

Origin and Geochemical Implications of Hopanoids in Saline Lacustrine Crude Oils from Huanghekou East Sag and Laizhouwan Northeastern Sag, Bohai Bay Basin

Congkai Niu, Dujie Hou,* Xiong Cheng, Xu Han, Yan Li, and Yaxi Li



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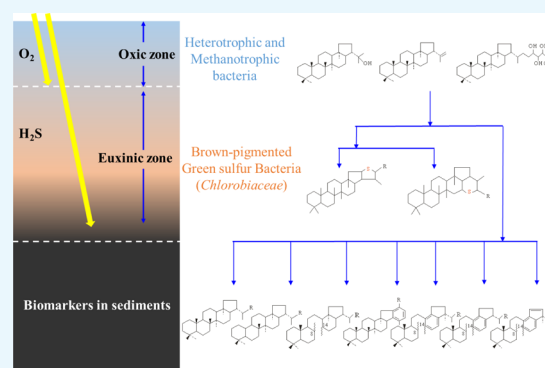


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ABSTRACT: A suite of low-mature crude oils (five high-sulfur oils and six low-sulfur oils) from the Huanghekou and the Laizhouwan Sags, Bohai Bay Basin, are analyzed to investigate the fate of the hopanoids. Abundant hopanes, such as secohopanes, 25-norhopanes, benzohopanes, aromatized secohopanes, and sulfide hopanes, are identified, and their carbon isotope compositions are determined. Varying ^{13}C isotope values of C_{31} hopane (-38.7 – -34.0‰) and C_{29-30} hopanes (-38.5 – -31.5‰) suggest different bacterial sources of these compounds. The presence of 25-norhopanes with enriched heavy carbon isotopes in severely biodegraded oils suggests that they are microbially mediated products. The detection of the isotopically depleted C_{29} and C_{30} D-ring-8,14-secohopanes (-45.6 – -41.2‰) indicates that secohopanes are from methane-oxidizing bacteria (methanotrophs). The presence of isorenieratane, lower aryl isoprenoid ratios, and a good correlation between the sulfur content and the gammacerane index indicate the presence of green sulfur bacteria (Chlorobiaceae) under photic zone euxinic conditions. Water column stratification results in good preservation of the organic matter, and it is in favor of diversity of aquatic microorganisms. The ratios of $\text{C}_{35}/\text{C}_{34}$ sulfide hopane, C_{35} sulfide hopane-2/ C_{35} sulfide hopane-1, and $\text{C}_{35}/\text{C}_{34}$ benzohopane are influenced by the reducing environments in this region. In addition, the D-ring monoaromatized 8,14-secohopanoid/(D-ring monoaromatized 8,14-secohopanoid + benzohopanes) and C_{31} – C_{35} secomoretanans/secohopanes are affected by the maturity. We hypothesize that the reducing environments and thermal effects are important markers for the hopanoid transformation, including the incorporation of inorganic sulfur in substituting functional groups, cyclizing, aromatizing, and opening ring C of the hopanoids.



1. INTRODUCTION

Hopanoids are ubiquitous in sediments and crude oils and are an important indicator for bacteria.^{1–3} Previous research studies have shown that hopanoids are more resistant to biodegradation,^{4–7} natural weathering processes,^{8,9} and thermal effects^{10,11} than *n*-alkanes, isoprenoids, and bicyclic sequiterpanes. In addition, the spatial configuration of hopanoids regularly evolves from the initial biological isomers to the following geological counterparts.¹² Therefore, many parameters of hopanoids are used to reflect the depositional environment,¹³ thermal maturity,¹⁴ and degree of biodegradation.¹⁵ Hopanoids have been used as molecular markers for oil–oil and oil–source rock correlation in petroleum geochemistry. A major source of hopanoids $> \text{C}_{30}$ in sediments is formed from the cleavage of the extended side chain of C_{31} – C_{35} bacteriohopanepolyols in some cyanobacteria and heterotrophic and methanotrophic bacteria.^{16–18} The number, position, and nature of functional groups in the extended side chain might represent different species of bacteria.^{19–22} For example, hopanoids with five functional groups at the side chain and an additional hydroxyl group at the C_{30} or C_{31}

position were found in cyanobacteria²⁰ and methanotrophic bacteria (with an amino group at C_{35}).²² Hopanoids $< \text{C}_{30}$ are mainly formed from diplopterol and diploptene without a long side chain at C-22, which can be found in almost all hopanoid-producing bacteria.²³ Because of the comprehensive impact of the origin, depositional environment, and diagenetic effects, the hopanoids occur in many novel types with various structures, such as hopenes,^{24,25} isohopenes,^{26,27} rearranged hopanes,^{28–32} aromatized hopanes,^{33,34} sulfur-containing hopanes,^{34–40} and nitrogen-containing hopanes.^{41,42} Intriguingly, most hopanoid precursors were found in aerobic bacteria,²³ although their biosynthesis does not require oxygen. However, some hopanoids were also found in anaerobic bacteria, such as *planctomycetes*,⁴³ *Petrosia* sp.,⁴⁴ *Desulfovibrio*,⁴⁵

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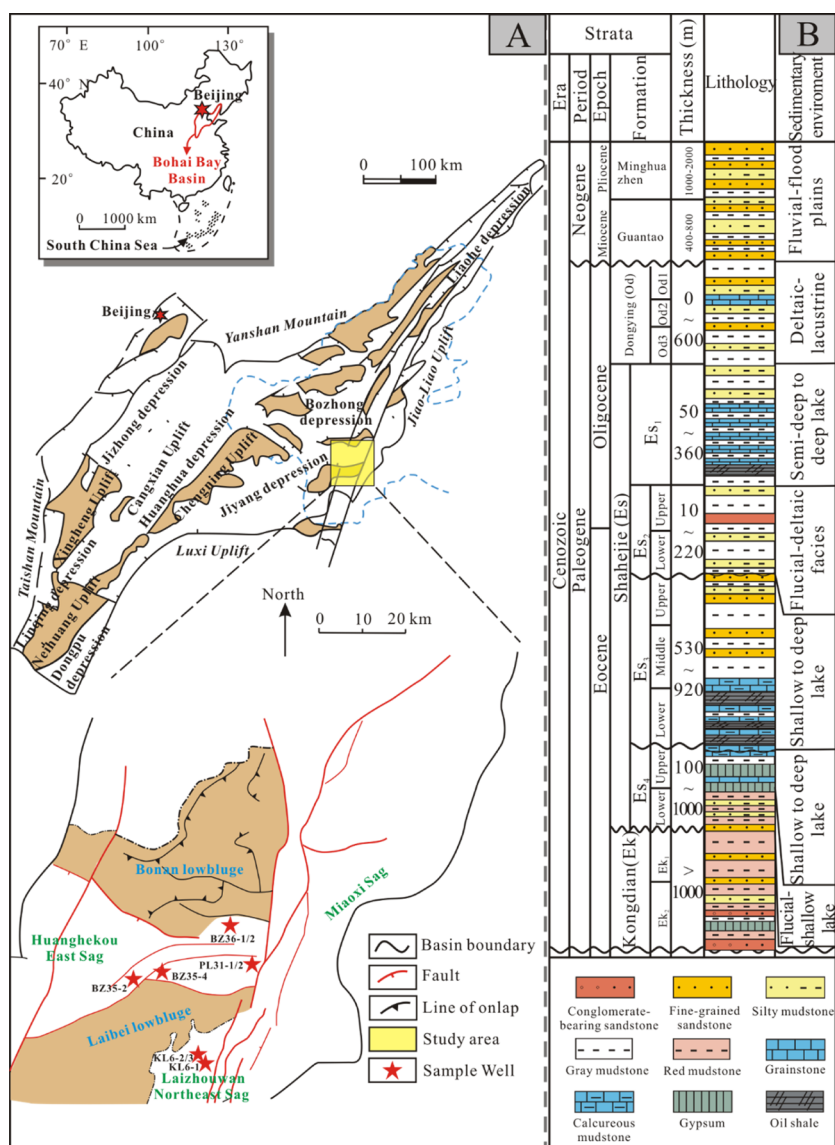


Figure 1. (a) Regional tectonic units of the Bohai Bay Basin, Huanghekou East Sag, and Laizhouwan Northeast Sag within the Bohai Bay Basin, structural units of the study area, and location of wells sampled here and (b) generalized lithology, stratigraphy, and depositional environment of the Bohai Bay Basin.

and *Rhodoplanes* spp.⁴⁶ Many novel hopanoids were reported following the development of technology. However, the origin and pathway of the diagenetic evolution of some novel hopanoids are obscure. Clearly, a deeper understanding of the fate of hopanoids during diagenesis needs more data from field and laboratory studies.

Compound-specific isotope analysis is the effective method to distinguish the organic matter with specific sources including chlorophyll, cuticular waxes, and bacteria.^{47–49} Stable carbon isotopic fractionation may be determined by the origin and fate of some compounds in the sedimentary environment.^{50,51} Compound-specific isotope analysis has been used for oil–oil and oil–source correlations.^{52,53} The carbon isotope of an individual hopane has been reported to have a special origin, including homohopane, homohop-17(21)-enes, benzohopanes, neohop-13(18)-enes, and aromatized hopanoids.^{14,24,54–56} Thermal maturation, depositional environment, and mixture of generated hydrocarbons might stymy the source analysis of petroleum.⁴⁹ However, the

fractionation effects in diagenetic reactions likely impact the carbon isotope values of the precursor molecules.⁵¹

Abundant hopanoids including hopanes, secohopanes, benzohopanes, aromatized secohopanes, and sulfide hopanes were identified in a suite of low-maturity crude oils from the Huanghekou East Sag and the Laizhouwan Northeastern Sag. Therefore, the analysis of the evolution of hopanoids under reducing lacustrine environments is possible in the study area. The aims of this study are to show the origin of hopanoids in reservoir oils using compound-specific carbon isotopic analysis to illustrate the fate of hopanoids sourced from different depositional environments and to predict the coeval growing environment of the dominant aquatic microorganism.

