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Identification of a novel C₃₃ botryococcane and C₃₃ botryococconone in the Maoming Basin, China

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ABSTRACT

A novel C₃₃ botryococcane in a sedimentary sample from the Maoming Basin in China was found to have a methyl group positioned β to the quaternary carbon C-10 in its structure based on IR, MS and NMR analyses and Lindeman-Adams modelling. This structure is inconsistent with the prediction from the currently accepted botryococcene biosynthetic pathway and different from the skeleton for the C₃₃ botryococcones suggested in the literature. A C₃₃ botryococcon-24-one with the same carbon skeleton as the C₃₃ botryococcane was also found to co-occur in the sediment sample. The location of the carbonyl group at C-24 is different from that of a previously reported botryococconone from an Australian collection of race B of *Botryococcus braunii* where it is positioned at C-15. We propose a biogeochemical pathway to explain the occurrence of this unique skeleton in the Maoming Basin sediment. The biochemical component of the pathway involves electrophilic or nucleophilic attack between two farnesyl diphosphates (FPPs) leading to direct or indirect formation of a C₃₀ botryococcene via a cyclobutane ring intermediate (1'-2-3-2' cyclisation). A Retro-Prins reaction and subsequent methylation gives rise to C₂₉ and C₃₃ botryococcones in sequence. The geochemical component involves first in-water-column (photo)-oxidation of the alkene to a C₃₃ botryococconone and subsequent reduction (hydrogenation) to stabilize the ketone as an alkane.

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1. Introduction

Botryococcones are geochemically important biomarkers for assessing depositional environment and geological age (Philp and Lewis, 1987; Guy-Ohlson, 1992; Volkman, 2014) as they are known to be synthesised only by race B of the freshwater unicellular green microalga *Botryococcus braunii* (Maxwell et al., 1968), which is known to contribute significantly to the botryococcones (and other botryococconoids) found in oil shales and oils. In field-grown or cultured *B. braunii*, botryococcones are found to have a carbon range of C₃₀–C₃₇ (Metzger et al., 1985a, 1985b; Okada et al., 1995; de Mesmay et al., 2008), and are often dominated by the

C₃₄ homologue(s) (Metzger and Largeau, 1999; Zhang et al., 2007; de Mesmay et al., 2008). This predominance of C₃₄ botryococcones in *B. braunii* is consistent with the C₃₄ botryococcones found in fossil samples. For example, C₃₄ botryococcones were first discovered in a Sumatran crude oil (Moldowan and Seifert, 1980) and subsequently in Australian coastal bitumens (McKirdy et al., 1986, 1994; Dowling et al., 1995) and the Kishenehn Formation in American Montana (Curiale, 1987). Grice et al. (1998) reported the occurrence of cyclobotryococcones and structurally related organic sulfur compounds (OSC) in hypersaline sediments of Miocene/Pliocene age from the Sdom Formation in the Dead Sea. A novel (partially reduced) C₃₄ botryococcene (1,6,17,21-octahydro botryococcene) from *B. braunii* was identified in a sediment from Sacred Lake, Mount Kenya (Huang and Murray, 1995). C₃₂–C₃₄ botryococcones were also detected in Florida Everglades freshwater wetland soil (Gao et al., 2007). C₃₃–C₃₇ botryococcones and partially reduced mono- and dicyclic unsaturated derivatives were reported in Holocene sediments from Lake Masoko in Tanzania (de Mesmay et al., 2008; Grossi et al., 2012).

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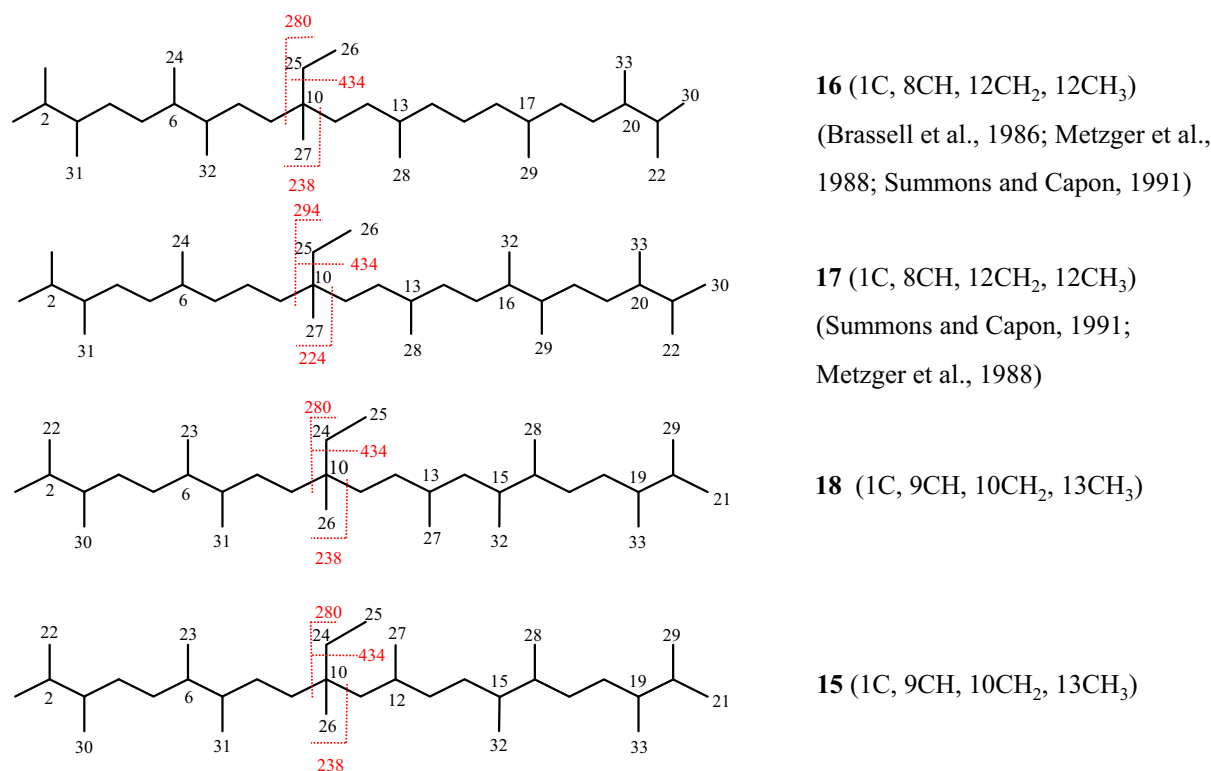


Fig. 1. The four possible carbon skeletons of C₃₃ botryococcanes deducible from the mass spectrum obtained by Brassell et al. (1986).

