

**Composition and origin of lipid biomarkers  
in the surface sediments from the southern  
Challenger Deep, Mariana Trench**

HONGXIANG GUAN<sup>1\*</sup>, LINYING CHEN<sup>2</sup>, MIN  
LUO<sup>2</sup>, SHENGYI MAO<sup>2</sup>, JIASONG FANG<sup>2</sup>,  
DUOFU CHEN<sup>2</sup>

<sup>1</sup> *Key Laboratory of Gas Hydrate, Guangzhou  
Institute of Energy Conversion, Chinese  
Academy of Sciences, Guangzhou 510640,  
China*

<sup>2</sup> *Shanghai Engineering Research Center of  
Hadal Science and Technology, College of  
Marine Sciences, Shanghai Ocean  
University, Shanghai 201306, China*

The surface sediments were collected from the southern Mariana Trench in water depths ranging between ca. 4900 and 7068 m in order to understand the origin of lipid biomarkers and the source of sedimentary organic matter. Among these samples, unresolved complex mixture (UCM) and resistant component tricyclic terpanes were detected whereas C<sub>27</sub>-C<sub>29</sub> regular steranes and hopanes indicative of crude oils were almost absent. This suggest that the organic matter was partially derived from degraded oils. The relative high concentrations of even predominant *n*-alkanes and low Carbon Preference Indices (from 0.29 to 0.45) indicated that the *n*-alkanes were mainly synthesized by diverse microorganisms inhabiting a petroleum contaminated environment, since the *n*-alkanes are the first removed compound class in biodegradation. Also detected were even-carbon numbered *n*-fatty acids C<sub>12:0</sub>-C<sub>22:0</sub> with maximum at *n*-C<sub>16:0</sub> and *n*-alcohols C<sub>12:0</sub>-C<sub>22:0</sub> with the *n*-C<sub>18:0</sub> accounting for 90% of the total *n*-alcohols. Characteristics of these biomarker patterns were attributed to a large input of algae and bacteria. The longer fatty acids and *n*-alkanes were more depleted in <sup>13</sup>C as a result of carbon chain elongation during lipid synthesis. In samples BC06, BC08, BC11-1 and BC13-1, GDGTs were mainly derived from marine Thaumarchaeota and may point to a significant input of methanogenic archaea in sample BC07.