2. GEOLOGICAL SETTING

Bohai Bay Basin in East China (Figure 1) is among one of the most petroliferous basins, with an area of about 4.2×10^4 km².^{57,58} The Bohai Bay Basin is an intracratonic rift basin that has experienced two major tectonic evolution stages including

Table 1. Biodegradation Level and Bulk Geochemistry Characteristics of the Investigated Lacustrine Oils from Bohai Bay Marginal Sags

sample ID	depth (m)	formation	PM level	sulfur content (%)	bulk oils $\delta^{13}\text{C}$ (‰)	fractions content (%)			
						saturates	aromatics	resins	asphaltenes
Laizhouwan Northeastern Depression									
KL6-1	1374.5–1440	N ₁ m ^L	2	nd	–26.3	57.58	26.26	8.85	3.23
KL6-2	2153–2181	E ₃ d ₂ ^L	1	0.31	–27.4	60.00	20.26	10.77	2.56
KL6-3	2561–2565	E ₃ s ₂	1	0.4	–27.3	57.35	20.41	13.47	3.47
KL6-4	2585–2603	E ₃ s ₂	1	0.52	–27.6	56.30	21.75	13.01	6.10
Huanghekou Eastern Depression									
BZ35-2	2685.5–2703.5	E ₃ d ₃	1	1.4	–26.8	41.75	24.50	15.00	17.25
BZ35-3	2912.2–2930	E ₃ s ₂	1	nd	–26.6	65.15	16.92	7.58	14.65
BZ36-1	1297–1321.5	N ₁ m ^L	5	2.95	–26.8	32.38	37.14	14.29	9.52
BZ36-2	1552–1570	N ₁ g	2	2.07	–26.5	37.17	34.87	15.46	7.24
PL31-1	1507–1529.5	N ₁ g	3	1.12	–26.1	46.10	30.50	11.35	4.61
PL31-2	1241.5–1259	N ₁ g	5	1	–26.1	43.62	36.38	14.47	2.77
BZ35-4	2449–2468.5	E ₃ d ₂ ^L	1	0.57	–26.2	58.53	24.04	10.46	3.30

the Paleogene syn-rift stage and the Neogene-Quaternary post-rift stage.^{59,60} A series of grabens and half-grabens developed along major NW- and NE-trending fault sets during the syn-rift stage.^{61,62} Three sedimentary sequences from bottom to top include the Paleogene Kongdian Formation (Ek), the Shahejie Formation (Es), and the Dongying Formation (Ed), which were deposited in the syn-rift stage (65–24.6 Ma).^{63,64} During the post-rift stage, the whole basin experienced a thermal subsidence, and the sedimentary sequences consist of the Neogene Guantao Formation (Ng), the Minghuazhen Formation (Nm), and the Pingyuan Formation (Qp) from bottom to top.⁶⁵

The Huanghekou East Sag and the Laizhouwan Northeastern Sag are located in the south area of the Bohai Sea.^{29,66,67} The two sags are separated by the Laibei low uplift and are controlled by NNE and NWW-EW striking extension faults.⁶⁸ The Huanghekou Sag is one of the margin sags in the Bohai Bay Basin and is a typical fault depression with a north fault and a south slope, which is bounded by structural units including the Bonan low uplift to the north, the Miaoxi South Sag, and the Laibei low uplift to the east and south.⁶⁹ The west branch of the Tanlu Fault passes through the middle of the Huanghekou Sag, separating the sag into the west sag and the east sag (Figure 1). The Palaeogene and Neogene strata are composed of the Ek, Es, Ed, Ng, and Nm in younger ages, of which Es and Ed are excellent-quality source rocks.⁶⁹ The good-quality source rocks in the Huanghekou East Sag are mainly controlled by tectonic, paleo-climate, paleo-productivity, and preservation condition of the organic matter.⁶⁹ The hot and humid climate is interpreted to cause the high level of productivity of algae and relative high organic contents in the source rocks in Es₃ in a semideep to deep lake environment. The Es_{1–2} source rocks are deposited in the environment of lower temperatures and are associated with algal blooming and have a high level of paleo-productivity. The Ed₁ source rocks with mixed sources of organic matter including algal and terrigenous plant matter are deposited in a warm and semihumid climate.⁶⁹ The reducing environments are in favor of the preservation of organic matter.⁶⁹ The Laizhouwan Northeastern Sag is also a typical marginal sag and it is formed by the uplift during the Ek-Es₄ stages.⁷⁰ The fluctuation of subsidence and uplift after the Es₄ stage results in large amounts of deposits, including Es₃, Es₂, Es₁, and Ed.⁷⁰ The organic sources of the Es₃ member are mainly lower aquatic

organisms and prokaryotes. The bottom water conditions fluctuate among the stratified, anoxic conditions and freshwater, weak reducing conditions.⁷¹ The organic matter in the Es₂ member is derived from less aquatic organisms in more saline and anoxic bottom water than that in the Es₁ member.⁷¹ Shales in the Ed₃ member are characterized by abundant terrigenous organic matter in the less saline and weaker reducing depositional conditions.⁷¹

3. SAMPLES AND METHODS

Full details have been reported elsewhere⁷² but are summarized as follows.

3.1. Samples. A suite of 11 crude oils were analyzed in this study (Table 1). Four samples were from the Laizhouwan Northeastern Sag and seven samples were from the Huanghekou East Sag. A detailed analysis of aliphatic hydrocarbons was presented in the study of Niu et al.⁷²

The samples obtained from the Laizhouwan Northeastern Sag were slightly biodegraded, corresponding to level 1 based on the Peters and Moldowan⁶ biodegradation scale (abbreviated as PM 1). The sulfur contents of these oils were lower than 0.5%. There were five high sulfur oils (HSOs: BZ35-2, BZ36-1, BZ36-2, PL31-1, and PL31-2) and two low sulfur oils (LSOs: BZ35-3 and BZ35-4) obtained from the Huanghekou East Sag (Table 1). The HSOs experienced a slight to heavy biodegradation with the PM level from 1 to 5, while LSOs were on PM level 1.

Isorenieratane was identified in HSOs, illustrating a photic zone euxinic paleoenvironment.⁷³ Meanwhile, the Gammacerane index (0.08–0.23, average 0.16), C₃₅S/C₃₄S hopane (0.79–1.10, average 0.91), and the ETR index (0.36–0.44, average 0.40) of HSOs represented brackish and saline water conditions. Furthermore, the organic matter mainly originated from algae and bacteria, and the maturity of all samples was in the low-mature to mature stage according to the identification of aliphatic biomarkers.⁷²

3.2. Separation of Aliphatic Hopanes, Benzohopanes, Secohopanes, and Sulfidic Hopanes. The details of the separation methods of aliphatic hopanes, benzohopanes, secohopanes, and sulfidic hopanes were described by Cheng et al.,⁷⁴ Jiang et al.,⁷⁵ Wang et al.,⁷⁶ and Wu et al.,⁷⁷ respectively. The methods were introduced briefly in the following.

The hopanes were mainly separated from the aliphatic fractions. First, the asphaltene was precipitated with excess *n*-hexane, and then the maltenes were separated into aliphatic and aromatic fractions and resins by column chromatography with *n*-hexane, *n*-hexane:dichloromethane (1:2, V/V), and dichloromethane as rinsing solvents, respectively. The column chromatograph (15 cm length and 6 mm i.d.) was filled with activated alumina. The alumina (neutral, 100–200 mesh) were activated at 450 °C for 5h. The saturated fraction was further separated into purified *n*-alkanes and branched/cyclic alkanes by urea adduction. The hopanes were all enriched in the second fraction for the analysis of individual carbon isotopes.

The benzohopanes and aromatic secohopanes were all enriched in the monoaromatic fraction. A total of aromatic hydrocarbons (ca. 1–10 mg) in a minimum volume of *n*-hexane were carefully introduced to the top of the prepared column chromatograph (15 cm length and 6 mm i.d.) over activated alumina (450 °C for 5h). The sample container was rinsed three times with 100 μ l *n*-hexane to fully transfer the aromatic hydrocarbons into the column. Then, *n*-hexane:dichloromethane (99:1, V/V, 5.5mL) was used to elute the monoaromatic hydrocarbons. Because of the volatility of the reagents, this experiment was best carried out at a room temperature of 25 °C.

The sulfide hopanes were separated by the methylation–demethylation method. Briefly, the crude oils (1 g) were dissolved in 5 mL of dichloromethane, and silver tetrafluoroborate (5 mol equiv) and methyl iodide (5 mol equiv) were added into the solvent. After that, the solution was stirred in the dark at room temperature for 24 h. The reaction products were centrifuged to separate the supernatant and precipitates, and the precipitates were removed by filtrating and evaporating dichloromethane using a rotating evaporator. The oily residue was mixed with 10 mL of *n*-hexane and kept in a refrigerator. Sulfonium salts were isolated by decanting the *n*-hexane supernatant after precipitation and centrifugation, and this procedure was repeated three times. The sulfonium salts were dissolved in 5 mL of acetonitrile and 10 mol equiv of 7-azaindole was added afterward, and then the solution was stirred in darkness for 24 h. The thiophenic fraction was extracted via *n*-hexane (10 mL, 3 times). 10 mol equiv of 4-dimethylaminopyridine was added to the residue and refluxed for 12 h. The sulfidic fraction was extracted by *n*-hexane (10 mL, 3 times).

3.3. Gas Chromatography–Mass Spectrometry Analysis. GC was performed on an HP6890 gas chromatograph equipped with a DP-5 MS-fused silica column (60 \times 0.25 m i.d. and film thickness 0.25 μ m) coupled to an HP 5973 mass selective detector (the ionization source operates at an electron beam energy of 70 eV). The aliphatic and aromatic fractions were analyzed in the same program in which the column was held at 40 °C for 2 min, which was then increased to 310 °C at 4 °C/min, with a final hold of 20 min. The program of sulfidic fractions was the same as that of saturate fractions, but the final hold was extended to 40 min. Helium was used as the carrier gas at a constant flow rate of 1.0 mL/min.

3.4. Gas Chromatography–Isotope Ratio Mass Spectrometry. Carbon isotopic analyses of *n*-alkanes, isoprenoids, and monoaromatic hydrocarbons were performed on a Thermo Scientific Delta V Advantage stable isotope ratio mass spectrometer system interfaced to a Trace Ultra gas chromatograph (Thermo Scientific; J&W DB-5 fused silica column: 30 \times 0.25 m i.d. and 0.25 μ m film thickness). The GC

oven temperature for the analysis of isotopic hydrocarbons was initially held at 50 °C for 2 min and increased to 310 °C at 1 °C/min, with a hold of 20 min. The GC oven temperature for the analysis of monoaromatic hydrocarbons was held at 50 °C for 2 min, increased to 290 °C at 2 °C/min, and then ramped to 310 °C at 1 °C/min, with a final hold of 20 min. The temperature of the isotope ratio mass spectrometry oxidation oven was 1000 °C. A standard mixture of *n*-alkanes with known ¹³C isotope values was analyzed over five samples to check the accuracy of the instrument. The isotopic compositions of all samples were relative to VPDB and were determined at least twice with a standard deviation for *n*-alkanes of $\pm 0.8\%$.

