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Isotopomers of chlorophyll nuclei: Theories and an application

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Abstract

In this review, we discuss site-specific carbon isotopic compositions of chlorophylls with special reference to their biosynthetic pathways in the cell. The carbon isotopic composition is ultimately inherited from glutamic acid, a major amino acid in the cell. Intramolecular (site-specific) carbon isotopic distributions of geoporphyrins, geological derivatives of the tetrapyrrole nuclei of various chlorophylls in sediments and sedimentary rocks, are thus potentially useful for investigating the carbon cycle in the geological past. As a case study, we report analytical results for the mean carbon isotopic compositions of the methine bridge carbons (i.e., C–5, C–10, C–15, and C–20 positions in the tetrapyrrole structure) in geoporphyrins derived from chlorophylls *a* and *c* in the Miocene Onnagawa Formation, Japan. The results suggest that β -carboxylation was not an important process for carbon assimilation of photoautotrophs during the formation of this sediment. Although substantial biogeochemical information is recorded as isotopomeric signatures of sedimentary tetrapyrroles, we are still just beginning to explore its potential, and further experiments are necessary to establish this approach as a robust tool for biogeochemistry.

INTRODUCTION

This review concerns with the isotopomers (i.e., site-specific or intramolecular carbon isotopic compositions) as well as bulk carbon isotopic compositions of chlorophyll and

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bacteriochlorophyll nuclei in terms of metabolic processes. Among the various types of tetrapyrroles produced by organisms, chlorophylls are far more abundant, widespread, and obvious biological molecules in photoautotrophic organisms. Chlorophyll is an essential part of the photosynthetic apparatus and functions as a light energy-harvesting pigment.

The advantages of chlorophyll as a tool for biogeochemical and geological studies are threefold. First, chlorophylls have various chemical structures that are correlated with major photoautotrophic groups. Therefore, the structural variations and relative abundances of chloropigments in the natural environment provide first-order diagnostic information on the components of the photoautotrophic community. For example, if abundant chlorophyll b or its derivatives are observed in a given horizon of lake sediment, you can tell that the green algae were among the primary photoautotrophs in the lake when the sediment was formed. Second, chlorophyll contains both carbon and nitrogen atoms whose stable isotopic compositions (i.e., ${}^{13}C/{}^{12}C$ and ${}^{15}N/{}^{14}N$ ratios) provide valuable information about biogeochemical processes associated with assimilation and metabolism of these elements, allowing reconstruction of the surface water environment where the compounds were produced. Because most biomarkers widely used for investigation of the contemporary environment or paleoenvironment do not contain nitrogen, nitrogen isotopic record of chlorophylls uniquely contributes to understanding environmental processes related to the nitrogen cycle. Third, the tetrapyrrole nuclei of chlorophylls are well preserved in sediments and sedimentary rocks, oil shales, and even crude oil (e.g., Treibs 1934, 1936; Blumer 1965; Baker and Louda 1986; Eckardt et al. 1991; Callot and Ocampo 2000; Kashiyama et al. 2007b). Therefore, they are of particular interest for reconstruction of biogeochemical processes in the geological past. Despite these merits, chlorophyll abundance and their isotopes have been measured in relatively a limited number of studies from only a few laboratories (Hayes et al. 1987; Madigan et al. 1989; Popp et al. 1989; Katase and Wada 1990; Bidigare et al. 1991; Chicarelli et al. 1993; Sachs et al. 1999; Beaumont et al. 2000; Ohkouchi et al. 2006; Sato et al. 2006). It is mainly because 1) it requires time-consuming, tedious chemical/analytical procedures for isolation and purification of the chlorophylls (and their derivatives) from the complex mixtures of organic molecules in natural materials, and 2) the measurement of stable isotopic compositions can only be achieved by a classical off-line method rather than a conventional on-line method (e.g., Hayes et al. 1990; Meier-Augenstein 1999).

