

## FATTY ACIDS OF GEOCHEMICAL SIGNIFICANCE IN MICROSCOPIC ALGAE\*

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**Abstract**—Fatty acids ranging from 14:0 to 18:4 were found in eleven different species of green and blue-green algae. In general, the unsaturated  $C_{18}$  fatty acids predominate in all the species of green algae analyzed while with some exceptions the  $C_{16}$  saturated acids are predominant in the blue-green algae. By the nature of their gas chromatographic patterns, there appears to be no direct correlation between the fatty acids and the hydrocarbons within the same species of algae. The fatty acids are distributed in a narrower molecular weight range ( $C_{14}$ – $C_{18}$ ) than the hydrocarbons ( $C_{15}$ – $C_{33}$ ) and do not show any significant qualitative variations between species. All assignments of structural identities are supported by gas chromatographic and mass spectrometric data.

### INTRODUCTION

DIFFERENT extant forms of ancient algae have been shown to contain hydrocarbon distributions similar to those found in sediments (see previous paper<sup>1</sup> for references). These same algae usually contain considerably greater amounts of fatty acids than aliphatic hydrocarbons. However, contrary to the hydrocarbons, these fatty acids are without exception more restricted in their molecular weight range.<sup>2–5</sup>

The geological significance of fatty acids in general has been discussed,<sup>4,6</sup> as has been their possible relationship to hydrocarbon production in sediments and in petroleum.<sup>6,9</sup> Algal fatty acids in particular, whether derived from either prokaryotes or eukaryotes, do not differ much in their distribution or relative amounts. It has been shown that changing environmental conditions will alter the fatty acid distribution of micro-organisms markedly;<sup>9,10</sup> however, these environmental changes do not seem to affect the molecular weight range.

Although the fatty acid compositions of several blue-green algae are rather simple in terms of their molecular structure, variations among different species have been noted.

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Also, a certain degree of correlation has been shown to exist between the degree of unsaturation of fatty acids in blue-green algae and their morphological complexity.<sup>2</sup> However, no such correlations apply to the aliphatic hydrocarbons.

Well-characterized fossil algae have been shown to be present in relatively large quantities in certain recent sediments.<sup>11-13</sup> Likewise, ancient sediments have been shown to contain different microscopic bodies whose morphology is analogous to that of several present-day species of blue-green and green algae.<sup>14,15</sup> To better understand the possible relationship of fatty acids to hydrocarbons as well as the origin of some of the lipids in sediments, the fatty acids of contemporary counterparts of several fossil algae have been studied in our laboratory.

## RESULTS

Table 1 gives the fatty acid ranges of the algae analyzed (see Table 1 in previous paper<sup>1</sup> for the sediments in which the fossil forms are found). The widest range is found in the green

TABLE 1. FATTY ACIDS FROM CONTEMPORARY REPRESENTATIVES OF ALGAE FOUND IN SEDIMENTS\*

Organism	Fatty acid Me esters	
	Range	Major
<i>Coelastrum microsporum</i>	14:0-18:4	18:1
<i>Chlorella pyrenoidosa</i>	14:0-18:3	18:3
<i>Scenedesmus quadricauda</i>	16:0-18:3	18:3
<i>Tetradron sp.</i>	16:1-18:4	18:1
<i>Anacystis cyanea</i>	15:0-18:3	16:0, 16:1
<i>A. nidulans</i>	14:0-18:1	16:0, 16:1
<i>Spirulina platensis</i>	14:0-18:3	16:0
<i>Lyngbya aestuarii</i>	16:0-18:3	16:0
<i>Nostoc sp.</i>	14:0-18:3	16:0
<i>Chroococcus turgidus</i>	16:0-18:3	16:0, 18:3
<i>Anacystis montana</i>	16:0-18:3	18:1

\* Refer to Table 2 in the preceeding paper for classification by genera and occurrence in sediments.

algae *Coelastrum microsporum*, 14:0 to 18:4. All of the algae analyzed fall within this range. Table 1 also shows the predominant fatty acids of each organism. In general, the green algae have a C<sub>18</sub> unsaturate as their major fatty acid, while all of the blue-green algae, except for *Anacystis montana*, have either a 16:0 or 16:1 as their major component. The relative percentage composition for each individual fatty acid is given in Table 2.

