

## Initial Reactions in Anaerobic Alkane Degradation by a Sulfate Reducer, Strain AK-01

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**An alkane-degrading, sulfate-reducing bacterial strain, AK-01, isolated from a petroleum-contaminated sediment was studied to elucidate its mechanism of alkane metabolism. Total cellular fatty acids of AK-01 were predominantly C even when it was grown on C-even alkanes and were predominantly C odd when grown on C-odd alkanes, suggesting that the bacterium anaerobically oxidizes alkanes to fatty acids. Among these fatty acids, some 2-, 4-, and 6-methylated fatty acids were specifically found only when AK-01 was grown on alkanes, and their chain lengths always correlated with those of the alkanes. When [1,2-<sup>13</sup>C<sub>2</sub>]hexadecane or perdeuterated pentadecane was used as the growth substrate, <sup>13</sup>C-labeled 2-Me-16:0, 4-Me-18:0, and 6-Me-20:0 fatty acids or deuterated 2-Me-15:0, 4-Me-17:0, and 6-Me-19:0 fatty acids were recovered, respectively, confirming that these monomethylated fatty acids were alkane derived. Examination of the <sup>13</sup>C-labeled 2-, 4-, and 6-methylated fatty acids by mass spectrometry showed that each of them contained two <sup>13</sup>C atoms, located at the methyl group and the adjacent carbon, thus indicating that the methyl group was the original terminal carbon of the [1,2-<sup>13</sup>C<sub>2</sub>]hexadecane. For perdeuterated pentadecane, the presence of three deuterium atoms, on the methyl group and its adjacent carbon, in each of the deuterated 2-, 4-, and 6-methylated fatty acids further supported the hypothesis that the methyl group was the terminal carbon of the alkane. Thus, exogenous carbon appears to be initially added to an alkane subterminally at the C-2 position such that the original terminal carbon of the alkane becomes a methyl group on the subsequently formed fatty acid. The carbon addition reaction, however, does not appear to be a direct carboxylation of inorganic bicarbonate. A pathway for anaerobic metabolism of alkanes by strain AK-01 is proposed.**

Alkanes are major components of petroleum fuels commonly found in polluted environments, and numerous studies of their biodegradability by microorganisms have been conducted. Most of the earlier studies focused on processes that take place under aerobic conditions (6), in which alkanes are initially oxidized by oxygen to alcohols by monooxygenases, with detailed accounts of the biochemistry and genetics (37, 38). In contrast, while alkane degradation under anaerobic conditions has been reported in early as well as recent literature (2, 7, 9, 27, 32), much less is understood about the mechanism(s) of degradation in the absence of oxygen. Some reports indicated that alkanes could be enzymatically dehydrogenated to alkenes (8, 19), and the reaction may have served as an initial reaction for anaerobic alkane degradation (29). These observed activities, however, could not be shown to support growth under anaerobic conditions (17).

In more recent studies, Aeckersberg et al. isolated and characterized a sulfate-reducing, alkane-degrading strain, Hxd3 (2). We have also recently reported the isolation and characterization of an alkane-degrading strain, AK-01, which is different from Hxd3 phylogenetically (35). Hxd3 appears to anaerobically oxidize alkanes to fatty acids, as evidenced by the influence of alkane carbon chain lengths on its cellular fatty acid composition (3). Other observations also suggest that an

odd-numbered carbon atom(s) may be initially added to or removed from the alkane chain terminus before the fatty acids are formed. On the other hand, evidence suggests that desaturation of the alkane to the corresponding 1-alkene, a hypothetical initial step previously proposed in the literature (29), does not occur during anaerobic alkane oxidation by Hxd3 (1, 3). Another alkane-degrading strain, Pnd3, reported by the same authors (3) also appears to anaerobically oxidize alkanes to fatty acids, though seemingly through a different initial reaction(s) (3, 14). These initial reactions are not yet understood.

In this study, we investigated the initial reaction(s) of alkane degradation by strain AK-01. This strain, which was isolated from a petroleum-contaminated estuarine sediment, is able to oxidize alkanes completely to carbon dioxide, with coupling to sulfate reduction (35). Early evidence also suggests that it anaerobically oxidize alkanes to fatty acids. By using stable-isotope-labeled alkanes as growth substrates, we clearly documented that alkanes are oxidized to fatty acids by this organism. Structural characterizations of the isotope-labeled fatty acids derived from the labeled alkanes demonstrated that exogenous carbon is initially added to the alkane chain subterminally at the C-2 position, with subsequent formation of fatty acids.

### MATERIALS AND METHODS

**Source of organism and culture conditions.** The alkane-degrading bacterial strain AK-01 was isolated from a petroleum-contaminated estuarine sediment collected from the Arthur Kill in New York (35). The medium used for maintaining AK-01 was the same as that described previously (35) except that no yeast extract was added. Liquid alkanes and 1-alkenes used as the substrates were sterilized separately by either autoclaving or filtration (0.22 μm pore size) before use. Long-chain fatty acids (as sodium salts) were prepared asexually as concentrated stock solutions and were autoclaved before being added (while still melted) to the medium. Solid 1-alkanols were autoclaved together with the medium. In most cases, subcultures of AK-01 grown on alkanes for about a

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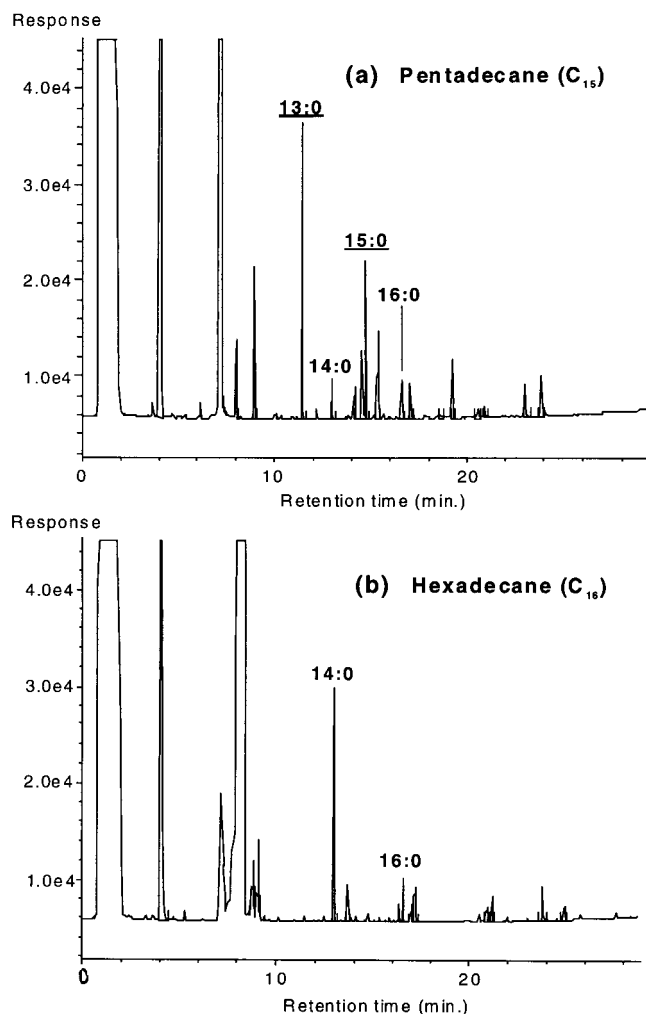