4. RESULTS

4.1. Hopanes in the Aliphatic Fraction. 4.1.1. Hopanes.

Hopanes are derived from bacterial membrane precursors and are widely distributed in sediments and petroleum.^{78–80} The integrated distribution of hopanes is shown in Figure 2. C₃₀

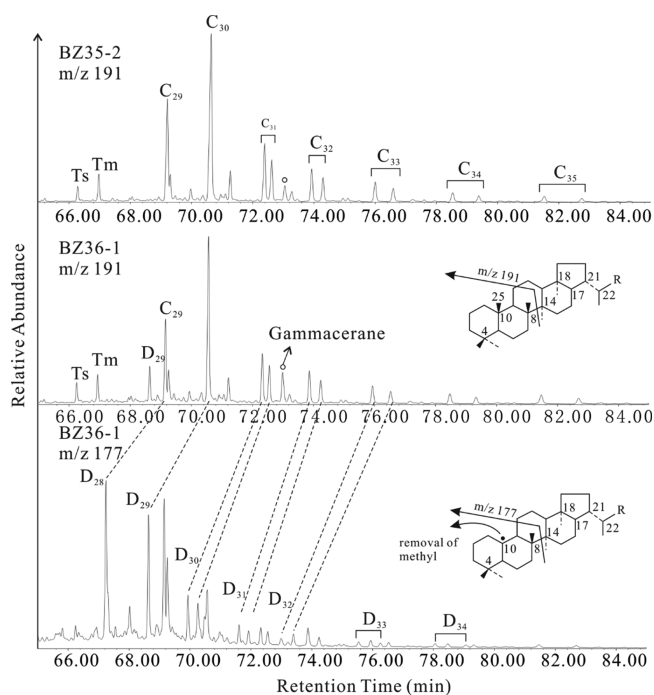


Figure 2. Representative mass chromatograms (m/z 191 and 177) showing the distribution of hopanes and 25-norhopanes in BZ35-2 and BZ36-1 samples. Ts: 8 α (H)-22,29,30-trinorhopane; Tm: 17 α (H)-22,29,30-trinorhopane; C₂₉–C₃₅: C₂₉–C₃₅ hopanes; D₂₈–D₃₄: C₂₈–C₃₄ 25-norhopanes.

17 α (H), 21 β (H) hopane is the dominant compound among its homologues. Hopanes range from C₂₇ to C₃₅ without C₂₈. The abundance of C₃₅ hopanes in HSOs is obviously higher than that of LSOs. Meanwhile, a higher gammacerane index of HSOs suggests stratification of the sedimentary water body.⁸⁰ In addition, the abundance of 17 α (H)-22,29,30-trinorhopane (Tm) in all samples is higher than that of 18 α (H)-22,29,30-trinorhopane (Ts), indicating the low-mature stage of the samples.

Hopanes are more resistant to biodegradation than *n*-alkanes, isoprenoids, and bicyclic sesquiterpanes.⁷⁴ In heavily degraded oils, demethylation of hopanes occurs to form 25-norhopanes.^{15,81} The 25-norhopanes are identified in the

Table 2. Carbon Isotope of Hopanoids^a

sample ID	C ₂₉ H (‰)	C ₃₀ H (‰)	(22S + 22R) C ₃₁ H (‰)	D ₂₉ H (‰)	C ₃₂ BH (‰)	C ₃₃ BH (‰)	C ₃₄ BH (‰)	C ₃₅ BH (‰)	C ₂₉ SH (‰)	C ₃₀ SH (‰)
Laizhouwan Northeastern Depression										
KL6-1	-37.44	-35.21	nd	nd	nd	nd	nd	nd	nd	nd
KL6-2	-37.49	-37.74	-38.66	nd	-35.27	-36.00	-32.65	-32.15	-44.65	-44.99
KL6-3	-33.90	-34.34	nd	nd	nd	nd	nd	nd	nd	nd
KL6-4	-38.05	-38.53	nd	nd	nd	nd	nd	nd	nd	nd
Huanghekou Eastern Depression										
BZ35-2	-33.93	-36.33	-37.28	nd	-32.01	-32.39	-30.96	-35.58	-43.94	-43.26
BZ35-3	-33.38	-31.47	-34.00	nd	nd	nd	nd	nd	-41.15	-42.27
BZ36-1	-37.44	-37.56	-36.77	-35.71	-31.27	-32.98	-34.22	-35.28	-45.60	-42.77
BZ36-2	-35.41	-35.12	-37.88		-32.41	-32.04	-31.38	-32.80	-44.06	-42.77
PL31-1	-35.06	-35.93	-35.78	-35.79	-31.34	-31.31	-31.31	-30.88	nd	nd
PL31-2	-37.59	-36.99	-35.40	nd	nd	nd	nd	nd	nd	nd
BZ35-4	-34.66	-33.88	-36.62	nd	-31.80	-31.95	-30.63	-32.18	-43.90	-43.56

^aC₂₉H: C₂₉, 17 α , 21 β (H)-hopane; C₃₀H: C₃₀, 17 α , 21 β (H)-hopane; (22S + 22R) C₃₁H: (22S + 22R), C₃₁, 17 α , 21 β (H)-hopane; D₂₉H: C₂₉, 25-norhopane; C₃₂BH: C₃₂, benzohopane; C₃₃BH: C₃₃, benzohopane; C₃₄BH: C₃₄, benzohopane; C₃₅BH: C₃₅, benzohopane; C₂₉SH: C₂₉, 8,14-D-ring monoaromatic secohopane; C₃₀SH: C₃₀, 8,14-D-ring monoaromatic secohopane; nd = not determined.

heavily biodegraded oils including PL31-2 and BZ36-1, suggesting that 25-norhopanes are formed by microbial demethylation of hopanes. In this study, a minor peak of C₂₉ 25-norhopane (D₂₉) can be observed in the *m/z* 191 mass chromatogram, and hopanes could be identified in the *m/z* 177 mass chromatogram of PL31-2 and BZ36-1, suggesting incipient biodegradation of hopane compounds.⁷⁴

In order to determine potential diagenesis relationships, the ¹³C isotope values of the series of hopanes and 25-norhopanes are measured (Table 2). The ¹³C isotope values of C₂₉ and C₃₀ hopanes range from -38 to -31‰, while $\delta^{13}\text{C}$ values of most C₃₁ (22S + 22R) hopanes are more depleted than those of C₂₉ and C₃₀ hopanes, with the range of -39 to -33‰. On the contrary, in PL31-2 and BZ36-1, ¹³C isotope values of 25-norhopanes are enriched in ¹³C isotope values (-35.8 and -35.7‰, respectively).

4.1.2. 8,14-Secohopane. Isomers of 8,14-secohopanes can be identified (Figure 3) by comparing the relative retention time and mass spectra in previous studies.^{82,83} All of these marked peaks in these chromatograms represent compounds with mass spectra containing a base peak of *m/z* 123 and a molecular ion appropriate for secohopanes. The C₂₉ and C₃₀ 8,14-secohopanes are recognized by the molecular ions at *m/z* 400 and 414 and *m/z* 123, 193, and 267 fragment ions. The mass spectra of C₂₉ and C₃₀ 8,14-secohopanes are listed in the Supporting Information (Figure S1a,b). Furthermore, six pseudohomologous series extending to C₃₅ are identified mainly by the molecular ion at *m/z* 428, 442, 456, 470, and 484 and the base peak at *m/z* 191. These compounds can be classified into secohopanes and secometretanes. Owing to the ring C opening, some novel 8,14-secohopanes configured at the C-8, C-14, C-17, and C-21 positions are found. In most cases, the peaks occur as pairs from C₃₁ to C₃₅ presumably because of the occurrence of diastereomers due to the chiral center at C₂₂.⁸³ Therefore, the abundances of 8 α , 14 α , 17 α , 21 β (H)-secohopanes and 8 α , 14 β , 17 β , 21 α (H)-secometretanes are higher than other counterparts.

4.2. Hopanes in the Aromatic Hydrocarbon Fraction.

4.2.1. Benzohopanes. Benzohopanes are separated from monoaromatic fraction, showing a relatively high abundance (Figure 4). These C₃₂–C₃₅ benzohopanes are suggested to be formed by the side-chain cyclization at C₂₀, followed by the

aromatization.^{84,85} They occur widely in high abundances in sedimentary rocks and crude oils in carbonate-rich and evaporitic environments.^{33,84,85} The C₃₂–C₃₅ benzohopanes with distributions similar to those of C₃₂–C₃₅ hopanes have the same base peak at *m/z* 191 with molecular ions at *m/z* 432, 446, 460, and 474. The mass spectra of benzohopanes are listed in the Supporting Information (Figure S1c–f). However, the ¹³C isotope values of benzohopanes vary from -36 to -31‰ (Table 2) and are more enriched than hopanes.

Benzohopanes may be ubiquitous components, especially abundant in confined and reducing environments.^{86,87} The predominance of the C₃₄ counterpart among benzohopanes has been reported in evaporitic rock samples; however, the carbon isotope depletion of C₃₂ to C₃₅ benzohopanes in oils is tentatively attributed to migration effects of hydrocarbons.⁸⁴

4.2.2. D-Ring Monoaromatic SHs, DSHs, and IDs. The isomers of 8,14-secohopanooids (SHs), D-ring monoaromatic 8,14-Seco-28-nor-hopanooids (DSHs), and indenyltrimanes (IDs) are identified in the *m/z* 365, 351, and 363 mass chromatograms by comparison of the relative retention time with those in the literature studies.^{87,88} The *m/z* 365, 351, and 363 mass chromatograms show the monoaromatic fraction from the oil sample in well BZ35-2 (Figure 4). SHs have abundant C₂₉ and C₃₀ compounds, while the abundance of C₃₁ SH is in minute abundance. The ¹³C isotope values of C₂₉ and C₃₀ SHs vary from -45 to -41‰ (Table 2), which is more depleted than that of hopanes and benzohopanes. DSHs are the demethylated counterparts of regular SHs.⁸⁷ The C₂₉ DSH is the dominant constituent in DSHs and the peaks of C₂₈ to C₃₀ DSHs occur as pairs. IDs at *m/z* 363 are a series of aromatic secohopanooids in oils, which are counterparts of the regular SHs but containing an extra C=C bond in the E-ring.⁸⁷ The typical mass spectra of SHs, DSHs, and IDs are listed in the Supporting Information (Figure S1(g–i)).