We have recently improved the isolation/purification protocol for chlorophylls and their diagenetic derivatives from sediments and particulate organic matter in the water column for precise, rapid determination of carbon and nitrogen isotopic compositions (e.g., Ohkouchi et al. 2005; Chikaraishi et al. 2005, 2008; Kashiyama et al. 2007a). Although the method still employs (automatic) fraction collection of chlorophylls using high-performance liquid chromatography (HPLC) followed by off-line measurement of isotopic compositions, it provides a strict analytical basis for studying the utility of these signatures. We have applied this method to several sedimentary rocks to better understand the surface water environment during their formation in the geological past (Kashiyama et al. 2008a, b). These results



	R ₃	R ₇	R ₈	7, 8-bond	R ₁₂	R_{13}^{2}	R ₂₀	R_{17}^{3}	17,18-bond
Chl a	$-CH = CH_2$	Me	Et	double	Me	-COO-CH ₃	Н	phytyl	single
Chl b	$-CH = CH_2$	-CHO	Et	double	Me	$-COO-CH_3$	Н	phytyl	single
Chl c	$-CH = CH_2$	Me	Et	double	Me	-COO-CH ₃	Н	Н	double
Chl d	-CHO	Me	Et	double	Me	-COO-CH ₃	Н	phytyl	single
BChl a	-CO-CH ₃	Me	Et	single	Me	-COO-CH ₃	Н	phytyl	single
BChl b	-CO-CH ₃	Me	= CH-CH ₃	single	Me	$-COO-CH_3$	Η	phytyl	single
BChl c	$\text{-}CHOH\text{-}CH_3$	Me	Et, Pr, iB	double	Me, Et	Н	Me	famesyl	single
BChl d	$-CHOH-CH_3$	Me	Et, Pr, iB	double	Me, Et	Н	Н	famesyl	single
BChl e	$\textbf{-CHOH-CH}_3$	-CHO	Et, Pr, iB	double	Et	Н	Me	famesyl	single
$\operatorname{BChl} g$	$-CH = CH_2$	Me	= CH-CH ₃	single	Me	$-COO-CH_3$	Н	famesyl	single

Fig. 1. Chemical structures of representative chlorophylls (*a*, *b*, *c*₁, and *d*) and bacteriochlorophylls (*a*, *b*, c_{1-3} , d_{1-3} , e_{1-3} , e_{1-

strongly suggested that diazotrophic (i.e., N₂-fixing) cyanobacteria were significant primary producers during the formation of certain petroleum source rocks.

In this review, we focus on the topic how the carbon isotopic composition of chlorophylls is controlled in the photoautotrophic cell, a fundamental concern when interpreting the isotope record of tetrapyrrole structures in natural materials including sediments. Furthermore, we demonstrate that the intramolecular carbon isotopic compositions of chlorophylls and their derivatives, such as sedimentary porphyrins (sometimes called geoporphyrins, petroporphyrins, or metalloalkylporphyrins), can provide critical information and profound insights into biogeochemical processes.

Chlorophylls and bacteriochlorophylls: structures and distribution

We begin with a brief overview of the chemical structures and natural distributions of



Fig. 2. The numbering scheme for chlorophyll *a* proposed by the International Union of Pure and Applied Chemistry (IUPAC). The same basic numbering scheme applies to all chlorophylls and bacteriochlorophylls.

chlorophylls, an important group of tetrapyrroles (i.e., compounds containing four pyrroles or nitrogen-containing five-membered rings) produced in the cell. The chlorophylls vary widely in chemical structure, from chlorophyll *a* to *d*, and from bacteriochlorophyll *a* to *g* (Fig. 1). With further variations within each of these chloropigments, nearly 100 chlorophylls are known today. In Fig. 2, we illustrate the atom numbering schemes for chlorophylls based on that introduced by International Union of Pure and Applied Chemistry (IUPAC) in 1979. The five rings of chlorophylls and bacteriochlorophylls are lettered A – E, and the carbon atoms of the macrocycle are numbered clockwise, beginning with ring A. In most cases, a magnesium ion is at the center of the molecule.

Among the various chlorophylls, chlorophyll a is found in all known eukaryotic, oxygenic photoautotrophs (Table 1), serving as the primary light-harvesting antenna pigment as well as a reaction center for capturing solar energy. Currently, the only known exception to this rule is the cyanobacterium *Acaryochloris marina*, which produces chlorophyll d as its sole chloropigment (Miyashita et al. 1996). In most marine microalgae, 0.25–5% of cellular carbon is associated with chlorophyll a (Geider 1987).