FIG. 1. Gas chromatograms (GC-FID) showing the fatty acid profiles of strain AK-01 grown on pentadecane (a) and hexadecane (b). The n-saturated fatty acids identified by their retention times are annotated. C-odd fatty acids are underlined.

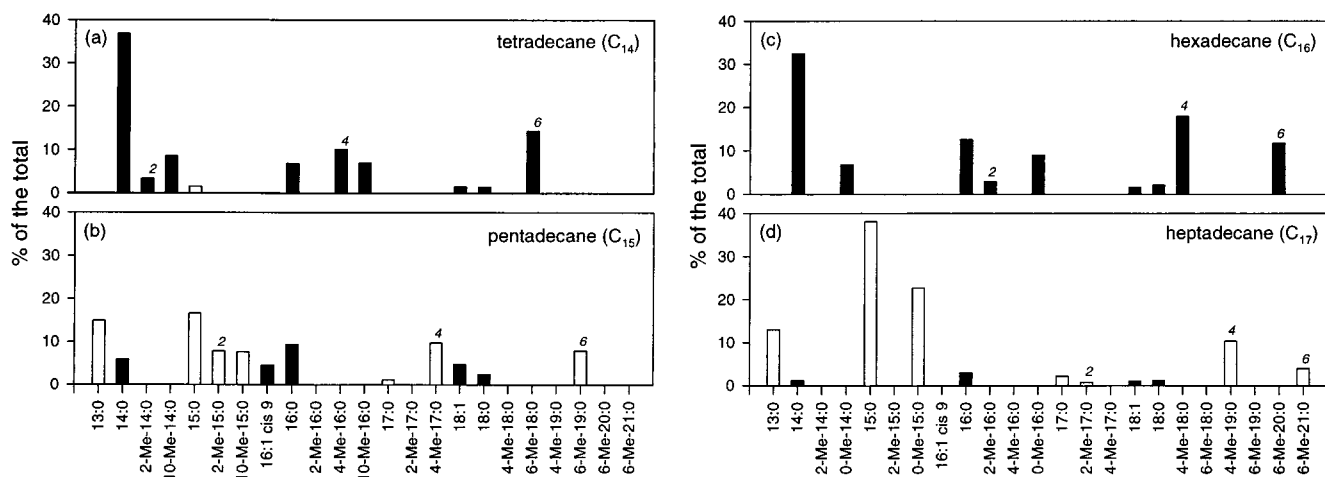


FIG. 2. Relative abundance of the total cellular fatty acids of strain AK-01 grown on tetradecane (a), pentadecane (b), hexadecane (c), and heptadecane (d). Only those which can be identified and comprise more than 1% of the total are shown. C-even fatty acids are represented by black bars, and C-odd fatty acids are represented by white bars. The 2-methyl, 4-methyl, and 6-methyl fatty acids are denoted by the numbers 2, 4, and 6, respectively. (The 2-Me-17:0 fatty acid is also shown, though it constitutes only 0.7% of the total.) No fatty acid was observed in the sterile controls.

month were used as the inocula, which differed in size with the experiment (see details for each experiment below). All cultures were incubated in the dark without shaking.

**Effect of alkanes on total cellular fatty acid composition.** Bottles containing 100 ml of medium were amended with 20  $\mu$ l of tetradecane, pentadecane, hexadecane, or heptadecane (99%; Sigma Chemical Co., St. Louis, Mo.) and inoculated with cultures of AK-01 maintained on hexadecane (5%, vol/vol). For each alkane, two experimental cultures and two sterile controls were set up. After growing for about a month, cultures were subcultured again in medium supplemented with the same alkane. After incubation for 38 to 51 days to reach the stationary phase (growth rates differed depending on the substrate), all cultures were sacrificed for extraction and analysis of total cellular fatty acids.

**Effect of other substrates on total cellular fatty acid composition.** Bottles containing 100 ml of medium were amended with the following substrates: 1-pentadecene (20  $\mu$ l), 1-hexadecene (20  $\mu$ l), 1-pentadecanol (20 mg), 1-hexadecanol (20 mg), pentadecanoate (0.5 mM), and hexadecanoate (0.5 mM). The media were then inoculated with cultures of AK-01 maintained on hexadecane (5%, vol/vol). For each substrate, two experimental cultures and two autoclaved controls were prepared. After incubation for 14 to 51 days to reach stationary phase, the cultures were sacrificed for extraction and analysis of total cellular fatty acids.

**Degradation of [1,2-<sup>13</sup>C<sub>2</sub>]hexadecane.** Cultures were prepared by adding 20  $\mu$ l of [1,2-<sup>13</sup>C<sub>2</sub>]hexadecane (isotopic enrichment, 99%; Isotec Inc., Miamisburg, Ohio) to 100 ml of medium and inoculating with a culture pregrown on pentadecane (5%, vol/vol). Two experimental cultures and one sterile control were prepared. For the sterile control, the inoculum was autoclaved before use. All cultures were incubated in the dark without shaking for 47 days and sacrificed for extraction and analysis of total cellular fatty acids.

**Degradation of perdeuterated pentadecane.** Cultures were prepared by adding 20  $\mu$ l of perdeuterated pentadecane (C<sub>15</sub>D<sub>32</sub>; isotopic enrichment, 98%; Cambridge Isotope Laboratories, Inc., Andover, Mass.) to 100 ml of medium and inoculating with a culture pregrown on hexadecane (5%, vol/vol). Similar to the study described above, two experimental cultures and one sterile control were prepared for the experiment, and the inoculum was autoclaved before use as the sterile control. All cultures were incubated in the dark without shaking for 39 to 47 days and then sacrificed for extraction and analysis of total cellular fatty acids.