Botryococcanes with an obvious odd carbon number predominance (C₃₁ and C₃₃), although rare, were also reported in the Chinese Maoming Basin (Brassell et al., 1986). The identification of these botryococcanes was solely based on a comparison of their mass spectra with that of a C₃₄ botryococcan standard (Moldowan and Seifert, 1980) and other botryococcan and botryococane homologues (Metzger and Casadevall, 1983; Metzger et al., 1985a), and therefore was not conclusive. In the case of C₃₃ botryococcan, two different structures (**16** and **17**; Fig. 1) were suggested by Brassell et al. (1986), Metzger et al. (1988) and Summons and Capon (1991). However, two equally possible structures (**15** and **18**; Fig. 1) are also consistent with the mass spectrum, and in particular the fragmentation pattern around the quaternary carbon (key ions at *m/z* 238, 280, and 434; Fig. 1). The four possible structures differ mainly in the location of the methyl group adjacent to the quaternary carbon C-10, i.e., C-28 (connected to C-13, γ to C-10) in **16** and **17**, C-27 (connected to C-13, γ to C-10) in **18** (connected to C-13, γ to C-10) and **15** (connected to C-12, β to C-10).

To firmly establish the structure of the predominant C₃₃ botryococcan in the Maoming Basin, we isolated the C₃₃ botryococcan from the saturated fraction. We report here the structural reassessment of this alkane and a ketone, with the same carbon number and skeleton isolated from the polar fraction, by a combination of MS and NMR techniques and discuss their genetic relationship and possible geochemical and biosynthetic pathways leading to their formation in the Maoming Basin.

2. Material and methods

2.1. Sampling, separation and purification

The Eocene sedimentary sample used in this study was collected from an outcrop of the Maoming Basin located in the southwest of Guangdong Province in China. It was collected from the same formation at the same location from which botryococcanes

were reported by Brassell et al. (1986) (Supplementary Fig. S1). The ground sample was Soxhlet-extracted with methanol/dichloromethane (1:9, v/v) for 72 h. The asphaltene fraction was centrifugally removed by precipitation in *n*-hexane. The maltene fraction was further separated into saturated, aromatic and polar fractions on a glass column (0.3 m \times 1 cm i.d.) packed with silica (80–100 mesh) activated at 120 °C for 12 h and alumina activated at 450 °C for 4 h in tandem (4:1, v/v), using sequential elution with *n*-hexane (80 mL), *n*-hexane/dichloromethane (40 mL, 1:1, v/v) and methanol (40 mL). The polar fraction was further sub-fractionated on a column (0.3 m \times 1 cm i.d.) packed with silica (100–230 mesh), activated at 120 °C for 12 h, by eluting with CH₂Cl₂/*n*-hexane (1:3 v/v, 40 mL, subfraction 1) and CH₂Cl₂/*n*-hexane (1:2 v/v, 40 mL, subfraction 2).

GC–MS analyses were then conducted on both subfraction 2 and the saturated hydrocarbon fraction. These fractions were then subjected to preparative gas chromatography (pGC) (Eglinton et al., 1996; Özek and Demirci, 2012; Zuo et al., 2013) with a DB-5 column (60 m \times 0.53 mm i.d. \times 1.5 μ m film thickness) on an Agilent 7890 GC coupled to a Gerstel-preparative fraction collector (PFC) to isolate and purify the two target compounds. The preparative conditions used were as follows: He was used as carrier gas at a flow rate of 3.0 mL/min. The column was held at 80 °C for 2 min, then heated to 300 °C at 30 °C/min, and held for 40 min at 300 °C. The purified botryococcanone (approx. 2.0 mg, see Supplementary Figs. S2 and S4) and C₃₃ botryococcan (approx. 3.5 mg, see Supplementary Figs. S1 and S3) were then subjected to FT-IR, HR-EIMS, and NMR analyses for further structural characterisation.

2.2. Instrumental analyses

FT-IR analysis of the purified botryococcanone was conducted on a FT-IR Tenor 27 spectrophotometer (Bruker Corporation, Billerica, USA) with the sample loaded as a KBr pellet. The spectrometer

was operated in absorption mode with absorptions in the range 475 cm^{-1} to 3675 cm^{-1} recorded at a resolution of 2 cm^{-1} .

HR-EIMS analyses of the botryococcane and botryococconone were conducted on a Thermo Finnigan MAT95XP mass spectrometer (Germany) to determine the accurate mass. The HR-EIMS system was operated in the electron impact ionization mode (42 eV) at a mass resolution of $R > 10,000$ (10% valley).

^1H and ^{13}C NMR spectral analyses were conducted on a Bruker 600M AVANCE III spectrometer (operated at 600.19 MHz for ^1H and 150.92 MHz for ^{13}C). Broadband ^{13}C -decoupled spectra were recorded in CDCl_3 , with TMS as internal standard. ^1H NMR chemical shifts were referenced to residual proton signal (7.26 ppm). ^{13}C NMR chemical shifts were referenced to the central line of ^{13}C multiplet (77.0 ppm). A combination of 1D and 2D experiments, ^1H - ^{13}C correlation spectroscopy (COSY), heteronuclear single quantum correlation (HSQC), and heteronuclear multiple bond correlation (HMBC) were performed to assign the individual resonances. Distortionless enhanced polarization transfer (DEPT) spectra were used to determine the multiplicity of each ^{13}C nucleus.