Chlorophyll *b* is the second-most important chlorophyll in terrestrial environments. Chlorophyll *b* differs from chlorophyll *a* only by one of the functional groups at the C– 8 position (a -CHO group in place of a CH₃-group, Fig. 1). In both chlorophylls *a* and *b*, a 20-carbon diterpenoid alcohol group (phytol) is esterified to the C–17 propionic acid attached to ring D. Higher plants and green algae contain chlorophyll *a* and chlorophyll *b* at a ratio of ~3: 1 (Table 1). Chlorophyll *b* acts as an accessory pigment that participates directly in the light reactions of photosynthesis.

Chlorophyll c homologues (i.e., chlorophylls c_1 , c_2 , and c_3) structurally differ from

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Table 1	Distribution of chlorophylls and bacteriochlorophylls in photoautotrophs (modified from
	Blankenship (2004))

Crown	Chlorophyll				Bacteriochlorophyll					
Gloup	а	b	С	d	а	b	С	d	е	g
Cyanobacteria	+			_						
Green algae	+	+								
Diatoms	+		+							
Brown algae	+		+							
Dinoflagellate	+		+							
Cryptomonads	+		+							
Red algae	+									
Higher plants	+	+								
Purple bacteria					+	+				
Green sulfur bacteria					-		+	+	+	
Green non-sulfur bacteria					-		+	+	+	
Heliobacteria										+

-Limited distribution

other chlorophylls in that they have a fully unsaturated, symmetric porphyrin nucleus rather than a chlorin nucleus (e.g., Zapata et al. 2006). These chlorophylls have propenoic acid at the C–17 position, instead of the phytol esterified propionic acid side chain of the other chlorophylls (Fig. 1). Chlorophyll *c* occurs in the primary aquatic photoautotrophs, including diatoms, dinoflagellates, brown algae, and haptophytes, accompanying chlorophyll *a* (Table 1).

Chlorophyll *d* was originally discovered in red macroalgae over 60 years ago, as the fourth type of chlorophyll (Manning and Strain 1943). The molecular structure of chlorophyll *d* is similar to chlorophyll *a*; a vinyl group at the C-3 position of chlorophyll *a* is replaced with a formyl group (Fig. 1). However, it was long erroneously considered a laboratory artifact. In 1996, it was re-discovered in the symbiotic cyanobacterium *Acaryochloris marina* from coralline (Miyashita et al. 1996). Recently, Kashiyama et al. (2008c) reported it in various natural aquatic environments, including freshwater and hypersaline lakes, strongly suggesting an ubiquitous distribution in the aquatic environment.

Bacteriochlorophylls are found in prokaryotic photosynthetic bacteria and heliobacteria. These photoautotrophs conduct anoxygenic photosynthesis; they use hydrogen sulfide rather than water as an electron donor, and thus produce elemental sulfur (S⁰) rather than molecular oxygen (O₂) as the byproduct of photosynthesis. Purple bacteria synthesize either bacteriochlorophyll *a* or *b* (Table 1). These bacteriochlorophylls are different from chlorophylls in that they have a bacteriochlorin rather than chlorin nucleus (Fig. 1). Furthermore, they contain a C₂₀ phytol side chain rather than a C₁₅ farnesol.

Bacteriochlorophylls c, d, and e are produced only by green photosynthetic bacteria, all

of which are strictly anaerobic (Table 1). Although they are termed "bacteriochlorophylls", their ring B contains a C-7-C-8 double bond as in chlorophylls; thus, these pigments are actually chlorins rather than bacteriochlorins (Fig. 1). Bacteriochlorophylls c and e have a methyl substituent at the C-20 methine bridge carbon (Glaeser et al. 2002). Bacteriochlorophyll g has a bacteriochlorin nucleus and is produced only by strictly anaerobic heliobacteria (Brockmann and Lipinski 1983).

Biosynthesis of tetrapyrroles in the cell

It is essential to comprehend the biosynthetic processes of chlorophylls in the cell for proper understanding of intramolecular isotopic signatures. The biosynthetic pathways of chlorophylls together with the bifurcation points for the syntheses of other tetrapyrroles including hemes, bilins, and corrins are outlined in Fig. 3. The biosynthetic processes of these tetrapyrrole structures in the cell have been intensively studied for more than 60 years. It has been demonstrated that 5-aminolevulinic acid or δ -aminolevulinic acid (ALA, Fig. 3) is a universal precursor in the biosynthesis of all tetrapyrrole compounds produced in the cell (e.g., Kikuchi et al. 1958; Beale 1993; Porra 1997).