**Alkane degradation in the presence of [<sup>13</sup>C]bicarbonate.** The mineral salt medium was prepared with NaH<sup>13</sup>CO<sub>3</sub> (isotopic enrichment, 99%; Cambridge Isotope Laboratories) instead of unlabeled sodium bicarbonate for growth of the cultures in this experiment. Each bottle containing 50 ml of medium was amended with 20  $\mu$ l of pentadecane or hexadecane and then inoculated with a hexadecane-grown culture (1%, vol/vol; a small inoculum size was used to minimize [<sup>13</sup>C]bicarbonate contamination from the inoculum). Duplicates of active cultures were prepared for each alkane. Cultures were incubated for 45 to 67 days and then sacrificed for extraction and analysis of total cellular fatty acids.

**Extraction of total cellular fatty acids.** Each culture was sacrificed by acidification with HCl (0.05 N final concentration), and cells were collected by filtering them through a Teflon filter (0.5  $\mu$ m pore size, 47 mm diameter; Gelman Sciences, Inc., Ann Arbor, Mich.). The collected cells were transferred to a screw-capped glass test tube with a Teflon-faced septum. A method to extract the total cellular fatty acids from the cells was adapted from a protocol for fatty acid analysis (Sherlock Microbial Identification System; MIDI, Inc., Newark, Del.).

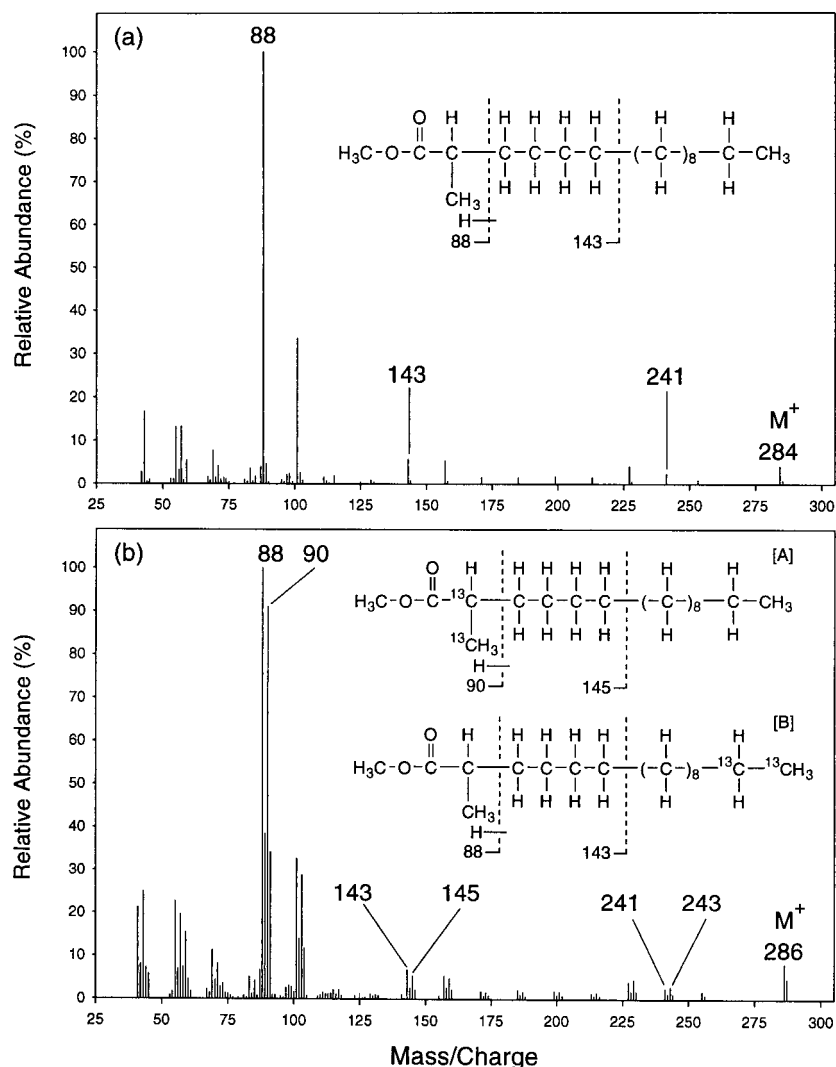


FIG. 3. Mass spectra of the methyl esters of the unlabeled (a) and  $^{13}\text{C}$ -labeled (b) 2-Me-16:0 fatty acids recovered from cultures of strain AK-01 grown on unlabeled hexadecane and  $[1,2-^{13}\text{C}_2]$ hexadecane, respectively. Chemical structures of the fatty acids represented by the mass spectra are shown as insets. The key diagnostic ion peaks are annotated with their  $m/z$  values. Structural compositions of the ion fragments represented by the annotated peaks are delineated. Intersection with a dotted line indicates a point of bond cleavage, and the ion fragment formed contains only the part of the molecule on the left side of the dotted line. (Note: the mass spectrum for the  $^{13}\text{C}$ -labeled 2-Me-16:0 fatty acid indicates the coexistence of two isotomers.)

To each tube with cells on the filter was added 2 ml of a NaOH-methanol solution (45 g of NaOH in 150 ml of methanol plus 150 ml of deionized water); the tubes were then capped and heated at  $100^\circ\text{C}$  for 30 min to release the cellular fatty acids by saponification and cooled. Then 4 ml of an HCl-methanol solution (325 ml of 6 N HCl and 275 ml of methanol) was added to each tube, and the tubes were heated at  $80^\circ\text{C}$  for 10 min to esterify the fatty acids. When the tube contents had cooled down, 1.25 ml of hexane-methyl *tert*-butyl ether (1:1) was added to extract the fatty acid methyl esters. The solvent extract was then washed with an aqueous NaOH solution (1.2%, wt/vol) to remove any unesterified fatty acids. Samples were stored in glass vials for subsequent analysis by gas chromatography-mass spectrometry (GC-MS).