GC-MS analyses of the saturated and polar fractions were performed on a Trace Ultra GC coupled to a Thermo DSQ-II mass spectrometer operating with an ionization energy of 70 eV and a m/z range of 50–600. An HP-5 column (30 m \times 0.25 mm i.d. \times 0.25 μm film thickness) was used. The oven temperature was ramped

from $80\text{ }^\circ\text{C}$ (hold for 2 min) to $295\text{ }^\circ\text{C}$ (hold for 20 min) at rate of $6\text{ }^\circ\text{C}/\text{min}$. He was used as the carrier gas with a constant flow of $1.2\text{ mL}/\text{min}$.

3. Results and discussion

3.1. Structural elucidation of the novel C_{33} botryococcane and botryococconone

Abundant C_{31} and C_{33} botryococcanes were found in the Maoming sediment extracts (Supplementary Fig. S1). The overall distribution of these hydrocarbons (including botryococcanes) in the saturated hydrocarbon fraction is very similar to that reported by Brassell et al. (1986) implying that we have essentially reproduced the profile of the saturated fraction obtained by these authors. A compound suspected to be the corresponding ketone of the C_{33} botryococcane was discovered and isolated from the polar fraction (see Supplementary Fig. S2 for the TICs of the polar fraction and subfraction 2). To confirm their skeletons, we first assigned the structure of the C_{33} botryococcane by NMR and then verified the skeleton of the C_{33} ketone by MS and NMR analyses.

C_{33} botryococcane. In order to ascertain its structure, in particular the substructure around the C-10 quaternary carbon (Fig. 1), the HMBC data were investigated in detail (Fig. 2, see also

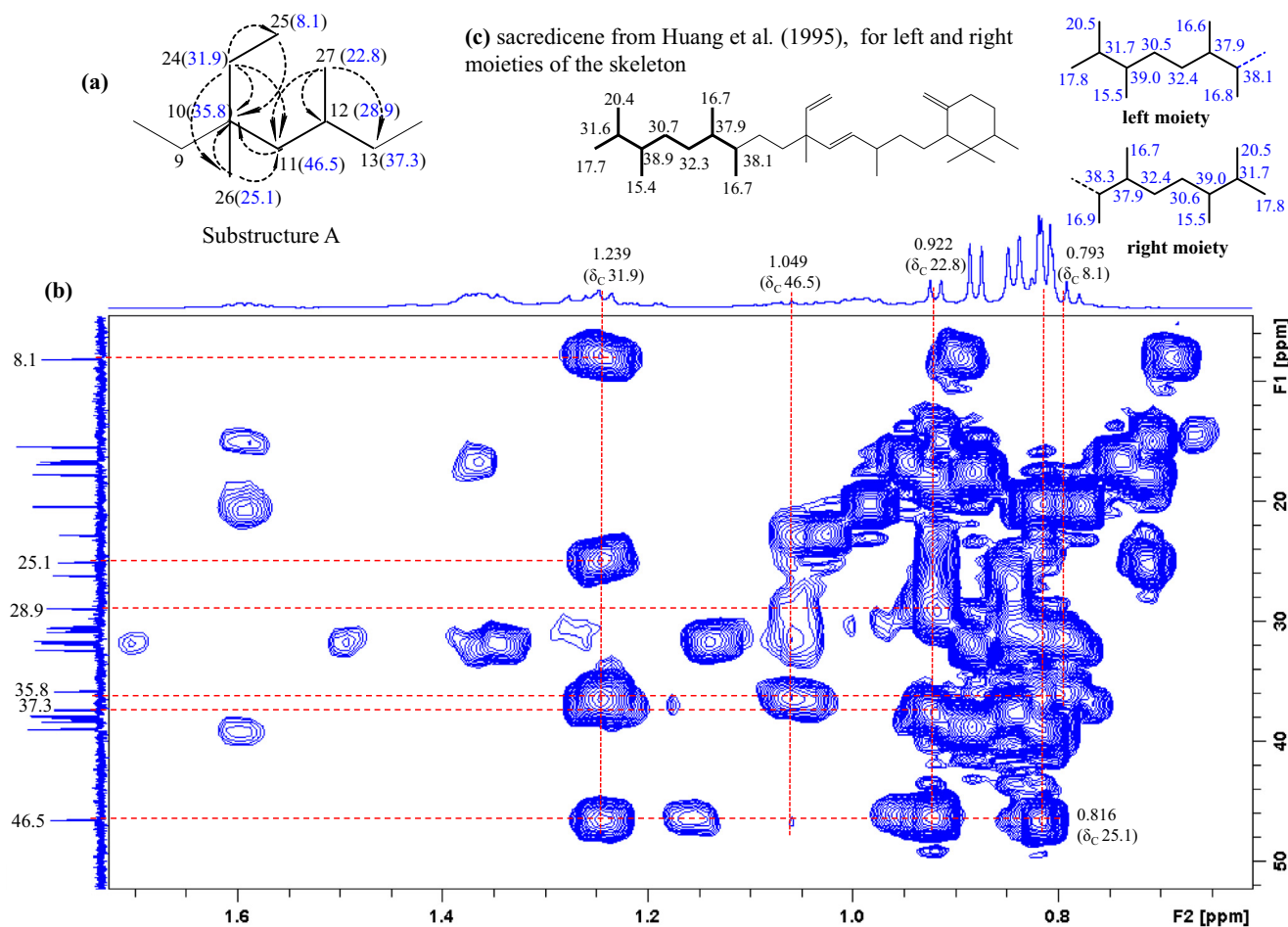


Fig. 2. The core substructure (a) around the quaternary carbon as deduced from the HMBC spectrum (b) for the C_{33} botryococcane (**15**) in this study. The position-specific chemical shifts in the left and right moieties in the C_{33} botryococcane identified here are very similar to that of the equivalent substructure of the sacredicene (c) identified by Huang et al. (1995).