4.3. Hopanes in Sulfidic Fraction. A homologous series of hexacyclic sulfides with a 17 α , 21 β (H) hopane carbon skeleton ranging from C₃₀ to C₃₅ are identified by Cyr et al.³⁶ in heavy oils of Northern Alberta. Schaeffer et al.³⁷ described the structural elucidation of sulfide hopanes and separated them into eight series. By comparing with the relative retention time and mass spectra,³⁷ the structures of compound-1 and compound-2 are represented in Figure 5a. The mass spectra

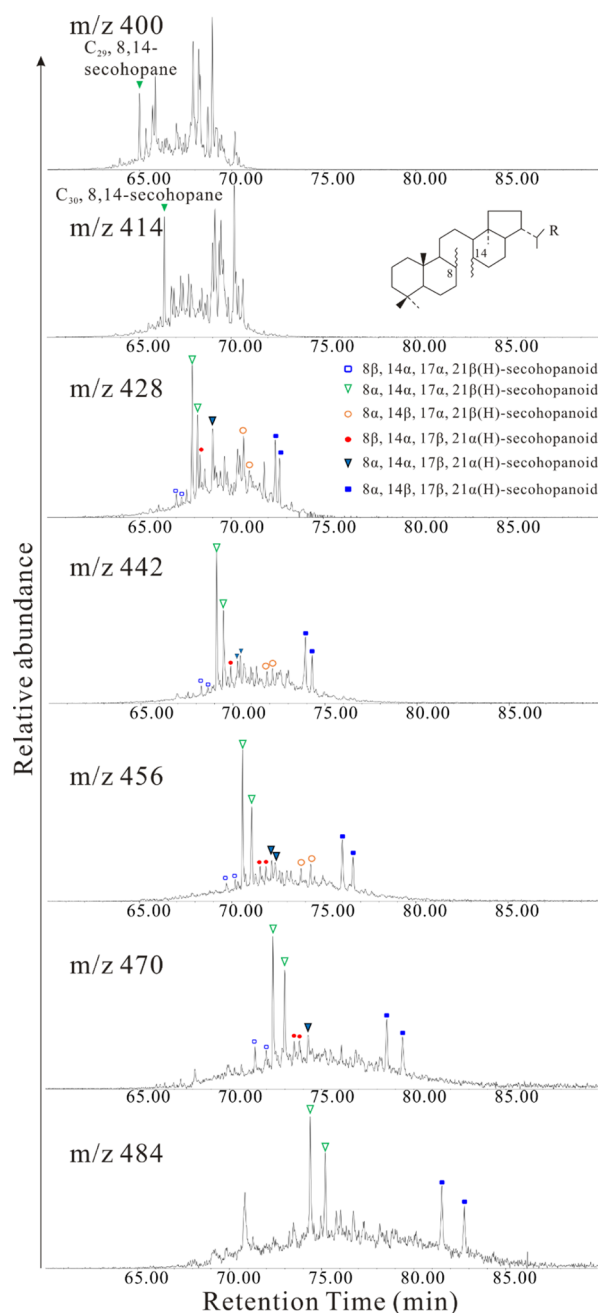


Figure 3. Representative mass chromatograms showing the distribution of 8,14-secohopanes from C_{29} to C_{35} in BZ35-2 samples.

are listed in the Supporting Information (Figure S1m–p). The spectra of the sulfides from these compounds show a base peak at the molecular ion and doublets of fragments at m/z 191 and m/z 235 (C_{31} homologue) + $14n$ due to the cleavage of ring C. The mass spectra of compound-1 and compound-2 (Figure S1m–p) show similarities; however, the presence of a fragment at m/z 122 in compound-2 with a higher intensity than in the case of compound-1 is a typical feature of these compounds.³⁷

A compound with a base peak at m/z 395 is co-eluted with compound-1 of C_{32} sulfide hopane ($C_{32}SH-1$, Figure 5b), resulting in a relatively high abundance of $C_{32}SH-1$. As shown in the mass spectrum of $C_{32}SH-1$ (Figure S1o), the fragments at m/z 395 and the molecular ion (m/z 484) are similar to the mass spectrum of compound B (Figure 5d), suggesting the co-elution of $C_{32}SH-1$ and compound B. The mass spectrum of

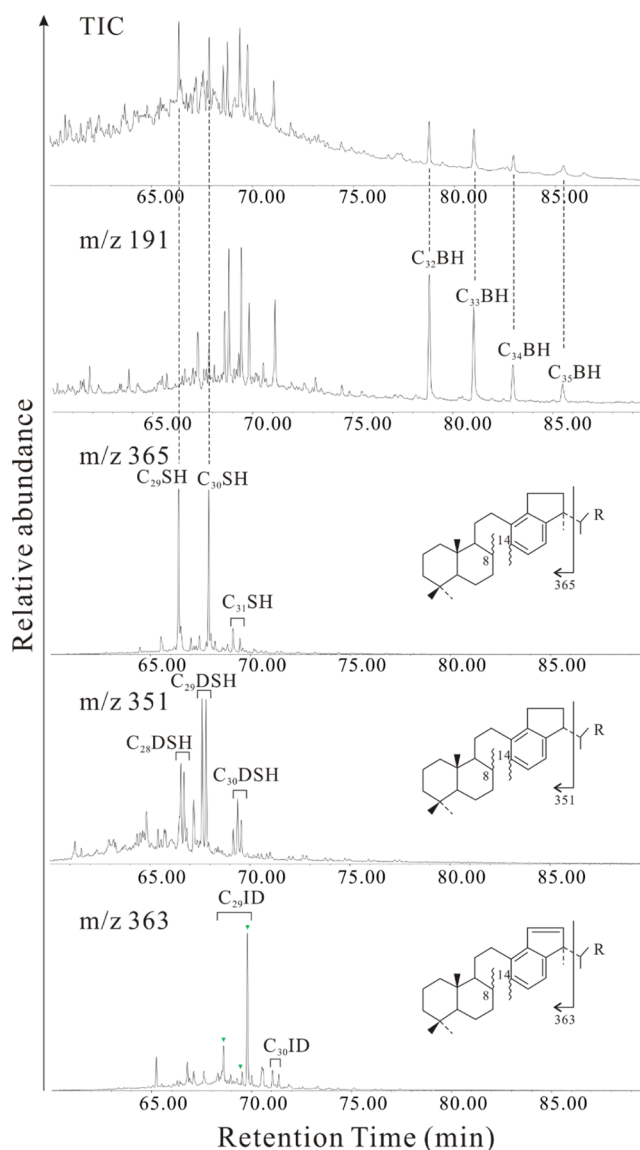


Figure 4. Representative mass chromatograms showing the distribution of benzohopanes (m/z 191), D-ring monoaromatic 8,14-secohopanoids (m/z 365), D-ring monoaromatic 8,14-seco-28-norhopanoids (m/z 351), and indenyltrimanes (m/z 363) in aromatic hydrocarbon of BZ35-2 samples.

$C_{32}SH-2$ is shown in Figure S1p, suggesting that $C_{32}SH-2$ is less influenced.

The mass spectra of compounds A–E are shown in Figure 5c–g. The spectra of these compounds show a base peak at m/z 395 and fragments at m/z 121, m/z 163, m/z 191, and m/z 75 (compound A homologue) + $14n$ ($0 \leq n \leq 4$). Compounds from A to E show molecular ions from M^+ 470 to 526. More details about the structure of these compounds are beyond the scope of this paper, which needs further study in future studies.

5. DISCUSSION

5.1. Sources of Hopanes. An important source of high-molecular-weight hopanoids ($>C_{30}$) is from the diagenetic transformation of bacteriohopanepolyols in membrane lipids in a wide range of prokaryotes including cyanobacteria and heterotrophic and methanotrophic bacteria.^{18,23,89} In addition, lower pseudohomologues (C_{30} or less) may be related to their

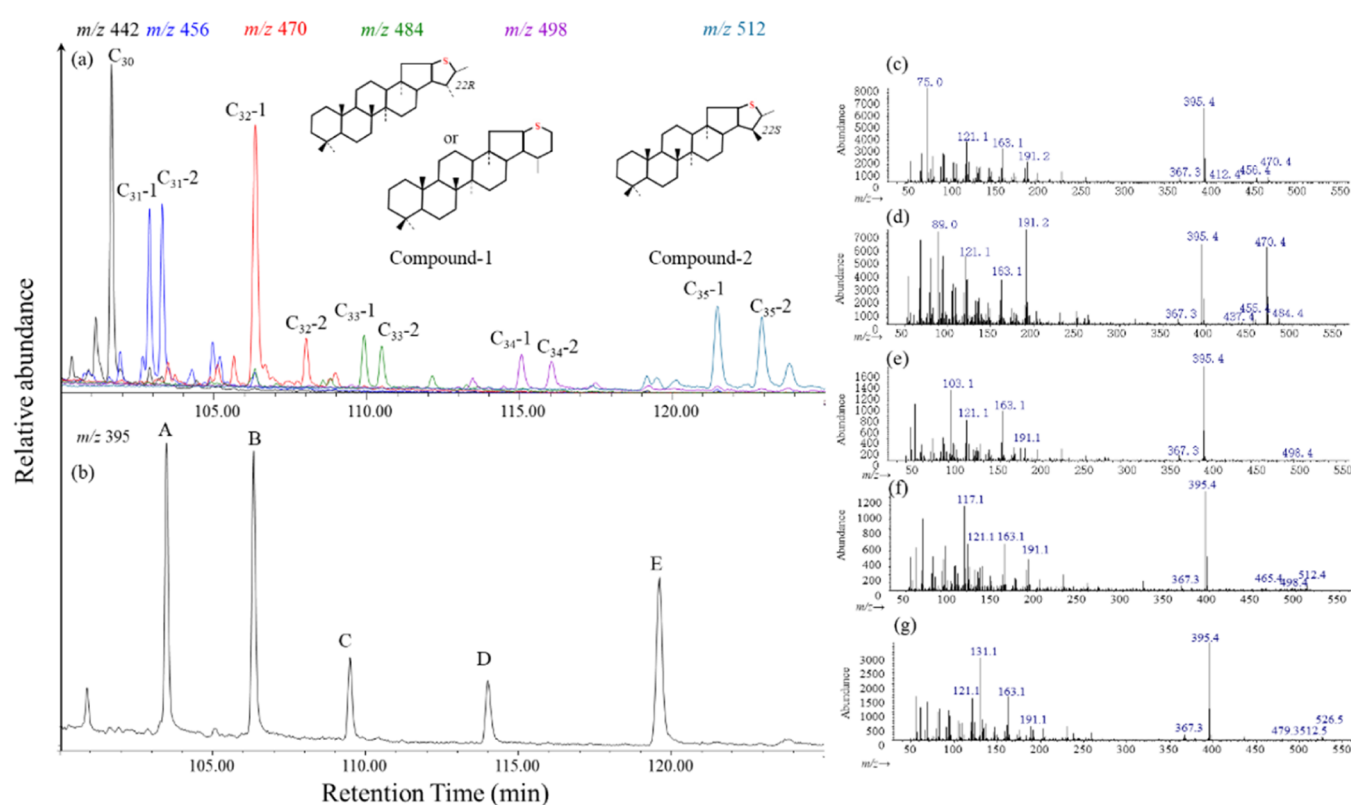


Figure 5. Representative mass chromatograms showing the distribution of sulfide hopanes from C_{30} to C_{35} in sulfide fractions of BZ35-2 samples, C_x -1: compound-1 of C_x sulfide hopane and C_x -2: compound-2 of C_x sulfide hopane (a); mass chromatograms showing the distribution of compound A to compound E in sulfide fractions of BZ35-2 samples (b); mass spectra of compound A to compound E (c–g).

C_{30} precursors such as diplopterol (hopan-22-ol) or diploptene (hop-22(29)-ene), found in nearly all hopanoid-producing bacteria.^{28,33,90,91} Hopanoids were once thought as biomarkers of aerobic bacteria (Talbot and Farrimond²³ and references therein) even though their biosynthesis does not require O_2 . However, the biosynthesis of the hopanoids has now been reported in a series of anaerobic bacteria.^{42,92} Therefore, abundant hopanes found in aliphatic, aromatic, and sulfidic fractions might imply that their diagenetic environment may be closely associated with active microbial activities.