**Analysis of fatty acid methyl esters.** Samples of fatty acid methyl esters were initially analyzed by GC-flame ionization detection (FID) with a gas chromatograph (model 5890, series II; Hewlett-Packard, Wilmington, Del.) equipped with a flame ionization detector and a DB-WAX column (30 m by 0.25 mm; film thickness, 0.25  $\mu\text{m}$ ; J&W Scientific, Folsom, Calif.). The column temperature was initially set at  $50^\circ\text{C}$  for 2 min, then increased to  $140^\circ\text{C}$  by  $15^\circ\text{C}$  per min and, finally, to  $220^\circ\text{C}$  by  $4^\circ\text{C}$  per min. The injector and detector temperatures were 250 and  $300^\circ\text{C}$ , respectively. The n-saturated and monounsaturated fatty acids were identified by comparing their retention times with those of commercially available authentic standards (Sigma Chemical Co.).

Samples were also analyzed by GC-MS with a 5890 series II gas chromatograph (Hewlett-Packard) equipped with a DB-5ms column (30 m by 0.25 mm;

film thickness, 0.25  $\mu\text{m}$ ; J&W Scientific) and a 5971 series mass-selective detector (Hewlett-Packard). The temperature program was as follows:  $60^\circ\text{C}$  for 1 min, increased by  $20^\circ\text{C}$  per min to  $200^\circ\text{C}$ , and then increased by  $4^\circ\text{C}$  per min to  $280^\circ\text{C}$ . The temperature for the injector and detector was  $280^\circ\text{C}$ . Identities of most n-saturated and unsaturated fatty acids were determined by comparing their retention times and mass spectra with those of commercially available authentic standards (Sigma Chemical Co.). Since no authentic standards were available for them, the various methylated and isotope-labeled fatty acids were identified by interpretation of their mass spectra based on information reported in the literature (11, 16, 26, 28, 33). Specific locations of  $^{13}\text{C}$  or deuterium atoms on the isotope-labeled fatty acids were determined by examining the diagnostically important ion fragments whose structural compositions had been previously determined (10, 12, 26, 37a). Each fatty acid in a sample was quantified based on its peak area in the total-ion chromatogram and was expressed as a percentage of the total fatty acids recovered. Preliminary analytical studies of a standard mixture of fatty acid methyl esters on the GC-MS instrument showed that those with chain lengths of  $\text{C}_{12}$  to  $\text{C}_{24}$  had similar response ratios based on the total mass of each compound injected.

**Nomenclature for fatty acids.** The fatty acid nomenclature recommended by the IUPAC-IUB (20) was adopted in this study. An n-saturated hexadecanoic acid is designated as 16:0, with the first number representing the number of carbons in the acyl group and the second number representing the number of



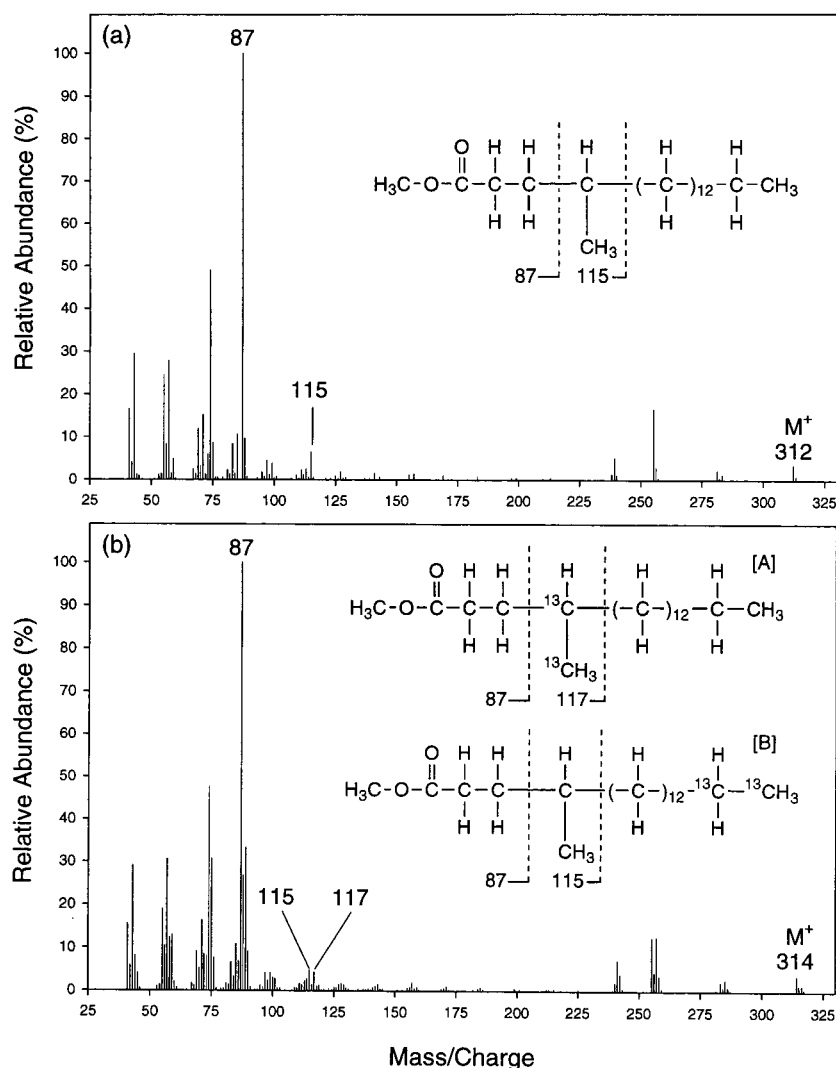


FIG. 5. Mass spectra of the methyl esters of the unlabeled (a) and  $^{13}\text{C}$ -labeled (b) 4-Me-18:0 fatty acids recovered from cultures of strain AK-01 grown on unlabeled hexadecane and  $[1,2-^{13}\text{C}_2]$ hexadecane, respectively. Chemical structures of the fatty acids represented by the mass spectra are shown as insets. (Note: the mass spectrum for the  $^{13}\text{C}$ -labeled 4-Me-18:0 fatty acid indicates the coexistence of two isotopomers.)

ecule (isotopomer B). Isotopomers are versions of the same molecule with different specific isotopic labelling patterns (33a).