Supplementary Table S1 and Supplementary Figs. S8–S10 for other NMR data). The correlations from H-24 (δ_{H} 1.239) to C-10 (δ_{C} 35.8), C-11 (δ_{C} 46.5), C-25 (δ_{C} 8.1) and C-26 (δ_{C} 25.1) indicated connections between C-10/C-11, C-10/C-24, C-10/C-26 and C-24/C-25 (Fig. 2a), which could also be further confirmed by the correlations from H-25 (δ_{H} 0.793) to C-10 (δ_{C} 35.8), H-26 (δ_{H} 0.816) to C-10 (δ_{C} 35.8), H-26 (δ_{H} 0.816) to C-11 (δ_{C} 46.5) and H-11 (δ_{H} 1.049) to C-10 (δ_{C} 35.8). Correlations from H-27 (δ_{H} 0.922) to C-11 (δ_{C} 46.5), C-12 (δ_{C} 28.9) and C-13 (δ_{C} 37.3) (Fig. 2b) established the connections C-11/C-12, C-12/C-13, and C-12/C-27. The values of the chemical shifts measured for the moieties left to C-8 and right to C-14 (Fig. 2c) are in good agreement with those reported by Huang et al. (1995) for a moiety of sacredicene, a botryococcane derivative sharing the same substructure **A** of the botryococcane we identified here. Thus, it is reasonable to assign the structure of C_{33} botryococcane as skeleton **15** (Fig. 1), which has a methyl group connected to a carbon β to C-10.

C_{33} botryococconone. The mass spectrum (Fig. 3b) of the C_{33} ketone (see Supplementary Fig. S2 for the TIC of the polar fraction) displayed two distinct ions: m/z 254 and 296, both having an extra 16 Da added to the characteristic ions of m/z 238 and 280 for the C_{33} botryococcane as identified by Brassell et al. (1986) (Fig. 3a, see also Summons and Capon, 1991). Since cleavages around the sole quaternary carbon (C-10) in the botryococcane lead to the formation of the characteristic ions m/z 238/239 and 280/281 (Fig. 3a), which correspond to the loss of the two alkyl chains (C-1 – C-9) and (C-11 – C-21) (Fig. 3a), respectively, the characteristic ions m/z 254 and 296 could be explained by a McLafferty rearrangement around the quaternary carbon of a C_{33} botryococconone with a carbonyl group positioned on the ethyl group (C-24) connected to quaternary carbon C-10 (Fig. 3c). Such a positioning of the carbonyl group is further supported by the very similar

fragmentation pattern of C_{33} ketone to that of the C_{33} botryococcane (Fig. 3a): base peak m/z 71 (loss of the alkyl group C_5H_{11} from C-3 and C-19, Fig. 3b), major peak m/z 127 (loss of C_9H_{19} at C-6 and C-16, Fig. 3b) and the other ions m/z 57, 85, 99, 113, 141, 155, 169, 183, 197, 211, and 225).

HR-EIMS analysis (see Supplementary Fig. S6 for a partial mass spectrum) showed that the ketone has an accurate mass of 478.5108, in agreement with a calculated mass of 478.5114 for a formula of $\text{C}_{33}\text{H}_{66}\text{O}$, indicating one degree of unsaturation. FT-IR absorption (Supplementary Fig. S7) at 1704.8 cm^{-1} also indicated the presence of a carbonyl group ($1705\text{--}1725\text{ cm}^{-1}$) and excluded the possibility of an aldehyde group due to the absence of absorption characteristic bands for an aldehyde group ($2820\text{--}2850$ and $2720\text{--}2750\text{ cm}^{-1}$). Therefore, the compound was initially identified as a C_{33} botryococcon-24-one (Fig. 3b).

In such a botryococconone skeleton, one would expect strong correlations in the NMR chemical shifts (see Supplementary Table S2 and Supplementary Figs. S11–S13 for detailed 1D and 2D data) between the carbonyl carbon at C-24 and its neighbouring carbons and hydrogens at C-25, C-26, C-11 and possibly C-9 (Fig. 4). This is exactly what was observed with the HMBC correlation spectrum (Fig. 4b): carbon C-24 (δ_{C} 214.3) is correlated with H-9 (δ_{H} 1.500), H-11 (δ_{H} 1.673 and δ_{H} 1.358), H-26 (δ_{H} 1.102) and H-25 (δ_{H} 2.125). Note that the δ_{C} (214.3) of the C-24 (Fig. 4a and b) is close to that of the carbonyl carbon C-15 (δ_{C} 210.0) of the C_{33} botryococconone identified by Summons and Capon (1991). The δ_{C} at 25.5 ppm for C-25 is consistent with the fact that C-25 is deshielded by the neighbouring carbonyl group (Breitmaier and Voelter, 1978; Cheng and Bennett, 1991; Cheng, 1994). Further, in the HMBC spectrum, it was observed that H-25 (δ_{H} 2.125) was correlated to the carbonyl carbon C-24 (δ_{C} 214.3), consistent with the expectation that a carbonyl at C-24 would give