There are two distinct mechanisms of ALA formation (Fig. 3). In mammals, yeasts, and fungi, ALA is formed from glycine and succinyl-CoA in a one-step condensation reaction catalyzed by the enzyme ALA synthase (ALAS). Because of the pioneering studies of David Shemin, who ingested ¹⁵N-labeled glycine to determine the half-life of hemoglobin (Shemin and Rittenberg 1946), this pathway is often called the "Shemin pathway". In contrast, in plants, algae, and most bacteria, ALA is synthesized from L-glutamic acid in multi-enzyme-catalyzed reactions (Fig. 3; e.g., Beale et al. 1975; Meller et al. 1975; Beale 1995). Glutamic acid is activated to glutamyl-tRNA in the first step, is then reduced to glutamate–1-semialdehyde (GSA), and subsequently forms ALA by GSA aminotransferase. This unique pathway is traditionally referred to as the "C5 pathway". However, it has been demonstrated that in some photosynthetic organisms including the *a* subgroup of purple bacteria and *Euglena*, ALA is known to be a rate-limiting step in the biosynthesis of tetrapyrroles. The activities of ALAS and GSA aminotransferase are strictly regulated by feedback inhibition.

In the Shemin pathway, the carbon skeleton of ALA is inherited from both succinyl-CoA and glycine; carbon atoms at the C-1 to C-4 positions of ALA derive from the C-1 to C-4 carbons of succinyl-CoA, whereas the C-5 position of ALA is derived from the C-2 position of glycine (Fig. 3). In contrast, the carbon skeleton of ALA produced through the C5 pathway is inherited only from glutamic acid. In this case, the carbons at positions C-1 to C-5 of ALA derive from the C-5 to C-1 positions of glutamic acid. In the final reaction of the C5 pathway, GSA is the only required substrate and ALA is the only product. Therefore, the stable carbon isotopic compositions at each of the positions in ALA should reflect those



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Fig. 3. Overview of the biosynthetic pathways of chlorophylls and hemes.

of glutamic acid.

Once ALA is formed, two molecules of ALA condense to form porphobilinogen in a process catalyzed by ALA dehydratase (Fig. 3). Four molecules of porphobilinogen are further condense to form tetrapyrrole uroporphyrinogen III (Fig. 3). Overall, eight molecules of ALA are condensed to form the tetrapyrrole structure of chlorophylls. Uroporphyrinogen III then undergoes a series of decarboxylation and reduction reactions, which are followed by insertion of the metal ion (i.e., Mg^{2+} , Fe^{2+} , or Zn^{2+}) into the tetrapyrrole ring to form protoporphyrin IX (Fig. 3). The biosynthetic pathway of Fechelated heme branches at this point. In chlorophyll synthesis, metallation is followed by construction of the isocyclic ring E, reduction of ring D, and attachment of the phytyl (or farnesyl) side chain (Fig. 1).

The same basic pathway is followed for synthesis of all types of chlorophylls and bacteriochlorophylls. The biosynthetic processes of the various chlorophylls branch in the final stages of chlorophyll formation. For example, chlorophyll *b* is synthesized by oxidation of the methyl group at the C-7 position to give a formyl group.

In the bacteriochlorophyll *a* synthesis, the vinyl group at the C–3 position is converted to an acetyl group in the final step. However, this process does not cleave the carbon skeleton; thus, the carbon isotopic composition of bacteriochlorophyllide *a* is the same as that of the immediate precursor. In the synthesis of the bacteriochlorophyll *e* homologues, additional carbon atoms are added to the C–8² and C–20 positions in the final steps (Senge and Smith 1995). It has been demonstrated that these extra carbon atoms in the bacteriochlorophylls *e* are derived from the methyl group of S-adenosylmethionine, a metabolic product of methionine.