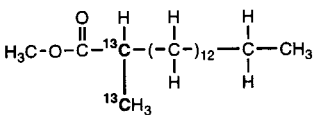
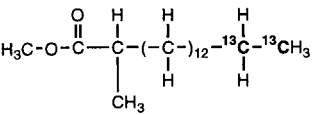
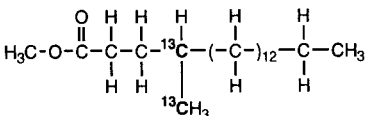
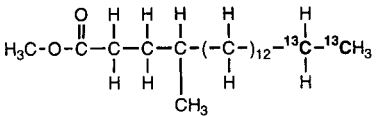
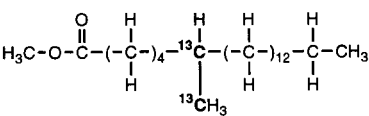
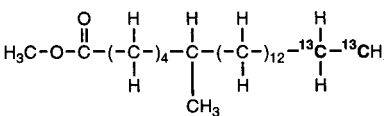
Further analysis of the mass spectrum of the  $^{13}\text{C}$ -labeled 2-Me-16:0 fatty acid showed that isotopomer B contained two  $^{13}\text{C}$  atoms between C-7 and C-16, indicating again the occurrence of two isotopomers. One (isotopomer A) contains two  $^{13}\text{C}$  atoms between C-1 and C-6 (thus,  $m/z = 145$ ), while the other (isotopomer B) has the two labels located between C-7 and C-16 (thus,  $m/z = 143$ ). These two labels in isotopomer B are most likely located at C-15 and C-16, their original locations on the  $[1,2-^{13}\text{C}_2]$ hexadecane. The presence of two  $^{13}\text{C}$  atoms at either end of the fatty acid, therefore, indicates that the initial attack occurs with the same likelihood at the labeled and the unlabeled termini of the hexadecane.

In addition, further examination of the mass spectra indicated that the two  $^{13}\text{C}$  atoms in isotopomer A of the 2-Me-16:0 fatty acid are located at C-2 and the 2-methyl group. As illustrated in Fig. 4a, the ion peak at  $m/z = 241$  for the unlabeled fatty acid (indicated in the mass spectrum in Fig. 3a) is formed

by the elimination of the segment of the molecule containing C-2, C-3, and the 2-methyl group (and an additional hydrogen, yielding a combined mass of 43) followed by the rearrangement of the remaining parts. This fragmentation mechanism is documented in the literature (12, 33, 37a). The corresponding ion peak in the mass spectrum of the  $^{13}\text{C}$ -labeled fatty acid appears as a doublet of peaks of comparable intensity at  $m/z = 241$  and 243 (Fig. 3b), indicating that isotopomer A contains two  $^{13}\text{C}$  atoms in the eliminated segment (Fig. 4b) while isotopomer B does not (Fig. 4c). The results show that the two  $^{13}\text{C}$  atoms in isotopomer A are located at C-2, C-3, and/or the 2-methyl group, not at C-1 or elsewhere (Fig. 4b). Furthermore, as shown earlier by examination of the doublet at  $m/z = 88$  and 90, none of the two  $^{13}\text{C}$  carbons on isotopomer A is located at C-3. Thus, they are located at C-2 and the 2-methyl group of isotopomer A.

Based on the results described above, we conclude that the 2-methyl group of the 2-Me-16:0 fatty acid is the original terminal carbon of the  $[1,2-^{13}\text{C}_2]$ hexadecane. This suggests that exogenous carbon is initially added onto the  $[1,2-^{13}\text{C}_2]$ hexade-

TABLE 1. Characteristics of the  $^{13}\text{C}$ -labeled 2-, 4-, and 6-methylated fatty acids recovered from cultures of strain AK-01 grown on  $[1,2-^{13}\text{C}_2]$ hexadecane

Cellular fatty acid <sup>a</sup>	Chemical structure and location of $^{13}\text{C}$ label for the fatty acid isotopmer <sup>b</sup> :		Relative abundance of isotopmers A and B <sup>c</sup>
	A	B	
2-Me-16:0			94:100
4-Me-18:0			91:100
6-Me-20:0			100:81

<sup>a</sup> Other recovered  $^{13}\text{C}$ -labeled fatty acids, namely 14:0, 10-Me-14:0, 16:0, and 10-Me-16:0, are not shown here. None of these fatty acids was found in the sterile control.

<sup>b</sup> Determined by interpretation of the mass spectrum for each fatty acid.

<sup>c</sup> Isotopmer A/B ratios. Results are estimations based on the relative intensities of the doublet of diagnostic ion peaks for each fatty acid.

cane subterminally at the C-2 position such that the original terminal carbon of the alkane then becomes the methyl group of the resulting 2-Me-16:0 fatty acid.

Mass spectral data on the 4-Me-18:0 fatty acid from the  $[1,2-^{13}\text{C}_2]$ hexadecane-grown cultures also indicate that the 4-methyl group is the original terminal carbon of the alkane. Figure 5 shows the mass spectra of the unlabeled (Fig. 5a) and  $^{13}\text{C}$ -labeled (Fig. 5b) 4-Me-18:0 fatty acid recovered from cultures grown on unlabeled hexadecane and  $[1,2-^{13}\text{C}_2]$ hexadecane, respectively. The presence of two  $^{13}\text{C}$  atoms in the latter fatty acid is indicated by the molecular ion at  $m/z = 314$  (Fig. 5b).

The  $^{13}\text{C}$ -labeled 4-Me-18:0 fatty acid also occurs as two isotopmers, one with the two  $^{13}\text{C}$  atoms located at C-4 and the 4-methyl group. As shown in Fig. 5a, the ion fragment at  $m/z = 87$  is formed by cleavage of the C-3-C-4 bond and includes C-1 to C-3 of the unlabeled 4-Me-18:0 fatty acid. In addition, the ion fragment at  $m/z = 115$  is formed by cleavage of the C-4-C-5 bond and includes C-1 to C-4 and the 4-methyl group of the molecule (12, 26, 33). The difference between the  $m/z$  values of these two ion peaks, which amounts to 28 mass units, represents the combined mass of C-4 plus the attached hydrogen atom and the 4-methyl group [ $-\text{CH}(\text{CH}_3)-$ ].