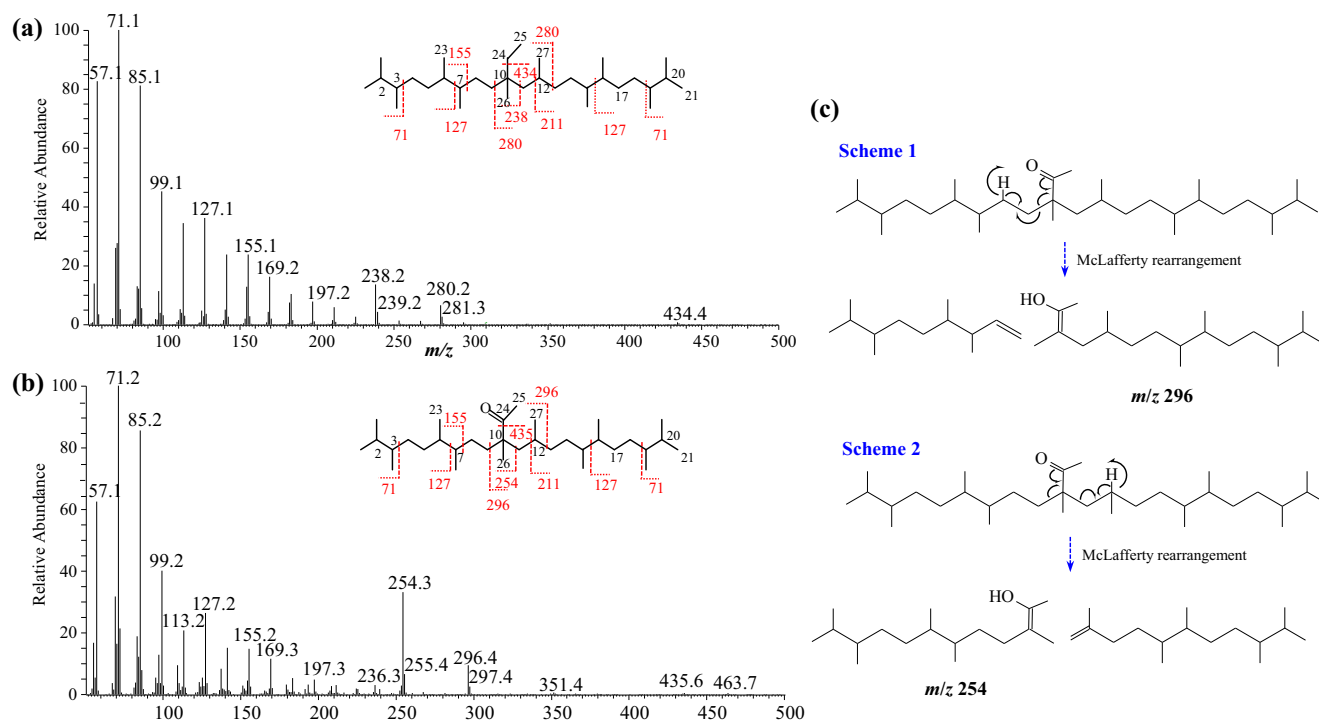


Fig. 3. The mass spectra for the novel C_{33} botryococcane (a) and C_{33} botryococconone (b) identified in this study; (c) McLafferty rearrangement leading to the formation of fragment ions m/z 254 and 296 of C_{33} botryococconone (b).

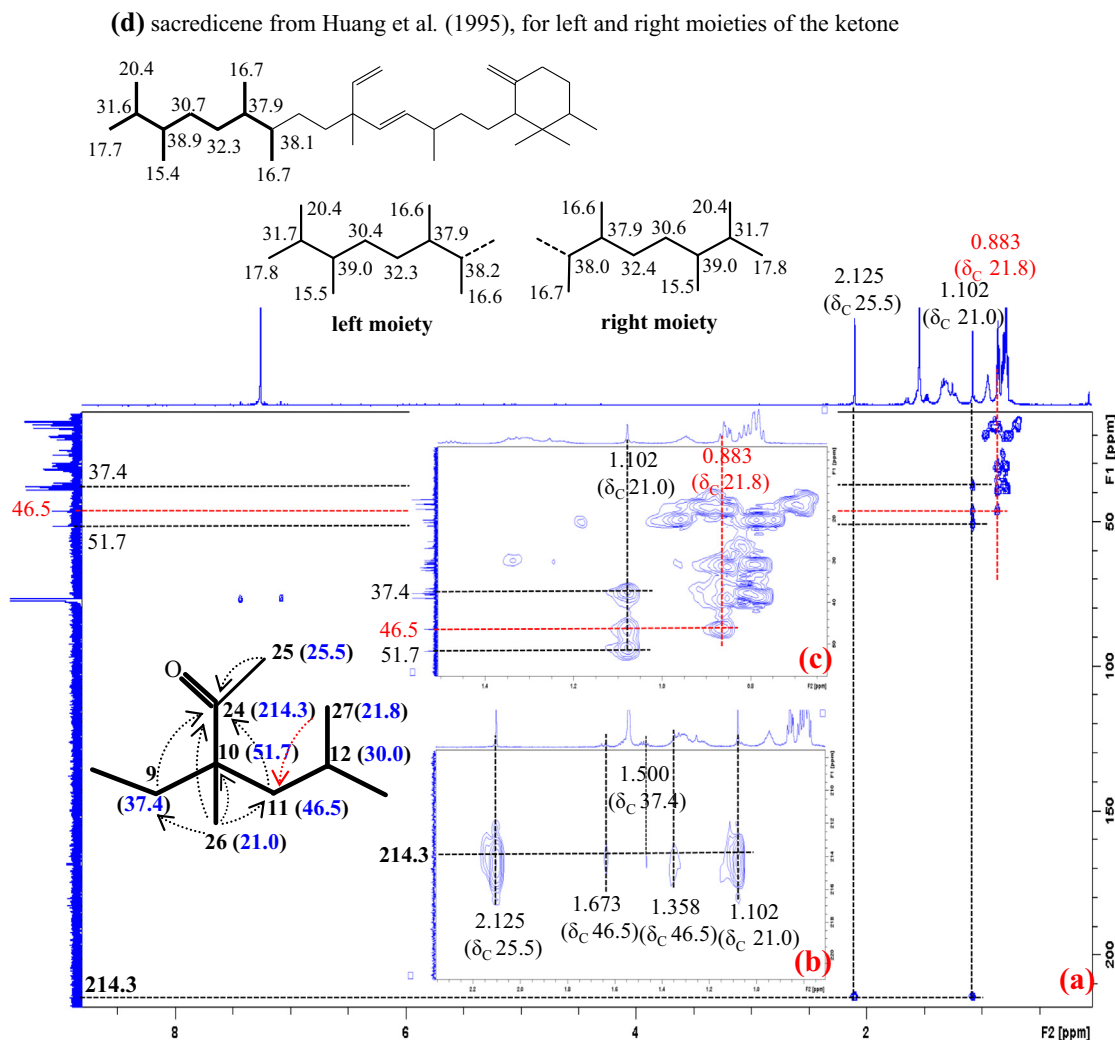


Fig. 4. Full (a) and zoomed-in partial (b and c) HMBC spectra of the identified C_{33} botryococcan-24-one showing the assignment of the carbonyl group to C-10 and a methyl group to C-12. The position-specific chemical shifts of the left and right moieties of the C_{33} botryococcan-24-one identified here (d) are also very close to that of the equivalent substructure of the sacredicene identified by Huang et al. (1995).

such a strong correlation (Fig. 4). The smaller values of δ_C at 21.0 ppm and δ_H at 1.102 ppm is consistent with the fact that C-26 is two bonds away from the carbonyl carbon. Therefore, the skeleton of the C_{33} botryococcanone can be established in which the carbonyl group is positioned on the ethyl group connected to the sole quaternary carbon (C-10), different from that of the botryococconone identified by Summons and Capon (1991) where it is located on C-15.