The $\delta^{13}C$ values of lipid biomarkers are strongly determined by the isotopic composition of their biosynthetic precursors and their biosynthesized pathway.^{33,93} Generally, it is assumed that the carbon isotopic fractionation in a diagenetic process is limited.⁵¹ The ^{13}C isotope value differences between C_{31} hopane and C_{29} – C_{30} hopanes ranging from -3.3 to 2.2% (Figure 6) might illustrate different bacterial sources of these compounds as these differences are larger than those from a single bacteriohopanepolyol precursor with a specific ^{13}C isotope value. There are abundant bacteriohopanepolyols with different chemical functional groups on the side chain, which are thought as potential indicators of various organisms. The differences in diagenetic transformation pathways result in the formation of homohopanes with different distributions.¹⁴ The commonly occurring bacteriohopane-32,33,34,35-tetrol is ubiquitous in sediments and soils, and it is found in a diverse suite of microorganisms (Rohmer¹⁹ and references therein). Therefore, it is not diagnostic for a specific bacterial community. It is ascertained that a wide range of ^{13}C isotope values between C_{31} hopane and C_{29} – C_{30} hopanes represent the different precursors of these compounds (Figure 6).

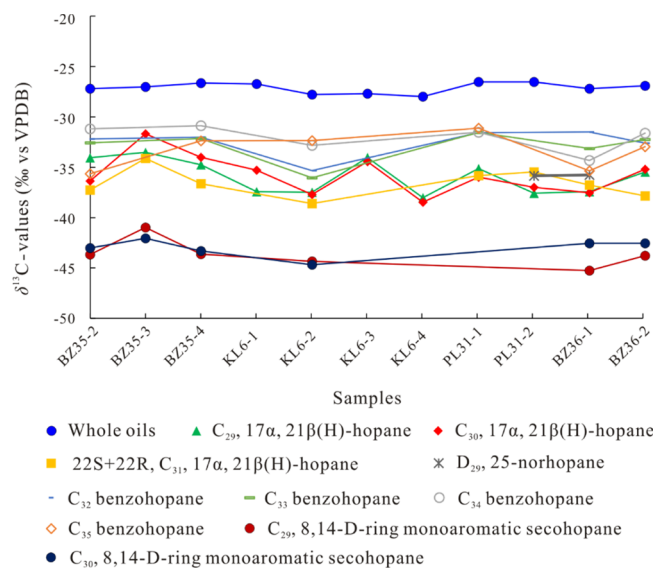


Figure 6. Distribution of the carbon isotopes of whole oils, hopanes, benzohopanes, and aromatic 8,14-secohopanes.

Diplopterol (hopan-22-ol) or diploptene (hop-22(29)-ene) was believed to be one of the precursors of a lower hopanoid ($<C_{30}$).⁹⁰ The ^{13}C isotope values of C_{29} and C_{30} hopanes may represent a similar origin, with the differences of ^{13}C isotope values from -0.8 to 0.7% in most samples. Oil samples in wells BZ35-2, BZ35-3, and BZ35-4 have greater differences of the $\delta^{13}C$ values between C_{29} and C_{30} hopanes. Sakata et al.⁹⁴ measured the carbon isotopic values of the cultured cyanobacterium *Synechocystis* UTEX 2470, finding that the

^{13}C isotope values of C_{30} diplopterol and diploptene are enriched (0.09–2.3‰) relative to that of bacteriohopanetrol. It is likely that the relative contribution of bacteriohopanetrol and diplopterol (diploptene) leads to the fluctuation of ^{13}C isotope values between C_{29} and C_{30} hopanes in BZ35-2, BZ35-3, and BZ35-4 samples.

The 25-norhopane series, also named as 10-demethylhopanes, generally occur in biodegraded oils and are widely used to assess the biodegradation level.⁶ The biological source of 25-norhopanes remains obscure. No microorganism is known to have the ability to biosynthesize 25-norhopanes.^{15,73,96} The 25-norhopanes and their counterparts mostly in biodegraded reservoir oils indicate that 25-norhopanes originate directly from microbial degradation of hopanes.^{74,81,96} However, the degradation compounds of hopanes are not only 25-norhopanes.^{97,98} Intriguingly, no 25-norhopanes were observed in enrichment cultures or pure aerobic bacteria.⁹⁸ Methanogens living in fresh or brackish water were thought to be probably responsible for the degradation of hopanes to 25-norhopanes.^{95,99} The 25-norhopanes were observed only in biodegraded oils (PL31-2 and BZ36-1 samples). However, no 25-norhopanes were detected in any of the m/z 177 of the non-biodegraded oils, perhaps indicating a major origin of microbial degradation. The ^{13}C isotope values of C_{29} 25-norhopanes in PL31-2 and BZ36-1 are both higher than those of C_{30} hopanes, with deviations of 1.1 and 1.8‰, respectively (Figure 6). It is likely that the 25-norhopanes are the microbially mediated demethylation products of hopanoids, which lead to the enrichment of the carbon isotopes. Dominance of the C_{29} 25-norhopane suggests incipient biodegradation of hopane compounds.⁷⁴

Benzohopanes were reported to be the transformation products of bacteriohopanetrol.^{84,85} Benzohopanes and C_{31} hopane are both products of a transformation of bacteriohopanepolyol. Nevertheless, varying differences of ^{13}C isotope values (1.5–6.5‰) between these two types of hopanes might represent (1) the mediation of bacterial activities and (2) the enrichment of the ^{13}C isotope in the aromatization of these compounds. The $\delta^{13}\text{C}$ values of benzohopanes are even isotopically heavier than those of the C_{29} – C_{30} hopanes and the C_{29} 25-norhopane. Liao et al.³³ found ^{13}C enrichments of 2–4‰ in aromatized hopanes via diagenetic alteration and successive dehydrogenation from the D-ring to the A-ring. In our study, the ^{13}C isotope enrichment of some compounds is even greater than 5‰, suggesting that the benzohopanes are not only produced by cyclization of bacteriohopanepolyol, but they are also derived from heterotrophic or cyanobacteria in the paleo-ecosystem during deposition of its source rocks.²⁸ However, the cyanobacteria-derived biomarkers, such as the 2-methylhopanes²¹ and 7-,8-monomethyl heptadecanes,¹⁰⁰ are absent in the samples, probably suggesting that cyanobacteria might have a minor contribution to the benzohopanes, perhaps indirectly indicating that the heterotrophic bacteria are responsible for the origin of the benzohopanes.

The D-ring-8,14-sec-hopanes in sedimentary organic matter have been reported for decades,^{83,101–103} which might be formed in the subsurface from bacterial C_{35} hopanoid precursors.⁸⁴ The heavily depleted ^{13}C isotopes (–45.6–41.2‰) of C_{29} and C_{30} SHs relative to hopanes and benzohopanes (Figure 6) tentatively suggest the mixed sources of SHs from methanotrophs and other bacteria.^{103,104} Methanotrophs utilizing methane as the carbon source that

produced depleted $\delta^{13}\text{C}$ values have been reported in previous research studies.^{103,104} Freeman et al.⁵⁰ interpreted the heavily depleted carbon isotopes of hopanes (up to –65‰) as sources of various amounts of methylotrophs, that is, recycling of carbon through methanotrophs. Volkman et al.¹⁸ reported a very light carbon isotope of C_{29} hop-17(21)-ene (avg. –50.8‰) from an oil shale in the Huadian Basin, which was attributed to the contribution of methanotrophs. However, the ^{13}C isotopes of C_{29} SH (–45.6–41.2‰) and C_{30} SH (–45–42.2‰) are less depleted than hopanes with a sole methanotroph contribution, indicating the mixed input of some unknown bacterial sources and possibly the methanotrophs.^{18,25} It is noteworthy that 3-methylhopanes are not the samples found in this study, which are similar to the methylhopanoids in the Huadian Basin, probably indicating that the original methanotrophs do not have the ability to biosynthesize 3 β -methylhopanoids.¹⁸

5.2. Diagenetic Factors Determining the Hopanoid Distributions. The formation of hopanoids is controlled by multiple factors, such as bacterial degradation, diagenesis, thermal maturity, and biological precursors.^{29,106}

Abundant sulfide hopanes are detected in HSOs (Figure 5). Due to the limited sulfur in microorganisms, the sulfur-containing compounds are thought to be formed by incorporated sulfur in functionalized hydrocarbons and carbohydrates under favorable water conditions.^{107–109} There are two mechanisms of incorporating an inorganic sulfur element into organic matter. First, the molecules are synthesized via intermolecular sulfur bridges to form micro-molecules, such as kerogen, asphaltenes, and resins.¹¹⁰ Second, some functional groups in the molecules are added or substituted by inorganic reducing sulfur to form organic sulfur compounds with thiane, thiolane, and thiophene rings.¹¹¹ Cyr et al.³⁶ suggested that the predominant peaks of sulfide hopanes at m/z 442, 456, 470, 484, 498, and 512 were derived from hopanoids bearing either a thiacyclopentane ring between C_{16} and C_{21} or a thiacyclopentane ring between C_{20} and C_{21} . The structures of sulfur hopanes are characterized by the linkage of the C_{30} and the C_{16} or the C_{20} carbon position via a sulfur bond. In low-maturity samples, the sulfurization is closely associated with the bacterial sulfate reduction (BSR) and other reactions in early diagenesis.¹¹²

Numerous studies reported the isomerization and formation processes of hopanes.^{14,18,25,91,113–117} A common sequence of reactions is conversion of bacteriohopanepolyol to hopene to hopane.²⁵ Nevertheless, rearranged hopanes and novel series of hopanes might have experienced the more complex transformation process under special conditions.^{91,118} In this study, there is a dominance of normal hopane distributions. However, the 8,14-sec-hopanes occurring in saturated hydrocarbons might suggest the special generation mechanism for sec-hopanes. The 8,14-sec-hopanes are considered to be presumably derived from both hopanes and moretanes via ring C opening during maturation or degradation because the C-8,14 bond has the lowest energy in the whole hopane skeleton.¹¹⁹ Wenger and Isaksen¹⁰³ reported that the 8,14-sec-hopanes were formed via a microbial alteration in the reservoir. Additionally, the compounds were directly formed during early diagenesis via transformation of bacteriohopanepolyol precursors,¹⁰⁶ with a high resistance to biodegradation.^{103,120} The 8,14-sec-hopanes were recognized in all samples, including biodegraded and non-biodegraded oils. Further, the C_{31} – C_{35} secomorethane/sec-hopane ratios have