All patterns appear similar and are in close agreement with other data on micro-organisms presented in the literature.<sup>2-5,8,9,16</sup> Fatty acid distributions of a number of cyanophycean algae have been reported.<sup>2,4,8</sup> The major variations in the fatty acid patterns relate to the

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relative amounts and degree of saturation. The identities of all the fatty acid methyl esters reported in Tables 1 and 2 have also been confirmed by mass spectrometry. Their mass spectrometric fragmentation patterns have already been discussed in the literature.<sup>3,16</sup> Considered as a whole, the C<sub>16</sub> acids appear to predominate in some species of blue-green algae, according to the computed ratio of total C<sub>16</sub>/C<sub>18</sub> fatty acids given in Table 2. The variations are great, however. From the relatively very high value for *A. nidulans* to the even and low values for *Lyngbya aestauri*, *Chroococcus turgidus* and *A. montana*. Conversely, the ratio of C<sub>16</sub>/C<sub>18</sub> fatty acids in the green algae (Table 2), shows without exception the C<sub>18</sub> fatty acids to be predominant.

TABLE 2. RELATIVE PERCENT FATTY ACID COMPOSITION OF SEVERAL MICROSCOPIC ALGAE\*

Fatty acid Me esters	Green algae				Blue-green algae						
	Chlor. pyr.	Tetr. sp.	Coel. micro.	Scen. quad.	Ana. nid.	Ana. cyan.	Spir. plat.	Nost. sp.	Lyng. aest.	Chroo. turg.	Ana. mont.
14:0	1.6		0.7		3.0		1.9	2.3			
14:1					4.0						
15:0						3.0					
16:0	21.4	18.3	13.1	13.0	30.7	23.0	45.0	36.3	25.8	28.2	21.3
16:1	2.0	2.6	1.3	4.0	33.3	20.0	6.0	10.0	15.4	12.3	3.1
16:2	5.8	2.7	4.6					1.0	3.2	3.8	
17:0					1.2						
17:1							2.0				
18:0		1.7			0.4	2.0		3.0			
18:1†	16.7	33.5	30.0	18.0	3.6	7.0	10.1	19.8	8.3	7.3	25.3
18:1‡			7.0	14.0							
18:2	18.7	11.1	13.0	7.0		10.0	16.0	8.0	16.8	16.4	17.8
18:3	26.4	18.3	16.0	29.0		5.0	13.0	6.5	18.2	25.7	19.4
18:4		6.3	7.0								
Total C <sub>16</sub> /C <sub>18</sub>	0.47	0.33	0.26	0.25	16.0	1.92	1.31	1.27	1.02	0.92	0.39

See Table 1 for species.

\* Differences to 100% made up by unidentified compounds.

† 9-*cis*-Octadecenoic acid (oleic).

‡ 9-*trans*-Octadecenoic acid (elaidic).

All of the algae except *A. nidulans* have a ratio of saturated to unsaturated C<sub>16</sub> in which the saturated form predominates, while the reverse is true for the C<sub>18</sub> acids. Also, the octadecanoic acids show somewhat larger amounts and variety of polyunsaturated forms in all of the algae analyzed, with the exception again of *A. nidulans*.

Both *Scenedesmus quadricauda* and *Coelastrum microsporum* contain two components with different retention times, that show the mass spectral characteristics of a mono unsaturated C<sub>18</sub> acid. By comparing the gas chromatographic peak retention time with that of the authentic *cis* and *trans* isomers of 9-octadecenoid acid (oleic and elaidic) they were respectively identified as *cis* and *trans* C<sub>18:1</sub> isomers. A similar situation applies to the mono unsaturated C<sub>16</sub> acid in the same two species of algae. Only two of the algae analyzed, both green, showed detectable amounts of tetraunsaturated C<sub>18</sub> fatty acids as seen in (Table 2).

## DISCUSSION

The data obtained on algae so far does not suggest a direct biosynthetic relationship between hydrocarbons and fatty acids.<sup>3,8</sup> Rather, if the hydrocarbons in the algal cells arise from the regular fatty acids, a variety of complex intermediate processes must be involved.

For instance, the marked predominance of the *n*-C<sub>17</sub> hydrocarbon (see first paper<sup>1</sup>) could be a point in favor of either of two direct pathways; elongation of the C<sub>16</sub> fatty acids or decarboxylation of the C<sub>18</sub> acids. However, depending on the prevailing mechanism, one would also expect to see substantial amounts of pentadecane, if the C<sub>16</sub> fatty acid were to undergo decarboxylation, or nonadecane if the C<sub>18</sub> acid were to elongate. The absence of significant amounts of these two hydrocarbons implies a more complex mechanism than simple elongation or decarboxylation of corresponding fatty acids. The same would apply to the high molecular weight hydrocarbons found in some species of algae,<sup>1,17,18</sup> since their corresponding fatty acids are absent. In this case, the mechanisms proposed for the condensation of two fatty acids involving the participation of alcohols and ketones or aldehydes<sup>19,20</sup> as intermediates do not seem to explain the facts either. An alternative mechanism would be that of the elongation-decarboxylation complex recently proposed by Kolattukudy.<sup>21</sup>

Considering the fatty acid distributions, any attempt made to correlate the variations in these distributions to phylogenetic positions is in our opinion still premature. Our patterns do not show significant and consistent variations between species or in some cases genera (Tables 1 and 2).