Next, we explain why there should be a methyl group connected to a carbon β to C-10 (Fig. 3b), rather than γ to C-10. As C-11 (δ_C 46.5) is correlated with H-27 (δ_H 0.883), it is reasonable to position the C-27 methyl (δ_C 21.8) on C-12. Such a positioning successfully explains why C-9 and C-11 differ in their δ_H (up to 0.3 ppm) and δ_C (up to 9 ppm) and why H-9 is a singlet (δ_H 1.500) while H-11 is a doublet (δ_H 1.673 and 1.358). The δ_H and δ_C difference between C-9 and C-11 should not come from the C-24 carbonyl oxygen as, according to Lindeman and Adams (1971), Grant and Paul modelling (Breitmaier and Voelter, 1978), the δ_C difference between carbons β to a $-\text{COCH}_3$ group is 1 ppm. As the two substructures left to C-8 and right to C-14 are symmetrical and that their position-specific δ_C values are very close to a substructure of the sacredicene reported by Huang

et al. (1995) (Fig. 4d), it is reasonable to assign the structure of C_{33} ketone to C_{33} botryococcan-24-one, with a methyl group β to C-10.

To further confirm the new skeletons of C_{33} botryococcanone and botryococconone in this study, we compared the Lindeman-Adams (Lindeman and Adams, 1971) modelled and measured carbon chemical shifts (Fig. 5) for the proposed new structures and that suggested by Brassell et al. (1986). The position-specific measured δ_C values (Fig. 5c and e) were very close to the calculated (empirical) values (Fig. 5b and d). By contrast, the measured δ_C values at C-11 (46.5 ppm) and C-16 (37.9 ppm) for the newly proposed skeleton **15** (Fig. 5e) differ significantly from the calculated δ_C values (C-11 (36.3 ppm) and C-17 (32.5 ppm)) for the suggested skeleton **15** (Fig. 5a) (Brassell et al., 1986).

3.2. Possible pathways for the formation of C_{33} botryococconone and botryococcanone

B. braunii is a phototrophic microalga usually found in surface waters. The commonly accepted pathway (A of Fig. 6) for the synthesis of (normal) botryococcones (**6**) in *B. braunii* starts with the condensation of two farnesyl diphosphates (FPP, **1**) to form

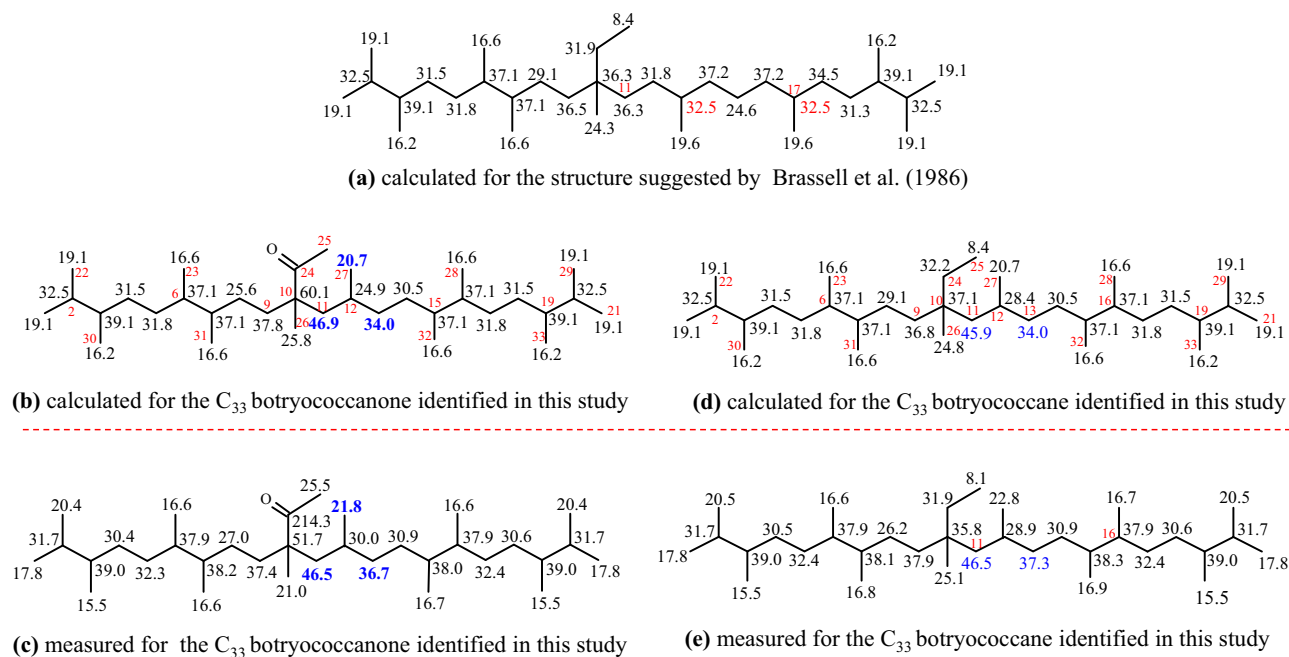


Fig. 5. The measured (c, e) and the calculated (b, d) δ_c values for the newly identified C₃₃ botryococcanone and C₃₃ botryococcan-24-one. For comparison, the Lindeman-Adam modelling was also applied to the skeleton (a) proposed by Brassell et al. (1986). The position-specific chemical shift differences (Z_{iso}) brought by the substituent group $-\text{COCH}_3$ (Breitmaier and Voelter, 1978) for the modelled positions C-8, C-9, C-10, C-11, C-12 and C-26 of between the C₃₃ botryococcanone and C₃₃ botryococcane were -3.5 , 1.0 , 23.0 , 1.0 , -3.5 and 1.0 ppm, respectively.

