

# Identification and geochemical significance of a series of rare lanostanes in the alkaline lacustrine source rock from the Mahu Sag, Junggar Basin NW China

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The organic-rich source rocks of high maturity from Lower Permian Fengcheng Formation (P<sub>1</sub>f) were deposited under an alkaline lacustrine environment in the Mahu Sag, Junggar Basin, Northwest China<sup>[1]</sup>. Eight P<sub>1</sub>f source rock samples were subjected to solvent extraction and saturated hydrocarbon fractions were collected by column chromatography. The fractions were analyzed by gas chromatography-mass spectrometry: *n*-alkanes, triterpenoid hydrocarbons and steranes were detected. A series of rare lanostanes, including a C<sub>23</sub> homologue not reported in the previous literature were present in this basin. The absolute concentration of short chain lanostanes bears a strongly positive correlation to those of long chain lanostanes, indicating their possible common biological origin and genesis. A well-defined relationship is evident between the content of lanostanes and the distributions of methyl heptadecanes, 2-methyl hopanes: data points with high 7-+8-methyl heptadecanes/C<sub>max</sub> and 2-methyl hopanes/C<sub>30</sub>-hopanes exhibit high lanostane contents. The 7-+8-methyl heptadecanes and 2-methyl hopanes are considered to be special molecule biomarkers sourced from cyanobacteria. Therefore, it is concluded that cyanobacteria also may be the diagnostic biological precursors of lanostanes in P<sub>1</sub>f source rock. In addition, the occurrence of relatively abundant lanostane may indicate the strongly reducing sedimentary environment, which is characterized by high content of phytane, β-carotanes and gammacerane<sup>[2]</sup>. The relative invariance of the content of lanostanes with maturity suggest that these compounds are not maturity dependent. Preliminary genesis analysis shows that the long chain (C<sub>30</sub>-C<sub>32</sub>) lanostanes in sediments was formed through diagenetic reactions from biological lanosterol via a reduction pathway and then were transformed to short chain lanostanes via demethylation.

[1]Cao, Xia, Wang, Zhi, Tang, & Li (2020), *Earth-Science Reviews*, 202, 103091.

[2] Chen, Philp, Fu, & Sheng (1989), *Geochimica et Cosmochimica Acta* 53, 2775-2779.