On the mass spectrum for the  $^{13}\text{C}$ -labeled 4-Me-18:0 fatty acid (Fig. 5b), the ion peak corresponding to that at  $m/z = 87$  for the unlabeled fatty acid (Fig. 5a) remains the same, indicating the absence of  $^{13}\text{C}$  label at C-1 to C-3. The peak corresponding to  $m/z = 115$ , however, appears as a doublet of peaks at  $m/z = 115$  and  $117$  of comparable intensity. Thus, the  $^{13}\text{C}$ -labeled 4-Me-18:0 fatty acid occurs as two isotopmers of comparable abundance, one labeled with two  $^{13}\text{C}$  atoms, at C-4 and the 4-methyl group (isotopmer A in Fig. 5b), and the other with no isotope label on these carbons. The two  $^{13}\text{C}$  atoms in the latter case are most likely located at C-17 and C-18 (isotopmer B in Fig. 5b). These results show that the 4-methyl group is the original terminal carbon of the  $[1,2-^{13}\text{C}_2]$ hexadecane and further underscore the point that exogenous carbon is added onto the alkane subterminally at the C-2 position. It

also appears that the 4-Me-18:0 fatty acid is formed after elongation of the 2-Me-16:0 fatty acid by two carbon units through fatty acid chain elongation reactions (31).

Table 1 summarizes the characteristics of the 2-, 4-, and 6-methylated fatty acids derived from the  $[1,2-^{13}\text{C}_2]$ hexadecane. For each fatty acid, the positions of the  $^{13}\text{C}$  atoms on each isotopmer and the relative abundance of the two different isotopmers are shown. None of these  $^{13}\text{C}$ -labeled fatty acids was found in the sterile control. Like the 2-Me-16:0 and 4-Me-18:0 fatty acids, the  $^{13}\text{C}$ -labeled 6-Me-20:0 fatty acid contains two  $^{13}\text{C}$  atoms, at C-6 and the 6-methyl group, in one isotopmer (mass spectrum not shown). Thus, it is a result of a similar carbon addition mechanism followed by chain elongation. Other C-even fatty acids (14:0, 10-Me-14:0, 16:0, and 10-Me-16:0), which are also  $^{13}\text{C}$  labeled, appear to be formed after further metabolism of the 2-Me-16:0 fatty acid (data not shown).

**Analyses of perdeuterated pentadecane-derived fatty acids.** Mass spectral analyses of the deuterated fatty acids also indicate that exogenous carbon is added onto the alkane chain subterminally to form a monomethylated fatty acid. Figure 6 shows the mass spectra of the methyl esters of the unlabeled (Fig. 6a) and deuterated (Fig. 6b) 2-Me-15:0 fatty acid recovered from cultures grown on unlabeled and perdeuterated pentadecane, respectively. As shown in Fig. 6a, the ion fragment,  $m/z = 88$ , of the unlabeled 2-Me-15:0 fatty acid is formed by the McLafferty rearrangement as described earlier (26, 33). The corresponding peak for the deuterated fatty acid (Fig. 6b) is shifted up by 4 mass units ( $m/z = 92$ ), indicating the presence of four deuterium atoms in the ion fragment. Since the methanol used for the fatty acid derivatization was not deuterated and one deuterium would have been contributed by the McLafferty rearrangement (12), we deduce that three deuterium atoms are located at C-2 and the 2-methyl group. Similar to the study with the  $[1,2-^{13}\text{C}_2]$ hexadecane, the results here indicate that the 2-methyl group of the deuterated 2-Me-15:0 fatty acid is the original terminal carbon of the perdeuterated pentadecane. Thus, the 2-Me-15:0 fatty acid is the result of

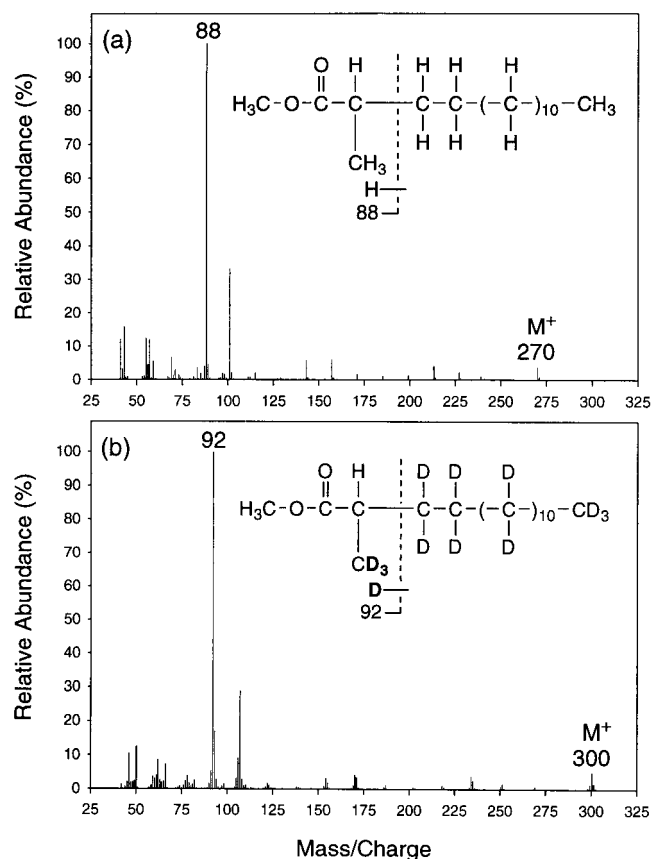


FIG. 6. Mass spectra of the methyl esters of the unlabeled (a) and deuterated (b) 2-Me-15:0 fatty acids recovered from cultures of strain AK-01 grown on unlabeled and perdeuterated pentadecane, respectively. Chemical structures of the fatty acids represented by the mass spectra are shown as insets. Interpretation of the mass spectra shows that there are three deuterium atoms, located on C-2 and the 2-methyl group of the deuterated 2-Me-15:0 fatty acid. The locations of all three deuterium atoms on the 2-methyl group as shown in Fig. 6b represent one possible arrangement.

subterminal carbon addition to the perdeuterated pentadecane at the C-2 position.