Intramolecular carbon isotopic compositions of chlorophylls

The biosynthetic pathways of chlorophyll synthesis described above indicate that the five carbon atoms in glutamic acid are not equally incorporated into the chlorophyll nucleus. For a chlorophyll *a* nucleus (i.e., chlorophyllide *a*), eight carbon atoms from positions C -1 through C-4 in glutamic acid are introduced, whereas only two carbon atoms from the C-5 position of glutamic acid are incorporated (Fig. 4). We should note that the methoxy group carbon (-OCH₃) at the C-13² position originates from a methyl group (me) derived from the S-adenosylmethionine pool. Overall, in a mechanistic sense, the carbon isotopic composition of the chlorophyll *a* nucleus (chlide *a*) can be expressed by the following equation (Ohkouchi et al. 2008):

$${}^{13}\delta_{\text{chlide }a} = (8^{13}\delta_{\text{C}-1} + 8^{13}\delta_{\text{C}-2} + 8^{13}\delta_{\text{C}-3} + 8^{13}\delta_{\text{C}-4} + 2^{13}\delta_{\text{C}-5} + {}^{13}\delta_{\text{me}})/35 \tag{1}$$

Although the methyl group derived from the S-adenosylmethionine pool may be somewhat depleted in ¹³C (Weilacher et al. 1996), it should not significantly affect the overall isotopic composition, because this carbon contributes only 2.8% (1/35) to the chlorophyll *a* nucleus. In all of the chlorophylls and bacteriochlorophylls, eight carbon atoms from positions C–1 through C–4 in glutamic acid are introduced, whereas only one or two carbons from the C–5 position in glutamic acid are incorporated.

In the plastids of photoautotrophic cells, L-glutamic acid is produced by transfer of the amide group of L-glutamine to 2-oxoglutarate, an important metabolite in the citric acid cycle (i.e., tricarboxylic acid cycle or TCA cycle), catalyzed by the enzyme glutamate





Fig. 4. Relationships between the carbon positions in glutamate, 5-aminolevulinate (ALA), and chlorophyll *a*. Pathways of glutamate biosynthesis via the citric acid cycle are also shown (modified from Ohkouchi et al. 2008).

synthase as part of the GS-GOGAT cycle (Coruzzi and Last 2000). In unusually nitrogenenriched environments (>100 mM NH₃), glutamic acid is also synthesized directly from 2-oxoglutarate by incorporation of ammonium by the enzyme glutamate dehydrogenase. In both cases, the carbon skeleton of glutamic acid is inherited from 2-oxoglutarate.

The carbon atoms at the C-4 and C-5 positions in glutamic acid originate from the

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methyl and carboxyl groups of acetyl-CoA, respectively. In the citric acid cycle, acetyl-CoA condenses with the four-carbon compound oxaloacetate to form citric acid (Fig. 4; Hayes 2001). 2-Oxoglutarate is formed from citrate by decarboxylation, which does not involve the carbons that originated from acetyl-CoA introduced immediately prior to this step (Fig. 4). Previous studies have indicated that the carboxyl group of acetyl-CoA is substantially depleted in ¹³C relative to the methyl group due to isotopic fractionation associated with oxidation of pyruvate to acetyl-CoA by the pyruvate dehydrogenase complex (O'Leary 1976; DeNiro and Epstein 1977; Monson and Hayes 1982; Hayes 1993).

The relatively small contribution of the carbon at the C–5 position in glutamic acid to the chlorophyll nucleus could result in its isotopic composition being somewhat enriched in ¹³C relative to the mean isotopic composition of glutamic acid. Because the isotopic fractionation associated with oxidation of pyruvate to acetyl-CoA is purely a kinetic process, the magnitude of the ¹³C depletion in the carboxyl carbon may vary widely (Monson and Hayes 1982; Melzer and Schmidt 1987). In case of a 10‰ ¹³C-depletion of the carboxyl group relative to the methyl group of acetyl-CoA, the chlorophyll nucleus can be calculated to be ~0.6‰ enriched in ¹³C relative to glutamic acid (or ALA).

In the citric acid cycle of photoautotrophic cells, not only acetyl-CoA but also oxaloacetate derived from irreversible β -carboxylation of phosphoenolpyruvate (PEP) substantially compensate for the metabolites removed from the citric acid cycle (Fig. 4; Sakata et al. 1997). In β -carboxylation of PEP, a bicarbonate carbon (HCO₃⁻) binds to the C-4 position of oxaloacetate (corresponding to the C-1 position of 2-oxoglutarate). Typically, HCO₃⁻ is enriched relative to CO₂ dissolved in seawater by ~10.5‰ at 10° C and ~13.3‰ at 25°C at equilibrium (Mook et al. 1974). Therefore, the carbon isotopic composition of the C-1 position of glutamic acid (i.e., the carboxyl group carbon) should be *significantly* enriched in ¹³C if β -carboxylation is an important process in the photoautotrophic cell.