Among the fatty acid patterns there are considerable differences in the degree of saturation of the C<sub>16</sub> and C<sub>18</sub> fatty acids and, as shown by recent literature, changes in growth parameters such as temperature can lead to shifts in the degree of saturation of fatty acids.<sup>9,22</sup> However, in our case all cultures were grown under controlled conditions which would tend to minimize any shifts in degree of saturation. In addition, phylogenetic interpretation of fatty acid patterns should correlate with observed variations in hydrocarbon distributions. We cannot at present decipher any meaningful correspondence along the two lines of evidence. The only marked difference we have noted that could be related to phylogenetic position or morphological complexity is the total C<sub>16</sub>/C<sub>18</sub> ratio which shows (Table 2) the blue-green (prokaryotes) to have a predominance of C<sub>16</sub> fatty acids and the greens (eukaryotes) to have a predominance of C<sub>18</sub> fatty acids. If this were really an indication of morphological complexity the low value for *Anacystis montana* (Table 2) would seem to indicate that this organism is a transitional form<sup>16</sup> between the blue-green and green algae.

In relation to their geochemistry, the specific involvement of algal fatty acids in sediments does not seem to be as well substantiated as that of the aliphatic hydrocarbons (cf. first paper<sup>1</sup>). However, fatty acids are among the most abundant components of living systems and in general have been often regarded as the possible precursors of petroleum and sedimentary hydrocarbons.<sup>6,7</sup>

Normal fatty acids are present in recent as well as Precambrian sediments and considering the large number of algae implicated in these sediments, the data presented here would

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certainly suggest that algae must have contributed in one way or another to the fatty acid distributions found in our lithosphere.<sup>6, 23, 24</sup> However, it is hard to evaluate the extent of their contributions.

The absence of the unsaturated acids in sediments as opposed to their abundance in living algae may be explained by their lower stability to a reducing geological environment. Likewise, the appearance of significant amounts of odd carbon numbered normal fatty acids with geological time, which parallels the appearance of even carbon numbered hydrocarbons, can also be related to diagenetic changes of the originally deposited algal debris.

On the other hand, the fatty acid distributions of contemporary algae fail to account for the presence in sediments of branched-chain acids<sup>23-25</sup> and of acids higher than C<sub>18</sub>.<sup>6, 7, 23, 24</sup> They do not account either for the presence of odd fatty acids in recent muds and soils. Isoprenoid fatty acids, which have been reported in recent sediments,<sup>23, 26, 27</sup> have not been detected in the algae studied in our laboratory. This, however, would not be unexpected since they most likely arise from slow diagenetic degradation of the algal chlorophylls. On the other hand there is one report in the literature on the presence of significant amounts of pristane in a green alga.<sup>8</sup>

Consideration of the contributions of bacteria and higher plants to the fatty acid patterns observed in Tertiary and Cambrian sediments would fill the gap left by algae. Concerning the Precambrian, the scarce data available in the literature appears to support the role of algae as their direct source, except for the C<sub>20</sub> and C<sub>22</sub> acids.<sup>6</sup>

In conclusion, it seems that the correlation of algal fatty acids with the normal fatty acids in sediments, contrary to the hydrocarbons, is less apparent, although nevertheless valid in the C<sub>12</sub>-C<sub>20</sub> range.

## EXPERIMENTAL

The growth parameters, source, and harvesting of the algae are given in the preceding paper. Sample preparations are performed in the same manner as described in Paper I,<sup>1</sup> except that the methanol fraction containing free fatty acids, glycerides and other lipids was taken to dryness and methyl esters were produced by treatment with BF<sub>3</sub>.<sup>28</sup> The fatty acid methyl ester fraction was concentrated by evaporation under N<sub>2</sub> and analyzed by GLC, using a Barber-Coleman 5000 gas chromatograph equipped with a hydrogen flame-ionization detector; and combined gas chromatography-mass spectrometry (LKB-9000 Gas Chromatograph-mass spectrometer).

Separation of the fatty acid methyl esters was achieved on a stainless-steel column (9.6 m × 0.3 cm, i.d.) packed with 15% EGS on 80/100 Chromosorb W AW, N<sub>2</sub> pressure of 703 g/cm<sup>2</sup>, battery setting of ×1 and attenuation of 10. The GLC conditions were designed specifically to detect any fatty acids in the C<sub>12</sub>-C<sub>30</sub> range. Dry weights of algae extracted ranged from 0.29 to 1.0 g.

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