Mass spectral analyses of the deuterated 4-Me-17:0 and 6-Me-19:0 fatty acids from cultures grown on perdeuterated pentadecane also indicated the occurrence of subterminal carbon addition. As shown in Table 2, the 2-, 4-, and 6-methylated fatty acids all contain three deuterium atoms, located on their methyl groups and the adjacent carbons, indicating that the methyl groups also contain the original terminal carbon of the perdeuterated pentadecane and that exogenous carbon is added to the deuterated pentadecane subterminally at C-2. This thus shows that the deuterated 4-Me-17:0 and 6-Me-19:0 fatty acids are formed by chain elongation of the 2-Me-15:0 fatty acid. Deuterium atoms are also found on the carbons between the carboxyl groups and methyl groups of the 4-Me-17:0 and 6-Me-19:0 fatty acids (Table 2). This may be effected by the initial carbon addition reaction in that the carbon compound being added to the perdeuterated pentadecane is also directly derived from the labeled alkane and thus still carries deuterium atoms. Other C-odd deuterated fatty acids, namely 15:0 and 10-Me-15:0, are also recovered and are likely formed by further metabolism of the 2-Me-15:0 fatty acid. The lack of deuterium atoms in the 10-methyl group of the deuterated 10-Me-15:0 (data not shown) indicates that this C-10 methyl

group is exogenously added, in contrast to those of the 2-, 4-, and 6-methylated fatty acids.

**Fatty acids formed in the presence of [ $^{13}\text{C}$ ]bicarbonate.** The 2-methylated fatty acids recovered from cultures of AK-01 grown on unlabeled pentadecane and hexadecane in the presence of [ $^{13}\text{C}$ ]bicarbonate were characterized by mass spectrometry. The mass spectra of the 2-Me-15:0 and 2-Me-16:0 fatty acids formed in the presence of [ $^{13}\text{C}$ ]bicarbonate appeared the same as those formed in the presence of unlabeled bicarbonate. The results indicate that the carboxyl groups of the 2-methylated fatty acids are not  $^{13}\text{C}$  labeled. Hence, in AK-01, the carbon addition at C-2 of the alkane chain does not appear to occur through the addition of an inorganic carboxyl group.

## DISCUSSION

The alkane-degrading strain AK-01 was isolated from petroleum-contaminated estuarine sediments. It grows anaerobically on alkanes ( $\text{C}_{13}$  to  $\text{C}_{18}$ ) and is able to oxidize them to carbon dioxide coupled to sulfate reduction (35). In the present study, the mechanism for anaerobic alkane degradation by the strain was determined. Although 1-alkene and 1-alkanol have been suggested as intermediates (29) and strain AK-01 is able to utilize both for growth (35), neither compound was found in the alkane-grown cultures in our experiments. That they are unlikely to be intermediates is also supported by data from studies of strain Hxd3 (1, 3).

Although no free fatty acid was extractable from whole cultures of AK-01 grown on alkanes, the impact of the alkane substrates' carbon numbers on the total cellular fatty acid composition (Fig. 2) is highly suggestive of alkanes being oxidized to fatty acids by the organism. This has been observed with the other alkane-utilizing sulfate reducers (3, 14) and also for microorganisms capable of aerobic alkane utilization (13, 21). The fatty acids formed after alkane oxidation can be directly

TABLE 2. Characteristics of the deuterated, 2-, 4-, and 6-methylated fatty acids recovered from cultures of strain AK-01 grown on perdeuterated pentadecane

Deuterated fatty acid <sup>a</sup>	Chemical structure and distribution of deuterium label <sup>b</sup>
2-Me-15:0	$\text{H}_3\text{C}-\text{O}-\overset{\text{O}}{\parallel}{\text{C}}-\overset{\text{H}}{\underset{\text{CD}_3}{\text{C}}}-\overset{\text{D}}{\underset{\text{D}}{\text{C}}}-\text{(-}\overset{\text{D}}{\text{C}}\text{-)}_{12}-\text{CD}_3$
4-Me-17:0	$\text{H}_3\text{C}-\text{O}-\overset{\text{O}}{\parallel}{\text{C}}-\text{(-}\overset{\text{H}}{\text{C}}\text{D}_2\text{H}_2\text{-)}-\overset{\text{H}}{\underset{\text{CD}_3}{\text{C}}}-\overset{\text{D}}{\underset{\text{D}}{\text{C}}}-\text{(-}\overset{\text{D}}{\text{C}}\text{-)}_{12}-\text{CD}_3$
6-Me-19:0	$\text{H}_3\text{C}-\text{O}-\overset{\text{O}}{\parallel}{\text{C}}-\overset{\text{H}}{\underset{\text{H}}{\text{C}}}-\text{(-}\overset{\text{H}}{\text{C}}\text{D}_2\text{H}_4\text{-)}-\overset{\text{H}}{\underset{\text{CD}_3}{\text{C}}}-\overset{\text{D}}{\underset{\text{D}}{\text{C}}}-\text{(-}\overset{\text{D}}{\text{C}}\text{-)}_{12}-\text{CD}_3$

<sup>a</sup> Other recovered deuterated fatty acids, namely 15:0 and 10-Me-15:0, are not shown here. No deuterated fatty acids were found in the sterile control.

<sup>b</sup> Determined by interpretation of the mass spectrum for each fatty acid.

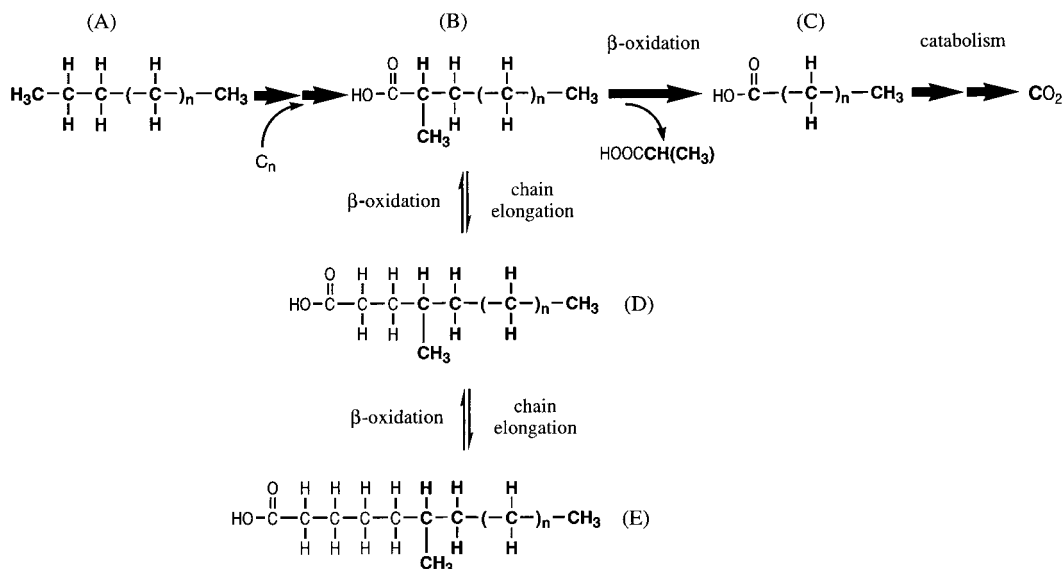


FIG. 7. Proposed pathway for anaerobic alkane metabolism by strain AK-01. (The original alkane atoms are boldfaced, and the major pathway is indicated by bold arrows.)

incorporated into the cellular lipids of the alkane-degrading organisms (5, 18, 22).