In their seminal paper, Abelson and Hoering (1961) reported for the first time the isotopic composition of the C–1 carbon in glutamic acid in photosynthetic microorganisms cultured in the laboratory. They found strong enrichment in ¹³C (+10.9 to +32.2‰), which was highly variable relative to the mean isotopic composition of the other carbon atoms in glutamic acid (i.e., C–2 through C–5), although their experiments were less precise and based on less carefully cultured organisms than current methods. The above theoretical considerations and experimental observations suggest that the isotopic composition of the tetrapyrrole) are enriched in ¹³C relative to the mean carbon isotopic composition of the entire chlorophyll nucleus, because they originate from the C–1 carbon in glutamic acid.

Geoporphyrins, geological derivatives of chlorophylls

Because chlorophylls are degraded either chemically or microbiologically, various





Fig. 5. Degradation (i.e., diagenetic) pathway of chlorophyll *a* in the water column and sediments.

chlorophyll derivatives have been widely observed in the water column and sediments. They include pheophytins which lack the magnesium ion in the center of the tetrapyrrole nucleus of chlorophylls (Fig. 5). Furthermore, pyropheophytins and pyropheophorbides have been reported in many sediments (Fig. 5, e.g., Villanueva et al. 1994; Harris et al.





1995; Nakajima et al. 2003). These "biological tetrapyrroles" are ultimately transformed to geoporphyrins, metallo-porphyrins substituted with various alkyl chains (Baker and Louda 1986; Eckardt et al. 1989; Keely et al. 1990; Kashiyama et al. 2010).

Geoporphyrins are "geological derivatives" of cellular tetrapyrroles including chlorophylls and hemes, and are sometimes termed either sedimentary porphyrins or petroporphyrins. Their origins were first postulated in the 1930s by Alfred Treibs from Hans Fischer's Munich school, who isolated deoxophylloerythroetioporphyrin (DPEP) and etioporphyrin (ETIO; Fig. 6) from a wide range of petroleum, shales, bitumens, and coal. Based on structural similarities, Treibs (1934, 1936) suggested that the DPEP and ETIO porphyrins originated from chlorophylls and hemes, respectively. Since then, the nomenclature, chemistry, synthesis, and distribution of geoporphyrins in nature have been summarized by several authors (Baker and Palmer 1978; Baker and Louda 1986; Callot and Ocampo 2000; Keely 2006). The purpose of this section is not to thoroughly review geoporphyrins but, rather, to give the readers some insight into the structure and nomenclature of geoporphyrins.

Since the Treibs hypothesis was proposed more than 70 years ago, over 70 species of geoporphyrins have been reported in various sediments, oils, and coals (Callot and Ocampo 2000). Sedimentary porphyrins occur mainly as VO(II) and Ni(II) complexes, although other metal complexes such as Cu(II) and Fe(III) have been frequently observed (Eckardt et al. 1989).

The structural variation in sedimentary porphyrins is far greater than in the original chlorophylls and hemes. In most geological samples, DPEP is most abundant (Baker et al. 1968). Because its exocyclic ring structure is closely related to the chlorophylls, it is generally accepted that DPEP must derive primarily from chlorophylls. Theoretically, all chlorophylls *a-d* and even bacteriochlorophyll *a* are potential precursors of sedimentary DPEP. However, it is likely that chlorophyll *a* is the dominant precursor, because it is far more abundant and widespread relative to other chlorophylls in the natural aquatic environment. In contrast, 17-nor-DPEP, a DPEP-type porphyrin lacking an alkyl chain at the C-17 position (Fig. 6), was postulated to be a derivative of chlorophyll *c*, because cleavage of the vinyl group at the C-17 position tends to occur after decarboxylation of the free acrylic chain (Callot et al. 1990). 17-nor-DPEP is widely distributed in geological samples (e.g., Huseby et al. 1996).

Isotopic composition of methine bridge carbon as an indicator of β -carboxylation

The above theoretical considerations suggest that the methine-bridge carbons in chlorophylls should be enriched in ¹³C relative to the other carbons in proportion to the β -carboxylase activity in the photoautotrophic cell. In other words, the isotopomeric (i.e., intramolecular or site-specific isotopic composition) information on these porphyrins potentially provides physiological information on photoautotrophs in the geological past.