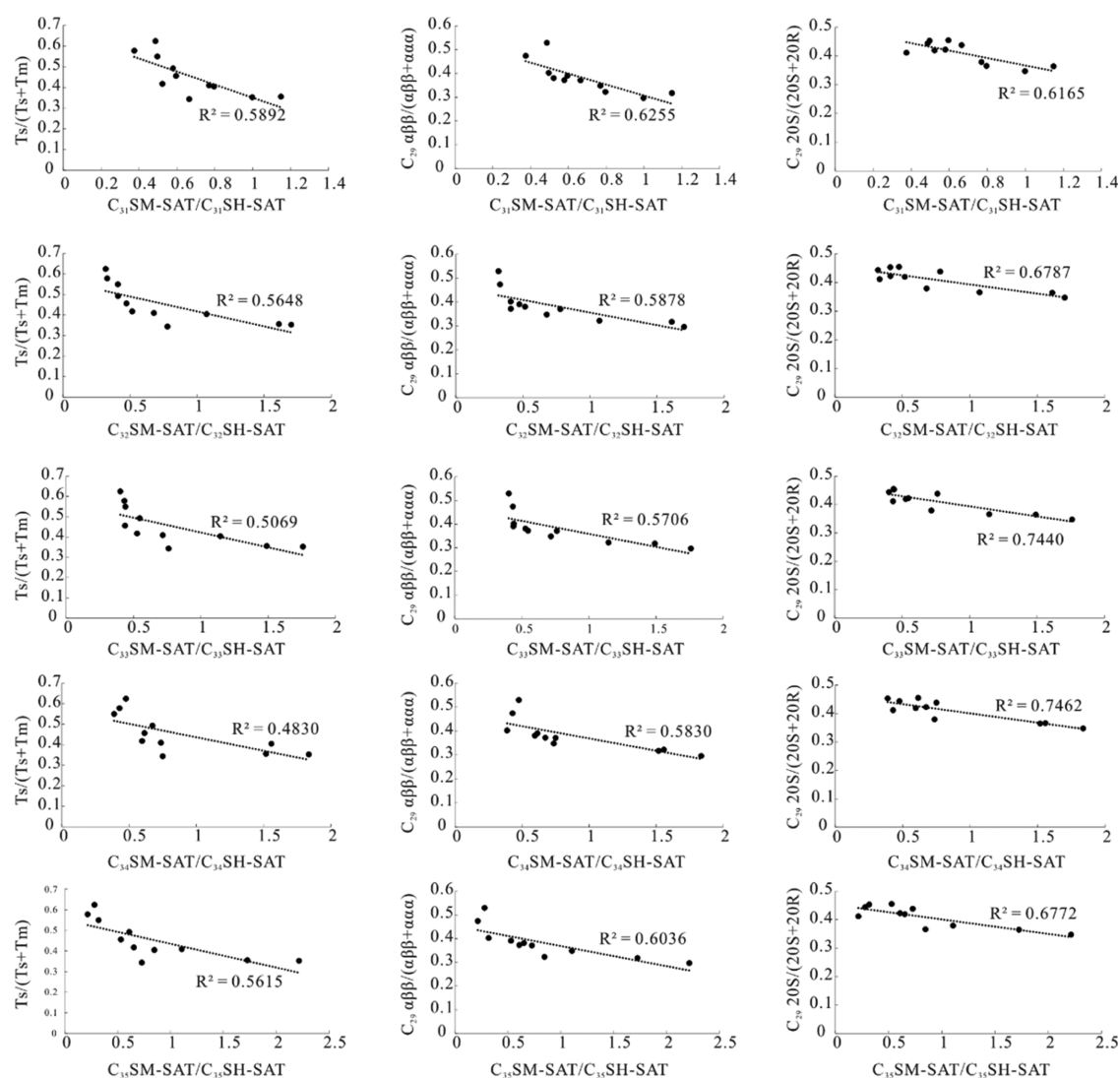


Figure 7. Correlations of C₃₁–C₃₅ SM-SAT/SH-SAT vs Ts/(Ts + Tm), C₂₉ αβ/(αα + αβ), and C₂₉ 20S/(20S + 20R). C₃₁–C₃₅ SM-SAT: C₃₁–C₃₅ 8,14-secomoretanes in saturates. C₃₁–C₃₅ SH-SAT: C₃₁–C₃₅ 8,14-secohopenes in saturates.

negative correlations with thermal maturity parameters including Ts/(Ts + Tm), C₂₉ ββ/(αα + ββ), and C₂₉ 20S/(20S + 20R) (Figure 7), suggesting the influence of maturity.

The benzohopanes are in high abundance in source rocks and crude oils under a reducing environment.^{84,85} Besides cyclization at C₂₀, Schaeffer et al.¹²¹ reported a novel series of benzohopanes cyclized at the C₁₆ position. These novel benzohopanes extending from C₃₁ up to C₃₅ and their mass spectra show a weak fragment at *m/z* 191 and a more pronounced *m/z* 197 + (*n* × 14). These benzohopanes always co-occur with the regular benzohopanes in immature sediments, particularly under carbonate-rich and evaporitic conditions.^{33,121} However, benzohopanes in monoaromatic hydrocarbons have the base peak of *m/z* 191 with the corresponding molecular ions and the lack of C₃₁ compounds, suggesting that the benzohopanes are all regular counterparts. Regular benzohopanes are usually formed and enriched at the early diagenesis stage,^{101,102} suggesting that the formation of compounds is not only influenced by thermal maturity but probably formed by the catalysis of the mineral matrix⁸⁴ or active microbial activities.³³ The enrichment of the ¹³C isotope

values might partly indicate that the diagenesis of these compounds is mediated by microbial reactions.

A series of SHs are particularly abundant in a confined environment⁸⁴ and are formed from bacteriohopanepolyols. However, the formation of aromatic 8,14-secohopenoids is likely to occur in the later stage of maturation.¹⁰² Compared to benzohopanes, the formation of aromatic 8,14-secohopenes needs aromatization of the D ring and opening the C ring of original hopanoid molecules, which requires more energy in the later stage of oil window.^{101,102} With an increase in maturity, SHs would convert to DSHs via demethylation and to IDs by further aromatization.⁸⁷ Both benzohopanes and aromatic 8,14-secohopenes are more resistant to biodegradation.¹⁰¹ In view of the differences of formation and degradation between benzohopanes and aromatic 8,14-secohopenes, He and Lu¹⁰² proposed a parameter, that is, D-ring monoaromatized 8,14-secohopenoids/(D-ring monoaromatized 8,14-secohopenoids + benzohopanes) (shortly, MAH) to evaluate the maturity of sediments and oils. It is suggested that the value of MAH at 0.3 may be the threshold of oil generation. However, in the low-mature oil samples, the MAHs are in the range of 0.83–0.94, which corresponds to the mature to high-

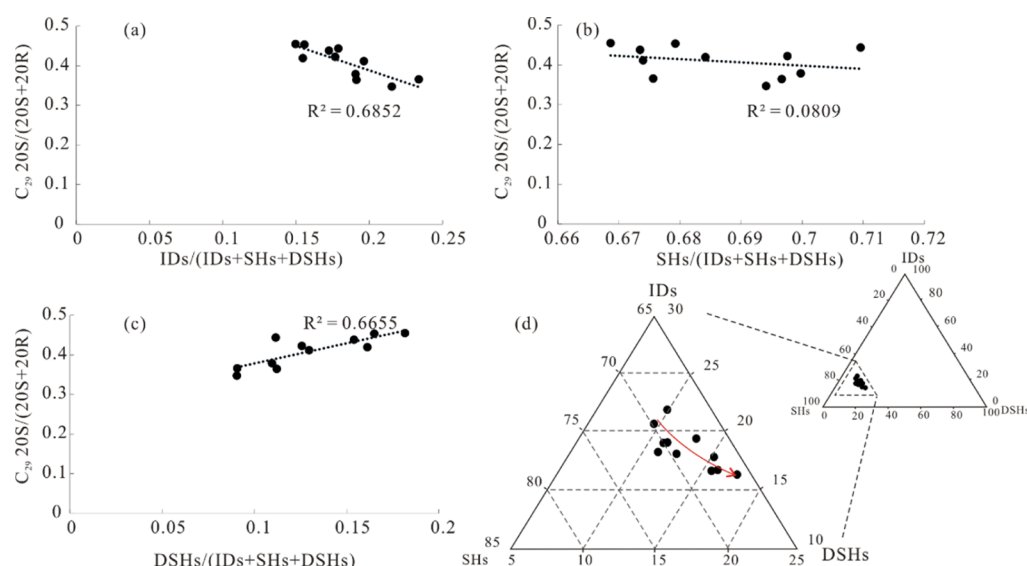


Figure 8. Correlations of $C_{29} 20S/(20S + 20R)$ vs $IDs/(IDs + SHs + DSHs)$ (a), $SHs/(IDs + SHs + DSHs)$ (b), $DSHs/(IDs + SHs + DSHs)$ (c), and ternary diagram of IDs , SHs , and $DSHs$ (d), showing the relative enrichment of IDs in low-mature samples.

mature stage based on He and Lu.¹⁰² Therefore, the bacterial reaction of the compounds cannot be excluded. In addition, the relative distributions of SHs , $DSHs$ and IDs are different from that reported in Killops et al.⁸⁷ In comparison with higher-maturity oils, there are abundant IDs in the relatively low-mature ones (Figure 8). This likely suggests that the formation of IDs might be mediated by microorganism reactions.^{87,102} The prominent peaks of SHs , $DSHs$ and IDs are $C_{30}SH$, $C_{29}DSH$, and $C_{29}ID$, respectively (Figure 4). The distribution of these compounds is similar to that of hopanes. However, a trace of aromatic 8,14-secohopanes ($>C_{30}$) is tentatively assigned in oils together with the depleted $\delta^{13}C$ values of C_{29} and C_{30} SHs , indicating the complexity of the diagenesis process and bacterial input. If methane is the main carbon source for the bacterial community, the ^{13}C isotope values of hopanoids and organic matter would be in the range of -70 to -90% .¹²² These results might imply that hopanoids from methanotrophs may have a contribution of SHs via microbial reactions during early diagenesis. Then, the C_{30} precursor hopanoids are altered and formed the dominance of $C_{30}SH$ in hopanes.

5.3. Depositional Environment. The high sulfur content and the identified abundant sulfur-containing hopanoids in HSOs together with isotopically depleted C_{29} and C_{30} SHs indicate that the depositional environment has an active carbon and sulfur cycle. The HSOs are mostly formed by the BSR and thermochemical sulfate reduction (TSR). TSR is proposed as the thermally driven reaction between a hydrocarbon and hydrogen sulfide.¹²³ The oils originated from the TSR reaction generally have high maturity and are rich in aromatic sulfur compounds including dibenzothiophenes and their derivatives.¹²⁴ This is obviously contradicted by our samples, which have abundant sulfide compounds formed by BSR in the low-maturity stage. A specific aryl isoprenoid, that is, isorenieratane, has been identified in the HSOs, implying the presence of sulfur bacteria in the photic zone euxinia of the water column.¹²⁵ Isorenieratane is exclusively biosynthesized by anaerobic, photosynthetic, green sulfur bacteria (*Chlorobactene*) living in organic-rich photic zones, where hydrogen sulfide is abundant for the

generation of green sulfur bacteria.^{126,127} The identification of isorenieratane and abundance of sulfur-containing compounds suggest the presence of sulfur-reducing bacteria and green sulfur bacteria (*Chlorobactene*) (Figure 9). In addition, abundant aryl isoprenoids are found in HSOs (Figure 9). Aryl isoprenoids are thought to be formed from the C–C bond cleavage of isorenieratene and β -isorenieratene with the isorenieratene from green sulfur bacteria, whereas β -isorenieratene was likely to be formed from the ubiquitous β -carotene.^{126,128} Schwark and Frimmel¹²⁹ proposed the aryl isoprenoid ratio (AIR, $\Sigma C_{13-17}/\Sigma C_{18-22}$) to assess the persistence of photic zone euxinia. An AIR > 3.0 suggests a short-term episodic photic zone euxinic condition, while an AIR < 3.0 indicates persistent photic zone euxinia. Xu et al.¹²⁷ used isorenieratane/ C_{18} aryl isoprenoid (Iso/ C_{18}) to evaluate the environment perturbations of Es_4U member source rocks in the Dongying Depression. The source rocks in a marine setting usually have Iso/ $C_{18} < 3.0$, whereas marine-related events (such as summer hurricanes or the East Asian monsoon) may lead to an abrupt increase in the Iso/ C_{18} ratio. The AIR ranges from 0 to 0.8 (lower than 3.0) for HSOs, which probably implies a photic zone euxinic environment. This is consistent with the presence of isorenieratane in the photic zone euxinia of the water column. This hypothesis needs to be further investigated. Furthermore, the good correlation ($R^2 = 0.86$) between the sulfur content and the gammacerane index ($Ga/C_{30}H$) indicates that the reducing depositional environment partly results in the incorporation of an inorganic sulfur element into organic matters (Figure 10).