The recovery of deuterated and  $^{13}\text{C}$ -labeled fatty acids in cultures of AK-01 grown on the perdeuterated and  $^{13}\text{C}$ -labeled alkanes, respectively, clearly confirms that alkanes are anaerobically oxidized to fatty acids. In addition, the carbon numbers of the labeled fatty acids also correlate with those of the alkane substrates. When  $[1,2\text{-}^{13}\text{C}_2]$ hexadecane was used as a growth substrate, the recovered  $^{13}\text{C}$ -labeled fatty acids were all C even (Table 1). When perdeuterated pentadecane was used, the recovered deuterated fatty acids were all C odd (Table 2). The results show that the fatty acids derived from the alkanes can be  $\beta$ -oxidized and mineralized to  $\text{CO}_2$  (24, 35) or further transformed through chain elongation and C-10 methylation (31) by AK-01.

Evidence presented here for a proposed mechanism of alkane oxidation is derived from mass spectral analyses of the resulting stable-isotope-labeled fatty acids. The mass spectra of the methyl esters of many unlabeled and deuterated fatty acids have been reported (11, 16, 26, 28, 33). The compositions and fragmentation mechanisms of various ion fragments, which contain different parts of the fatty acid molecule, are also well studied (10, 12, 26). Based on this information, many fatty acids can be identified and the locations of  $^{13}\text{C}$  and deuterium atoms on the isotope-labeled fatty acids can be determined.

Three lines of evidence support a mechanism by which exogenous carbon is added to an alkane subterminally as an initial step in anaerobic alkane metabolism. First, the 2-Me-16:0, 4-Me-18:0, and 6-Me-20 fatty acids derived from the  $[1,2\text{-}^{13}\text{C}_2]$ hexadecane each contains two  $^{13}\text{C}$  atoms, located at the methyl group and the adjacent carbon (Table 1). The results indicate that exogenous carbon is added subterminally to the hexadecane at the C-2 position, such that the terminal carbon of the alkane becomes the 2-, 4-, and 6-methyl groups of the fatty acids formed subsequently.

Second, the 2-Me-15:0, 4-Me-17:0, and 6-Me-19 fatty acids derived from the perdeuterated pentadecane each contains three deuterium atoms, located on the methyl group and the adjacent carbon (Table 2). This also supports the same mechanism, namely subterminal carbon addition to the perdeuter-

ated pentadecane at C-2, resulting in the original terminal carbon of the alkane forming the 2-, 4-, and 6-methyl groups of these fatty acids.

Third, the 2-, 4-, and 6-methylated fatty acids are found only in cultures grown on alkanes, and their chain lengths always correlate with those of the alkane substrates (Fig. 2). These uncommon monomethylated fatty acids are not observed when AK-01 is grown on other aliphatic substrates (1-alkenes, 1-alkanols, and fatty acids), and they are not observed in other sulfate-reducing bacteria (36). Therefore, these fatty acids are formed specifically by the initial reactions of alkane oxidation. Furthermore, it appears that the 2-methylated fatty acids are initially formed from the alkanes and can be subsequently elongated by two carbon units to form the 4-methylated and 6-methylated fatty acids through the fatty acid elongation reactions (Tables 1 and 2) (31). The nature of the carbon compound being added onto the alkane chain, however, remains unknown at this time. The absence of the 2-, 4-, and 6-methylated fatty acids in AK-01 cultures grown on 1-alkenes also indicates that carbon addition is not mediated by the formation of a terminal double bond in the alkane.

Although carboxylation has been demonstrated as an initial reaction of aromatic hydrocarbon metabolism under anaerobic conditions (30, 39), results from our experiment with  $[^{13}\text{C}]$ bicarbonate show that the subterminal carbon addition is probably not due to a direct carboxylation. On the other hand, it has been demonstrated that toluene is anaerobically metabolized by addition of fumarate to form benzyl succinate and then benzoyl coenzyme A (4). Hence, we cannot discount the possibility that a compound with more than one carbon is added to an alkane to form the first intermediate, which is then transformed into a 2-methylated fatty acid.

The anaerobic attack on alkanes by AK-01 involves carbon addition at the C-2 position. Such selectivity for the subterminal carbon may be caused by the conformational structure of the degrading enzyme. Studies with model systems support the hypothesis that the terminal carbons of alkanes are selectively hydroxylated by cytochrome P-450 because of steric hindrance at its active site (23). The degrading enzyme of AK-01 may have a less sterically hindered active site, such that the subter-



minal carbon at C-2 can be attacked. Furthermore, as a secondary carbon, its C—H bond is weaker than that of a terminal, primary carbon (bond dissociation energy, 94.5 versus 98 kcal/mol for propane, for example) (15) and thus should be more readily broken.

Based on the collective observations in this study, a proposed pathway for anaerobic oxidation of alkanes by strain AK-01 is summarized in Fig. 7. An alkane (A) is initially attacked subterminally by the addition of a carbon compound at the C-2 position to form a 2-methylated fatty acid (B). This fatty acid then undergoes  $\beta$ -oxidation (24, 25, 34) to form an *n*-saturated fatty acid (C) which can be further  $\beta$ -oxidized and eventually mineralized to CO<sub>2</sub> (35). In addition, the initially formed 2-methylated fatty acid (B) can also be elongated to form the 4-methylated (D) and 6-methylated (E) fatty acids.

In summary, we provide direct evidence that alkanes are oxidized to fatty acids by strain AK-01 under strictly anaerobic conditions. Subterminal addition of a carbon(s) to the hydrocarbon chain as an initial reaction in anaerobic alkane degradation is also demonstrated through a detailed characterization of these fatty acid metabolites. This carbon addition reaction represents a novel mechanism by which alkanes can be activated without oxygen, in contrast to the well-known hydroxylation reaction mediated by monooxygenases in aerobic organisms.

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