In the present study, we conducted chemical degradation of tetrapyrroles with chromium oxide (Quirke et al. 1980; Grice et al. 1996; Chikaraishi et al. 2008) to investigate their intramolecular isotopic compositions. Chromic acid oxidation of porphyrins produces several types of maleimides (1*H*-pyrrole-2,5-diones) as well as carbon dioxide from the methine bridge carbons (Fig. 7; Chikaraishi et al. 2008). Therefore, the mean isotopic compositions of the four methine bridge carbons of the porphyrins can be estimated using the following mass balance equation, including the isotopic compositions of the bulk tetrapyrroles and the maleimides, the oxidation products of the tetrapyrroles:

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Fig. 7. Chromic acid degradation of DPEP and 17-nor-DPEP to form maleimides.

$${}^{13}\delta_{\text{methine}} = (32^{13}\delta_{\text{porphyrin}} - 28^{13}\delta_{\text{maleimide}})/4$$
$$= 8^{13}\delta_{\text{porphyrin}} - 7^{13}\delta_{\text{maleimide}}$$

(2)

We evaluated whether isotopic fractionation is associated with chromic acid oxidation in the laboratory experiments. However, we did not observe variations in the estimated mean isotopic compositions of the methine bridge carbons with the maleimide yield of the chromic acid oxidation experiments. Although a more rigorous evaluation is required in the future, based on this evidence, we tentatively concluded that the isotopic fractionation associated with C-C bond cleavage during chromium oxide degradation of tetrapyrrole structures can be neglected.

We applied the above approach to sedimentary porphyrins isolated from the Onnagawa Formation, which was formed in the middle Miocene in a semi-enclosed basin and is distributed in northeast Japan. The Onnagawa Formation mainly consists of diatomaceous deposits with a total organic carbon content of up to 5% by weight; detailed lithology and stratigraphy are described in Tada (1991). From our rock collection, we selected two samples (BA138a and BA138b) for this study (Kashiyama et al. 2008b).

Detailed analytical procedures are described in Kashiyama et al. (2007a) and Chikaraishi et al. (2008). Briefly, the sediments were Soxhlet-extracted with chloroform/methanol (70: 30, v/v). The total extracts were separated with silica gel column chromatography to isolate vanadyl (VO)-porphyrins. Using a two-step HPLC method, we carefully isolated and purified VO-DPEP and VO-17-nor-DPEP from the VO-porphyrins using a computer-controlled fraction collector. Structural assignments for these purified porphyrins were achieved using a combination of absorption spectrometry with a photodiode array detector,

 Table 2
 Carbon isotopic compositions (‰) of bulk porphyrins from two sediment samples from

 Onnagawa Formation and maleimides produced by their chromic acid oxidation. Estimated carbon isotopic compositions of methine bridge carbon are also shown in the bottom of the table.

	B	A138a	BA138b
	VO DPEP	VO 17-nor-DPEP	VO DPEP
Porphyrin	- 17.1	- 16.7	- 17.9
2-Eethyl-3-methyl-maleimide	- 16.4	- 16.4	-17.1
2-Methyl-maleimide	n.d.	- 15.7	n.d.
Methine bridge carbon	-22.0	-22.3	-23.5

molecular ion monitoring by single-stage quadrupole mass spectrometry, and single-crystal X-ray diffraction patterns (Kashiyama et al. 2007b, 2008a). Carbon isotopic compositions of individual bulk porphyrins were determined using an elemental analyzer/isotope ratio mass spectrometry (EA/IRMS) with improved sensitivity (Ohkouchi et al. 2005; Ogawa et al. 2010). We have previously reported the detailed carbon isotopic compositions of VO-DPEP and VO-17-nor-DPEP for these samples in Kashiyama et al. (2008b) and they are also shown in Table 2.

We conducted chromic acid oxidation on aliquots of these purified porphyrins to form maleimides. The structural assignments of the maleimides produced by chromic acid oxidation were achieved by gas chromatography/mass spectrometry and comparison with synthetic maleimide standards (GC/MS; Chikaraishi et al. 2008). Carbon isotopic compositions of individual maleimides were determined using gas chromatography/ combustion/isotope ratio mass spectrometry (GC/C/IRMS; Chikaraishi et al. 2008).