presqualene diphosphate (PSPP, **2**). The condensation proceeds with the formation of a cyclopropylcarbinyl cation intermediate (**3**) by the attack of C1' of one molecule of FPP (**1**) to the C-2 double bond of another FPP (Rilling and Epstein, 1969; Rilling et al., 1971). Migration of C1'-3 and C1'-2 to C1'-1 and subsequent reduction by NADPH leads to the formation of squalene (not shown here; Van Tamelen and Schwartz, 1971; Coates and Robinson, 1972; Zhang and Poulter, 1995; Jarstfer et al., 1996; Blagg et al., 2002) while a reductive C1'-2 fission of the cyclopropane ring of **3** with NADPH leads to the formation of normal C₃₀ botryococcene (**5**) with a C1'-3 linkage between the two FPPs (Huang and Poulter, 1989; Poulter, 1990). In this pathway, the formation of a botryococcenol with an $-\text{OH}$ group located at various positions (hydroxybotryococcene, HBO; **4** as an example) as stable products or unstable intermediates is highly likely since in vitro cultivation experiments (Zhang and Poulter, 1995; Jarstfer et al., 1996, 2002; Pan et al., 2015) with recombinant squalene synthase (SQase) showed that HBO is a stable product although it is yet to be isolated from wild *B. braunii* populations. It is also highly likely that the often detected botryococcenones (**5** and **6**) from laboratory cultured or wild *B. braunii* are the in vivo dehydration product of HBO as the dehydration and subsequent reduction of alcohol by SQase has been confirmed by Pan et al. (2015), even though the experiment was conducted with bacteria. Successive methylation of **5** with S-adenosylmethionine-dependent methylase (SAM) can introduce up to 7 methyl groups (Metzger et al., 1985b, 1986) to give botryococcenones in the carbon range of C₃₀–C₃₇ (**6**). The C1'-3 linkage means that no methyl group α or β to the quaternary carbon C-10 is possible as none of the α or β positions is methylated. Since the here identified C₃₃ botryococcane (**15**) and C₃₃ botryococcanone (**13**) both have a methyl group positioned β to the quaternary carbon C-10, it calls for a pathway different to the current one (**A** in Fig. 6) to explain the presence of these two new compounds in the Maoming Basin.

Here we propose a biosynthetic pathway (**B** in Fig. 6) leading to the synthesis of botryococcene with this unique skeleton followed by a geochemical pathway (**C** of Fig. 6) that converts them into the identified C₃₃ botryococcanone and botryococcanone. The pathway starts with the condensation of two FPPs (**1** in Fig. 6) by two nucleophilic attacks (one from the double bond between C-2 and C-3 to C1' and the other from the double bond C-2' and C-3' to C-3) leading to the formation of a cyclobutylcarbinol cation (**7**, 1'-2-3-2' cyclisation) with the positive charge located on the tertiary carbon (C3'). The cyclobutylcarbinol cation (**7**) can be also indirectly formed from the cyclopropylcarbinyl cation intermediate (**3**) (Thulasiram et al., 2008, see Fig. 6). Subsequent C1'-2' opening of the cyclobutane ring caused by electron migration shifts the positive charge to C1' (**8**). A Retro-Prins reaction (Al-Qallaf et al., 2000) could then give rise to the formation of C₂₉ botryococcene (**10**) (Fig. 6) directly from **8** with the loss of a hydroxymethyl cation (**9**). Methylation of **10** leads to the formation of C₃₃ botryococcene (**11**).

As the surface water is most likely to be oxic and can be penetrated by sunlight, once released to the surface water following algal death or active excretion, photo-mediated oxidation of **11** is likely. Indeed, Rontani and co-workers (e.g., Rontani et al., 1987, 2011; Rontani and Giusti, 1988; Rontani, 2001) provided evidence for photo-mediated oxidation of unsaturated lipids in the water column although the exact mechanism for such oxidation is still debatable. We suggest that the first product of (photo)oxidation is C₃₃ botryococcan-24-one (**12**) as the ethenyl group of **11** is more easily oxidised (to a ketone) than those backbone-connected double bonds in the skeletons. Deeper in the water column or in the sediment, this oxidative process is expected to be gradually overtaken by reduction of the double bonds in **12** due to the decrease in oxygen and light levels, leading to the here identified C₃₃ botryococcanone **13**. Further reduction of **13** converts it to a hypothetical C₃₃ botryococcanol (**14**). Subsequent dehydration and

