [H] refers to reducing equivalents which cannot be specified in many cases. [CO] refers to carbonyl bound to CO dehydrogenase. For literature see [2, 18]; for active species of " $\text{CO}_2$ " utilized by these enzymes see [24] and literature quoted herein.

No.	$\text{CO}_2$ fixing enzyme	(E.C.)	$\text{CO}_2$ fixation reaction
1	Ribulose 1,5-bisphosphate carboxylase	4.1.1.39	$\text{CO}_2 + \text{D-Ribulose 1,5-bisphosphate} + \text{H}_2\text{O} \rightarrow 2 \times 3 \text{ phospho-D-glycerate}$
2	Formate dehydrogenase	1.2.-.-	$\text{CO}_2 + 2[\text{H}] \rightarrow \text{formate}$
3	CO dehydrogenase	1.2.99.2	$\text{CO}_2 + 2[\text{H}] \rightarrow [\text{CO}] + \text{H}_2\text{O}$
4	Pyruvate synthase	1.2.7.1 (?)	$\text{CO}_2 + \text{acetyl CoA} + 2[\text{H}] \rightarrow \text{pyruvate} + \text{coenzyme A}$
5	2-Oxoglutarate synthase	1.2.7.3 (?)	$\text{CO}_2 + \text{succinyl CoA} + 2[\text{H}] \rightarrow 2\text{-oxoglutarate} + \text{coenzyme A}$
6	Isocitrate dehydrogenase ( $\text{NADP}^+$ )	1.1.1.42	$\text{CO}_2 + 2\text{-oxoglutarate} + 2[\text{H}] \rightarrow \text{isocitrate}$
7	PEP carboxylase	4.1.1.31	$\text{HCO}_3^- + \text{phosphoenolpyruvate} \rightarrow \text{oxaloacetate} + \text{phosphate}$
8	PEP carboxykinase (pyrophosphate)	4.1.1.38	$\text{CO}_2 + \text{phosphoenolpyruvate} + \text{P}_i \rightarrow \text{oxaloacetate} + \text{pyrophosphate}$

*A. eutrophus*, which assimilates CO<sub>2</sub> via the Calvin cycle [16], discriminated <sup>13</sup>C by approximately -26‰. *Dbm. autotrophicum*, which assimilates CO<sub>2</sub> probably via a reductive acetyl CoA pathway [17–20], discriminated the heavy isotope even stronger (-36‰). *Ac. woodii* forms acetate via acetyl CoA from 2 CO<sub>2</sub> and 4 H<sub>2</sub> by a similar reductive acetyl CoA pathway [6, 21, 22]; this is reflected by a similar strong isotope discrimination (-40‰). In contrast cell carbon of this acetate forming bacterium which mostly derives from acetyl CoA had a Δδ<sup>13</sup>C value of -20 to -27‰. This relatively low value is unexpected because *Dbm. autotrophicum* and *Ac. woodii* use a similar CO<sub>2</sub> fixation mechanism. The interpretation of the data has to take into account that in the acetogenic bacterium two pathways are competing for acetyl CoA, i.e. conversion to acetyl phosphate and acetate for ATP synthesis [6], and assimilation into cell carbon via pyruvate, citrate, malonyl CoA and other acetyl CoA metabolizing reactions of intermediary metabolism [22]. Since approximately 12 times more acetyl CoA is converted to acetate than assimilated this eventually will lead to an accumulation of [<sup>13</sup>C]acetyl CoA due to the isotope effects exerted by phosphotransacetylase and acetate kinase. Consequently cell carbon will be enriched in <sup>13</sup>C relative to the acetate excreted. As a control acetate formed from ethanol by aerobic acetic acid bacteria in vinegar had a Δδ<sup>13</sup>C value of -25.3‰ (data not shown), if the δ<sup>13</sup>C value of atmospheric CO<sub>2</sub> was used as reference. This val-

ue corresponds to the Δδ<sup>13</sup>C value of C<sub>3</sub> plant material [reviewed in 1]. *Dbr. hydrogenophilus*, which assimilates CO<sub>2</sub> via acetyl CoA formed in a reductive citric acid cycle [23], exhibited a rather low <sup>13</sup>C discrimination (-10‰); a similar value was observed with *T. neutrophilus* which seems to assimilate CO<sub>2</sub> via a similar mechanism [9, unpublished results].

In conclusion the autotrophic organisms showed remarkable differences in their ability to discriminate the stable carbon isotopes. Three groups with similar fractionation were recognized each probably characterized by a distinct autotrophic CO<sub>2</sub> fixation mechanism. This is an additional argument – although indirect – for the existence of alternatives to the Calvin cycle in prokaryotes. Table III summarizes representative carbon isotope fractionation data in bacteria. Table II summarizes the main CO<sub>2</sub> fixation reactions in autotrophic organisms. We would like to emphasize three points. First, biological carbon isotope fractionation data should be interpreted with caution. One has to be aware of several CO<sub>2</sub> fixation mechanisms; even not all CO<sub>2</sub> fixation pathways in bacteria and their isotope effects may be known. Second, the Calvin cycle most likely was not the predominant autotrophic pathway in the early anaerobic biosphere, and attempts to relate isotope fractionation in geological specimens to this CO<sub>2</sub> fixation mechanism must be rejected on the basis of this study. A combination of the alternative CO<sub>2</sub> fixation pathways would lead to a similar discrimination as exerted by ribulose-1,5-bisphosphate carboxylase.

Table III. Stable carbon isotope <sup>13</sup>C/<sup>12</sup>C fractionation reported for autotrophic bacteria.

Organism	Δδ <sup>13</sup> C <sub>cell</sub> [‰]	References
<i>Alkaligenes eutrophus</i>	-26.1	this paper
<i>Desulfobacterium autotrophicum</i>	-35.5	this paper
<i>Desulfobacter hydrogenophilus</i>	-10.4	this paper
<i>Thermoproteus neutrophilus</i>	- 8.2	this paper
<i>Acetobacterium woodii</i>	-15 to -26.8	this paper
<i>Chromatium</i> , strain D	-22.5	[25]
<i>Rhodospirillum rubrum</i>	-20.5	[25]
<i>Chlorobium limicola thiosulfatophilum</i>	-12.2	[25]
<i>Methanobacterium thermoautotrophicum</i>	+5 to -34	[26]
<i>Chlorobium phaeovibrioides</i>	- 3.5	[27]
<i>Chlorobium vibrioforme</i>	- 3.9	[27]
	- 3.8	[27]
<i>Chromatium vinosum</i>	-19.6	[27]
<i>Rhodospirillum rubrum</i>	-12.4	[27]
<i>Rhodospseudomonas capsulata</i>	-10.6	[27]
<i>Thiomicrospira</i> sp.	-24.6	[28]
<i>Thiobacillus neapolitanus</i>	-25.1	[28]



Third, physical parameters such as temperature, pH of solution, salinity, limiting concentrations of CO<sub>2</sub>, and especially reactions of energy metabolism effectively competing for CO<sub>2</sub> (CO<sub>2</sub> reduction to methane or acetate) greatly influence the size of isotope discrimination by autotrophs. These parameters are completely unknown for the early biosphere. Their effect on biological isotope discrimination should be

studied in bacteria. No rationale for the magnitude of isotope fractionation can be given at present, since the isotope effects of the individual CO<sub>2</sub> metabolizing reactions are mostly unknown.

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