After chemical degradation of VO-DPEP, we observed 2-ethyl-3-methyl maleimide, a degradation product of rings A, B, and D of the tetrapyrrole structure (Fig. 7). In contrast, chromic acid degradation of VO-17-nor-DPEP produced both 2-ethyl-3-methyl maleimide and 2-methyl maleimide; the former is a product of rings A and B, and the latter is a product of ring D of the tetrapyrrole structure (Fig. 7). The measured carbon isotopic composition of 2-ethyl-3-methyl maleimide from VO-DPEP was -16.4‰ for sample BA138a and - 17.1‰ for sample BA138b. Furthermore, those of 2-ethyl-3-methyl maleimide and 2-methyl maleimide from VO-17-nor-DPEP were -16.4‰ and -15.7‰, respectively, for sample BA138a. We were not able to conduct maleimide analysis for VO-17-nor-DPEP from BA138b due to insufficient sample.

Using equation 2 above, we estimated the mean isotopic composition of the methine bridge carbons for these porphyrins. As shown in Table 2, the mean isotopic composition of the methine bridge carbons was -22.0% for VO-DPEP and -22.3% for VO-17-nor-DPEP from BA138a, whereas that of VO-DPEP from BA138b was -23.5%. The uncertainty of these estimated isotopic numbers derives not only from instrumental analytical errors (normally $\sim 0.3\%$ (1 σ) for EA/IRMS and $\sim 0.5\%$ (1 σ) for GC/C/IRMS), but also from

propagation of these errors. Nevertheless, the notable ¹³C enrichment of the methinebridge carbons should be robust, and we conclude that the methine bridge carbons were substantially (5–6‰) depleted in ¹³C relative to the mean bulk isotopic composition of the original porphyrins.

This isotopomeric evidence clearly suggests that β -carboxylation was not a significant physiological process for photoautotrophs during deposition of the Onnagawa Formation. It is well known that β -carboxylation is a widespread process in C4 and CAM plants (e.g., Hatch and Slack 1970). PEP carboxylase, an enzyme catalyzing this process, has also been widely observed in other organisms including cyanobacteria. Although it is still uncertain which factor controls β -carboxylation activity, 10–20% of the carbon in the cyanobacterial cell has been estimated to be assimilated through β -carboxylation (e.g., Coleman and Colman 1981; Owttrim and Colman 1986). Therefore, our isotopomeric observations in this study seem to be inconsistent with previous results (Kashiyama et al. 2008b). Kashiyama et al. (2008b) estimated an average δ^{15} N value for the entire photoautotrophic community of -2 to +1‰ based on δ^{15} N values for DPEP (-6.9 to -3.6‰, n=7), considering empirical isotopic data indicating that the tetrapyrrole nuclei of chloropigments are depleted in ¹⁵N by \sim 5% relative to the cells. This finding suggests that N₂-fixing cyanobacteria were important primary producers during the deposition of the Onnagawa Formation. Alternatively, as also pointed out by Kashiyama et al. (2008b), endosymbiosis between diatoms and cyanobacteria may be a mechanism that explains not only the isotopomeric evidence reported here but also the carbon and nitrogen isotopic compositions of the individual (bulk) porphyrins from these sediments.

Concluding Remarks

We investigated the intramolecular isotopic compositions of tetrapyrrole nuclei of chlorophylls with an emphasis on current knowledge of metabolic processes occurring in photoautotrophic cells. Because their chemical structures, synthetic pathways, and regulatory processes have been intensively studied, chlorophylls are ideal candidates for understanding the carbon cycle in surface water environments. Fortunately, geoporphyrins, geological products of chlorophyll nuclei, are well preserved in the sedimentary record, and thus their isotopic signatures can provide robust evidence regarding the surface water environment in the geologic past.

As suggested by this study, there are strikingly different isotopic ratios among carbon positions within a single molecule depending on the metabolic pathway, and thus, substantial information is recorded in intramolecular isotopic signatures. Therefore, intramolecular carbon isotopic compositions, as well as bulk isotopic compositions, are potentially quite useful for investigating the detailed processes associated with carbon cycling in the euphotic zone of the water column on geological timescales. Nevertheless, there still is not enough evidence strongly supporting the mechanistic approach to interpreting intramolecular isotopic compositions due to our current lack of information. In the authors' view, it is required to rigorously evaluate and understand the relationships between metabolic processes and isotopic fingerprinting, that will undoubtedly bring a bright future to the study of isotopomers of organic molecules.

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