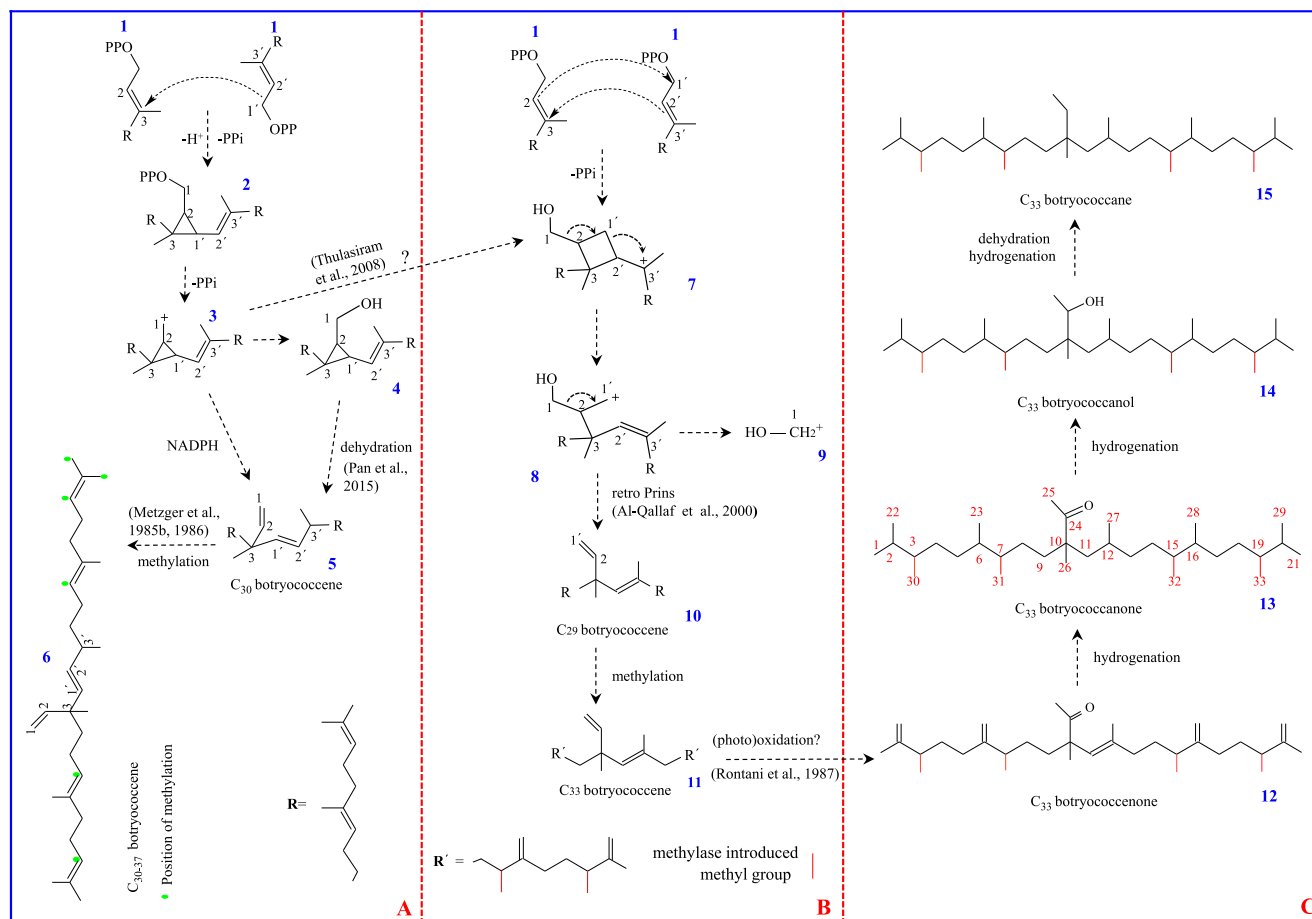


Fig. 6. Proposed biosynthetic and geochemical pathways for the formation of the new C_{33} botryococcane and C_{33} botryococconone with unique carbonyl group location and methyl group positioned β to the quaternary carbon in the skeleton. **A** is the currently accepted biochemical pathway for biosynthesis of normal botryococcenes; **B** is the proposed biochemical pathway for the formation of the C_{33} botryococcene; **C** is the geochemical pathway for the formation of the C_{33} botryococconone and botryococcane. The proposed pathway does not exclude other possibilities that may lead to the formation and preservations of these two new botryococconoids in the Maoming Basin.

hydrogenation of **14** leads to the formation of C_{33} botryococcane (**15**).

4. Conclusions

Accepted (normal) botryococcene biosynthetic pathways involve the formation of a cyclopropylcarbiny cation intermediate and predicts the absence of a β -positioned methyl group to the quaternary carbon in the skeleton. This prediction is, however, inconsistent with the firm structural establishment (1D and 2D NMR techniques, and Lindeman-Adams modelling) of a novel C_{33} botryococcane and a co-occurring C_{33} botryococcon-24-one with a uniquely positioned methyl group (β to the quaternary carbon) isolated by preparative gas chromatography.

We propose a biosynthetic pathway where direct or indirect cyclobutanation ($1'-2-3-2'$ cyclisation) is a necessary step leading to the synthesis of C_{33} botryococcene with the unique skeleton of the identified alkane and ketone from two FPPs: electron migration induced $C1'-2'$ opening of the cyclobutane ring and subsequent Retro-Prins reaction with a loss of a hydroxymethyl cation gives rise to the formation of a C_{29} botryococcene. The C_{33} botryococcene formed by successive methylation in vivo could then be (photo)oxidised to a C_{33} botryococconone in (oxic) shallow water, and

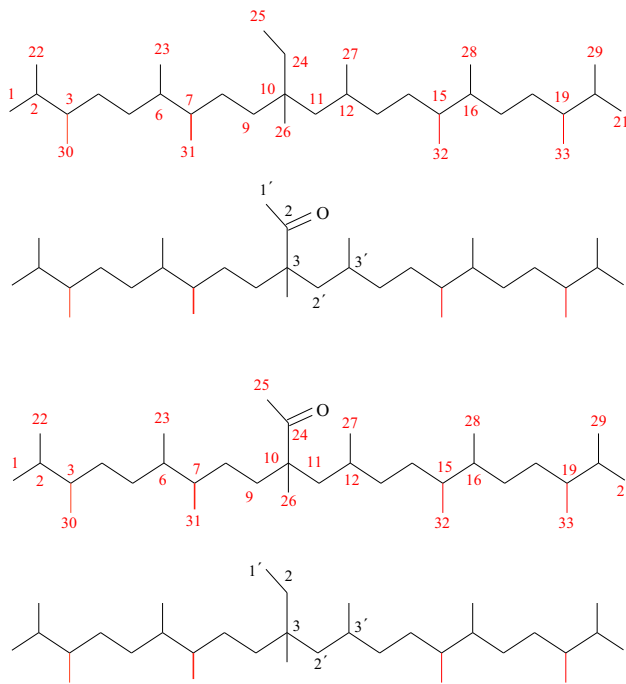
reduced (hydrogenated) to a C_{33} botryococcane in (anoxic) deep-water or sediments.

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Appendix A

Carbon numbering used for interpretation of NMR data and mass spectra, and speculation about the biosynthetic pathway for the C_{33} botryococcane and C_{33} botryococconone.



(a) The numbering of carbons in the C₃₃ botryococane as used for interpreting the NMR data and mass spectrum.

(b) The numbering of carbons in the C₃₃ botryococane as used in the tentative biosynthetic pathway (Fig. 6).

(c) The numbering of carbons in the C₃₃ botryococanone as used for naming and interpreting the NMR data and mass spectrum.

(d) The numbering of carbons in the C₃₃ botryococanone as used in the tentative biosynthetic pathway (Fig. 6).

Appendix B. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <https://doi.org/10.1016/j.orggeochem.2018.07.005>.

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