The sulfur-reducing bacteria can reduce the elemental sulfur to hydrogen sulfide, and the green sulfur bacteria would utilize it as an electron donor to oxidize it back to elemental sulfur.¹³⁰ The living conditions of green sulfur bacteria are an anaerobic and a transparent water column.¹³¹ Isorenieratane is the diagenetic product of isorenieratene, which originates from brown pigmented *Chlorobiaceae*.¹³¹ Brown-pigmented green sulfur bacteria can adapt to the low irradiance and blooms of the bacteria occur at a 2–80 m depth of the water column (most water depths are less than 17m).⁷³ However, there is a low abundance of isorenieratane in HSOs. The three possible

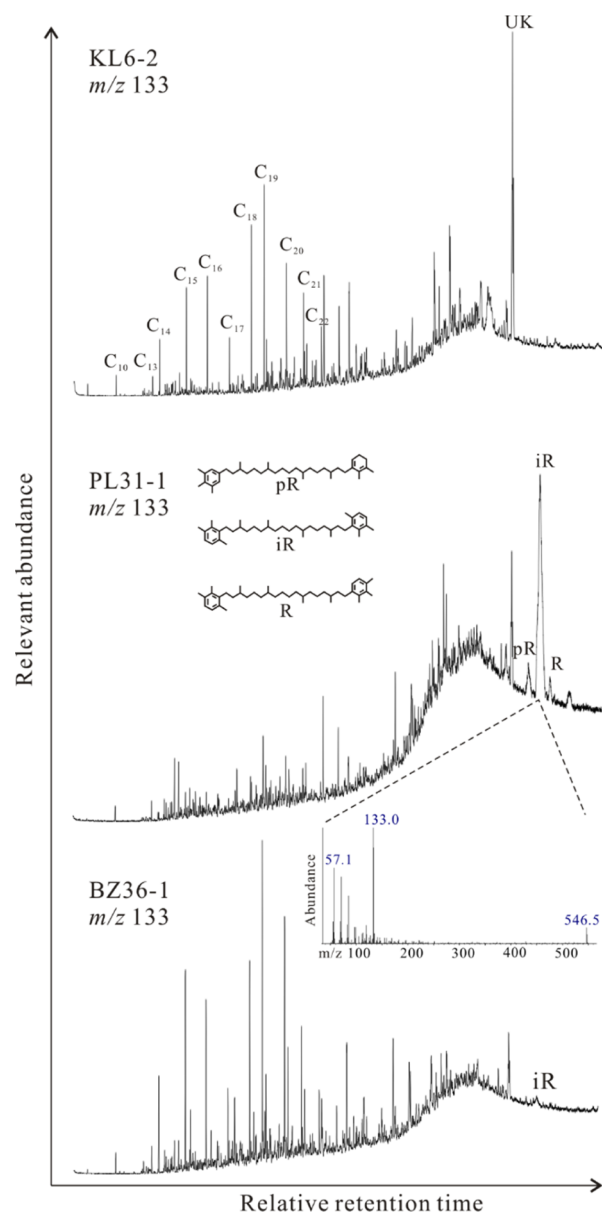


Figure 9. Representative m/z 133 mass chromatograms, showing the distributions of monoaryl isoprenoids (C_{10} – C_{22}), paleorenieratane, isorenieratane, and renieratane (R) which are typical biomarkers for green sulfur bacteria (*Chlorobiaceae*). UK: unknown compound.

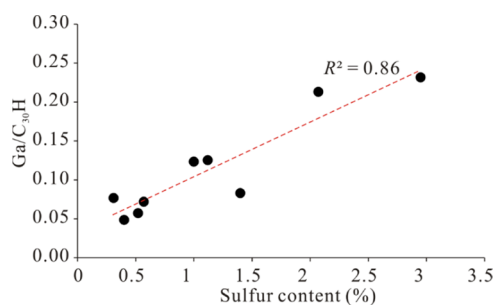


Figure 10. Cross-plot of the sulfur content vs $Ga/C_{30}H$, indicating the positive influence of a reducing environment on the sulfur content.

reasons for the presence of low concentrated isorenieratane are tentatively explained. First, the effects of migration might result

in the loss of some micromolecular biomarkers including isorenieratane. Second, isorenieratane might be degraded when the corresponding source rocks experience the diagenesis process. Third, the high primary productivity at the upper chemocline might influence the wavelength distribution of light. Thus, the relative distribution of green sulfur bacteria might be influenced. Clearly, the occurrence of brown-pigmented green sulfur bacteria means a reducing water column. Together with the HSOs and abundance of organic sulfur compounds, indicating the good preservation of organic matters. Moreover, the ratio of pristane (Pr) to phytane (Ph) is quite low (0.85–1.17) among the HSOs in a reducing sedimentary environment.¹³² It is worth nothing that isorenieratane is also identified in the LSOs and the Pr/Ph ratio is in the range of 1.1–1.3, implying that the paleo-water column is also under a reducing condition. However, no sulfide hopanes in biomarkers are identified in LSOs. The BSR reaction might probably also occur at the corresponding reducing water column. Nevertheless, hydrogen sulfide may be preferentially consumed by abundant ferric ions in sediments to produce early diagenetic pyrite, resulting in a low sulfur content of organic matter.¹³³

The isotopically depleted ^{13}C isotope values of aromatic secohpanes in all samples indicate the occurrence of a methane cycle during the depositional period, suggesting that methane is used by special microbes as the main carbon source and energy.^{122,134} Volkman et al.¹⁸ reported that C_{29} neohop-13(18)-ene and C_{30} hop-17(21)-ene had very negative ^{13}C isotope values (–50.8 and –43.7‰, respectively) in the Huadian shale under reducing depositional conditions. This is attributed to the depletion of ^{13}C isotope values to the contribution of methanotrophs even though 3β -methylhopanes and other biomarkers indicative for the methanotrophic bacterial activity were not detected. Similarly, Xie et al.¹³⁵ found that stable carbon isotopes of C_{29} and C_{30} 17α , $21\beta(H)$ -hopanes had a range of –45.6 to –61.5‰ in the Lucaogou formation oil shale from the Junggar Basin, considerably lighter than the n -alkanes (ca. –34‰), suggesting the contribution of methanotrophs. The ^{13}C isotope values of C_{29} secohpanes (–45.6–41.2‰) and C_{30} secohpanes (–45–42.3‰) are very similar to the isotopes' depleted values of hopanes in sediments in which methanotrophs use methane as a carbon source.¹⁰⁵ Methane is produced by methanogenic archaea, which are anaerobic bacteria,¹³⁶ while free-living aerobic methanotrophs are often concentrated at the microaerophilic interface between oxic and anoxic zones (chemocline).¹³⁷ Probably, the water column stratification might lead to the coexistence of anaerobic bacteria (methanogenic and sulfur-reducing bacteria) and aerobic bacteria (methanotrophs). The gammacerane index of HSOs is in the range of 0.12–0.23, suggesting the stratification of a water column and possibly the brackish water conditions.

5.4. Geological Implications of Hopanoids. Besides the regular parameters of hopanes, secohpanes, aromatic 8,14-secohpanes, and sulfide hopanes have also been altered during the diagenetic process.^{84,102}

In our samples, the MAH parameters also have good correlations ($R^2 = 0.67$ and 0.62 , respectively) with maturity parameters of saturated hydrocarbons including $Ts/(Ts + Tm)$ and $C_{29} \beta\beta/(\alpha\alpha + \beta\beta)$, although the MAH parameter is in a narrow range of 0.83–0.94 (Figure 11). The ratio of moretane/hopane is the typical parameter to indicate the maturity of samples because of the transformation of $\beta\alpha$ -

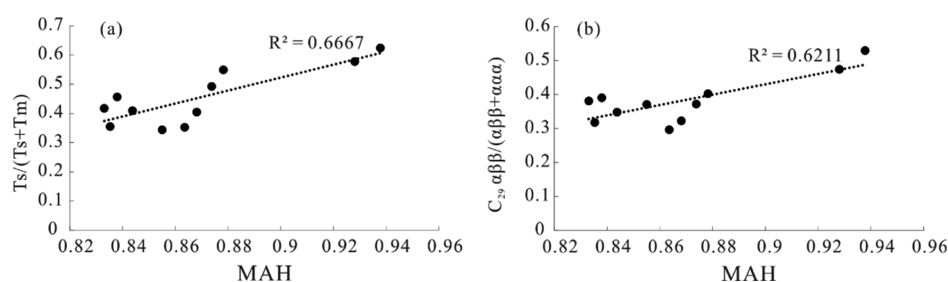


Figure 11. Correlations of MAH vs $Ts/(Ts + Tm)$ (a) and $C_{29} \alpha\beta\beta/(\alpha\alpha\alpha + \alpha\beta\beta)$ (b), showing good correlations between two parameters.

moretane to $\alpha\beta$ -hopane under thermal effects.¹⁰⁶ However, only some studies reported the application of 8,14-secohopanes in saturated hydrocarbons.^{82,119} The ratios of C_{31} – C_{35} secoretanes/secophanes all have good negative correlations with maturity parameters including $Ts/(Ts + Tm)$, $C_{29} \beta\beta/(\alpha\alpha + \beta\beta)$, and $C_{29} 20S/(20S + 20R)$ (Figure 7), suggesting that the distributions of secoretanes and secophanes are also influenced by mature effects in the studied samples in this region. The complexity of these compounds might indicate that the distributions are controlled by comprehensive factors. At least, the thermal effect might influence the evolution of 8,14-secohopanes in saturated hydrocarbons.

Aromatic 8,14-secohopanes were thought to be formed via thermal degradation of hopanes.^{87,101} The sequence of formation is SHs to DSHs to IDs.⁸⁷ In contrast, IDs are more enriched in lower-maturity samples, while DSHs are enriched in higher-maturity samples (Figure 8). This suggests that other reactions might have participated in the formation of aromatic 8,14-secohopanes except thermal maturity.

