TRACING THE DIAGENETIC ORIGIN OF NEOPROTEROZOIC 26-ALKYLSTERANES

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Timing, conditions and triggers surrounding the rise of complex life remain amongst the most highly debated topics in geobiology. In the absence of large body fossils, prior to emergence of the Ediacara biota, molecular biomarkers are our only means to reconstruct the composition of communities during the Neoproterozoic (Brocks et al., 2017; Hoshino et al., 2017; van Maldegem et al., 2019). Recently, novel steranes with a methylation at position C-26 were described to be present in several Neoproterozoic sedimentary rocks (Brocks et al., 2015; Zumberge et al., 2018). These molecules were suggested to derive from sponges, thereby possibly representing the oldest evidence for animals in the geological record (Zumberge et al., 2018). While most recently, one of these purported sponge markers, 26-methylstigmastane, has also been observed in modern Rhizaria (Nettersheim et al., 2019), biological precursors of the C₂₈ steroid Cryostane have not yet been detected in extant organisms.

We found elevated abundances of 26-alkylsteranes in two Neoproterozoic sedimentary sequences. Apart from cryostane, methylated at position C-26, we also detect a pseudohomologous series with C-26-alkylation extending up to C₇ in both investigated sample sets. Alongside these 26-alkylsteranes, the samples also contain abundant 3β-alkylsteranes. When quantifying the relative abundance of each alkylated molecule (C₁–C₇) for both 3β- and 26-alkylsteranes a significant correlation (R²=0.98) is observed between the different homologous series samples with abundances that rapidly decrease with each additional methyl addition. This suggests that these alkylsteranes were generated through similar mechanisms.

Thirty years after their discovery (Summons and Capon, 1988), the formation mechanisms of 3β-alkylsteranes still remain enigmatic. Once thought to possibly represent membrane lipids of unknown bacteria that modify eukaryotic steroids by adding sugars to the C-3 position (Dahl et al., 1992; 1995), they are usually attributed to bacterially mediated processes during diagenesis. Pyrolysis experiments with various sterols have now revealed that alkyl-steranes readily form through diagenetic processes without biology and that the alkylation position is indeed a function of hydroxyl functionalization. Our data suggest that neither 3β- nor 26-alkylsteranes derive from specific alkylated precursors, negating cryostane as a metazoan marker. More importantly, our findings suggest that extended alkylsteranes allow the reconstruction of past side-chain functionalization in ancient steroids, which may open a new realm of biological diagnosticity.

References