Besides the thermal effects on the distribution mode of hopanoids, the depositional environments also greatly impact the formation of hopanoids,¹⁰⁶ especially the sulfide hopanes. The decreasing relative abundance from C_{30} hopane to C_{35} hopane is normal in oil samples. However, the relative abundance of C_{35} hopane in HSOs is higher than in LSOs, even slightly higher than the C_{34} hopane in HSOs, suggesting a reducing depositional environment. However, the relative abundance of sulfide hopane is completely different from the distribution of hopanes (Figure 12). The preferential incorporation of inorganic sulfur into bacteriohopanepolyol derivatives leading to preservation of the intact carbon skeleton^{14,37} might result in the dominance of C_{35} sulfide hopane (Figure 12). The ratio of C_{35}/C_{34} sulfide hopane is in the range of 2.4–2.9, obviously higher than C_{35}/C_{34} hopane (0.82–1.00), suggesting that the incorporation of sulfur might protect the original carbon skeleton of bacteriohopanepolyol. The co-elution of compound B and C_{32} SH-1 results in the high abundance of C_{32} sulfide hopane (Figure 5). The ratio of C_{35}/C_{34} sulfide hopane has a good correlation with $C_{35}S/C_{34}S$ (Figure 13a), indicating that the long side-chain sulfide hopanes also experience a similar diagenesis evolution of hopanes in the low-mature to mature stage. Similarly, $C_{35}SH-2/C_{35}SH-1$ with the range of 0.90–1.56 has good correlations ($R^2 = 0.99$ and 0.88, respectively) with $C_{35}S/C_{34}S$ and $Ga/C_{30}H$ (Figure 13b,c), indicating that this ratio could be influenced by the reducing depositional environment. The sulfide hopanes are formed under the sulfur-containing bottom water column. The redox water environment might influence the distributions of compounds. Because of the low-mature to mature stage of the samples, the $C_{35}SH-2/C_{35}SH-1$ ratio is

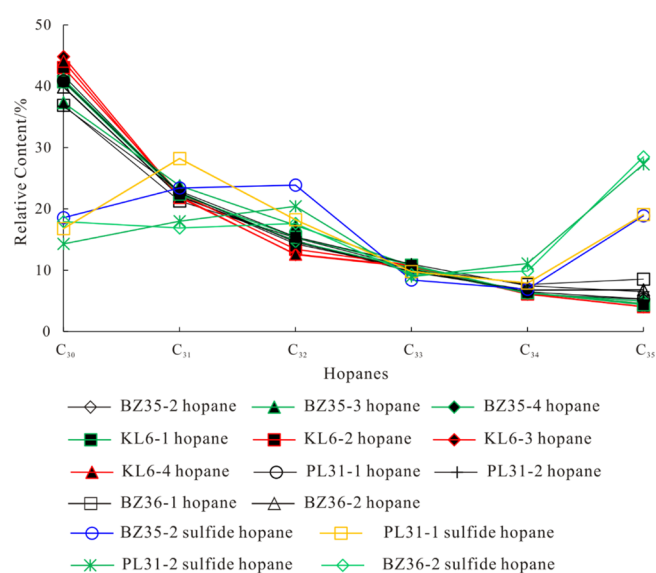


Figure 12. Relative distribution of C_{30} – C_{35} hopanes and sulfide hopanes.

mainly affected by the reducing depositional environment. In addition, the ratio of C_{35}/C_{34} benzohopane in the range of 0.30–0.94 also has a good correlation ($R^2 = 0.79$) with that of the $C_{35}S/C_{34}S$ hopane (Figure 13d). It is proposed that the long side-chain hopanoids in a reducing depositional environment may have experienced a similar evolution process, leading to good correlation between C_{35} and C_{34} hopanoids, although the hopanoids have different molecular structures.

Hopanoids are abundant in sediments and crude oils, and the compounds are very resistant to biodegradation.^{4,5,7} The parameters of hopanes can be used as an oil–source correlation, for maturity evaluation, and for depositional environment analyses, such as $C_{31}22R/C_{30}H$, $Ts/(Ts + Tm)$, $C_{29}Ts/(C_{29}H + C_{29}Ts)$, $C_{35}S/C_{34}S$, $C_{31}/C_{30}H$, $C_{29}/C_{30}H$, C_{31} – C_{35} 22S/(22S + 22R), and others.¹⁰⁶ In addition, the secophanes, aromatic 8,14-secohopanes, and sulfide hopanes are influenced by the thermal effects and depositional conditions of related source rocks. These compounds can provide more detailed information for the formation of high-sulfur crude oils. This knowledge provides a refined possibility of oil–oil and oil–source correlations. The BSR-derived high-sulfur crude oils were sourced from the source rocks under reducing paleo-conditions with water column stratification. This organic sedimentary environment is in favor of the growth of diverse microorganisms and preservation of the organic matter during the diagenetic process, resulting in the influence of some ratios, including C_{35}/C_{34} sulfide hopane, $C_{35}SH-2/C_{35}SH-1$, and C_{35}/C_{34} benzohopane. The parameter of C_{31} –

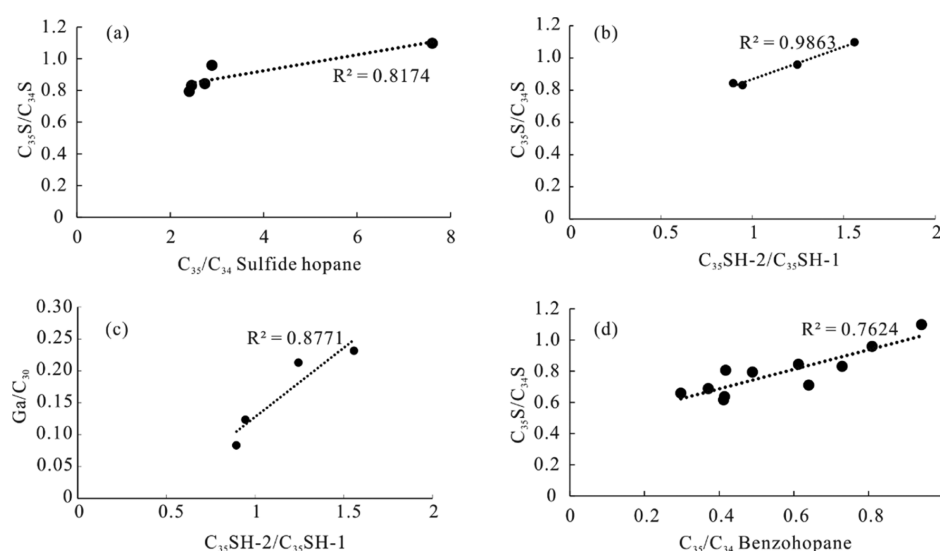


Figure 13. Cross-plots of C_{35}/C_{34} sulfide hopane vs $C_{35}S/C_{34}S$ (a); $C_{35}SH-2/C_{35}SH-1$ vs $C_{35}S/C_{34}S$ (b) and $Ga/C_{30}H$ (c); and C_{35}/C_{34} benzohopane $C_{35}S/C_{34}S$ (d). $C_{35}S/C_{34}S$: C_{35} 22S, 17 α (H), 21 β (H)-hopane/ C_{34} 22S, 17 α (H), 21 β (H)-hopane; $C_{35}SH-2/C_{35}SH-1$: compound-2/compound-1 of C_{35} sulfide hopane.

C_{35} secomoretanes/secohopanes is tentatively affected by the thermal effect in samples in the region.

6. CONCLUSIONS

Abundant hopanoids are identified in oils, including hopanes, secohopanes, 25-norhopanes, benzohopanes, aromatized secohopanes, and sulfide hopanes. The very depleted ^{13}C isotope values of C_{29} and C_{30} D-ring-monoaromatic-8,14-secohopanes indicate the presence of methane-oxidating bacteria (methanotrophs). The bacteria sulfate reduction results in the formation of HSOs. Perhaps, the stratification of the paleo-water column results in diverse microorganisms living in the oxic and anoxic environments.

The thermal effects and depositional environments influence the distributions of the parameters of secohopanes, aromatic 8,14-secohopanes, and sulfide hopanes. The ratios of C_{31} – C_{35} secomoretanes/secohopanes are mainly controlled by the maturity in the low-mature to mature stage in the studied samples in this region, showing the good negative correlations between C_{31} – C_{35} secomoretanes/secohopanes and maturity parameters of the saturates. Long side-chain hopanoids in a reducing depositional environment may experience a similar evolution process in the low-mature to mature stage, resulting in good correlations between C_{35}/C_{34} sulfide hopane, compound-2/compound-1 C_{35} sulfide hopane, C_{35}/C_{34} benzohopanes, and $C_{35}S/C_{34}S$ hopanes and the gammacerane index. The hopanoids in the saturates, aromatics, and sulfide fractions provide comprehensive information for high-sulfur crude oils, which is suitable for further oil–source rock correlations and exploration of HSOs in the Huanghekou Eastern Sag and the Laizhouwan Northeastern Sag.

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsomega.1c02762>.

Mass spectra of C_{29} 8,14-secohopanes, C_{30} 8,14-secohopanes, C_{32} benzohopane, C_{33} benzohopane, C_{34} benzohopane, C_{35} benzohopane, C_{29} D-ring monoar-

omatic 8,14-secohopanoide, C_{30} D-ring monoaromatic 8,14-secohopanoide, C_{28} D-ring monoaromatic 8,14-sec-28-nor-hopanoide, C_{29} D-ring monoaromatic 8,14-sec-28-nor-hopanoide, C_{29} indenyltrimane, C_{30} indenyltrimane, Compound-1 of C_{31} sulfide hopane, Compound-2 of C_{31} sulfide hopane, peak at Compound-1 of C_{32} sulfide hopane, and Compound-2 of C_{32} sulfide hopane (PDF)

■ AUTHOR INFORMATION

Corresponding Author

Dujie Hou – School of Energy Resources and Key Laboratory of Marine Reservoir Evolution and Hydrocarbon Accumulation Mechanism, Ministry of Education, China University of Geosciences (Beijing), Beijing 100083, China; orcid.org/0000-0003-2001-4082; Email: houdj313@163.com

Authors

Congkai Niu – School of Energy Resources and Key Laboratory of Marine Reservoir Evolution and Hydrocarbon Accumulation Mechanism, Ministry of Education, China University of Geosciences (Beijing), Beijing 100083, China
Xiong Cheng – School of Energy Resources and Key Laboratory of Marine Reservoir Evolution and Hydrocarbon Accumulation Mechanism, Ministry of Education, China University of Geosciences (Beijing), Beijing 100083, China; orcid.org/0000-0002-2991-1137

Xu Han – School of Energy Resources and Key Laboratory of Marine Reservoir Evolution and Hydrocarbon Accumulation Mechanism, Ministry of Education, China University of Geosciences (Beijing), Beijing 100083, China

Yan Li – School of Energy Resources and Key Laboratory of Marine Reservoir Evolution and Hydrocarbon Accumulation Mechanism, Ministry of Education, China University of Geosciences (Beijing), Beijing 100083, China

Yaxi Li – Consulting and Research Center, Ministry of Natural Resources, Beijing 100035, China

Complete contact information is available at:

<https://pubs.acs.org/doi/10.1021/acsomega.1c02762>

Notes

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