

# EVOLUTIONARY LINEAGES OF ALKENONES RECORDED IN CRETACEOUS AND PALEOCENE SEDIMENTS FROM THE TRANSKEI BASIN (IODP SITE U1581)

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C<sub>37</sub>-C<sub>40</sub> alkenones derived from haptophyte algae [1] were identified in Campanian to Paleocene (~76-60 Ma) sediments recovered from the Transkei Basin (Hole U1581B), offshore South Africa, during IODP Expedition 392 [2]. These occurrences extend the temporal continuity of alkenones and expand their paleogeographic range to southern high latitudes (~58°S) during this time interval [3-7], while their distributions exhibit several transitional features (**Fig. 1**). Cretaceous samples contain C<sub>40:3</sub> and C<sub>40:2</sub> alken-3-ones, confirming disparities between alkenone distributions for this period [4-7] and extant haptophytes. C<sub>40</sub> alkenones have only once been reported in marine species from phylogenetic Group III [8] but are common constituents of species in phylogenetic Group II [9,10]. The high relative abundance of C<sub>40:2</sub> alken-3-one [4-6] and other aspects of alkenone compositions in pre-Eocene (>56 Ma) sediments appear typical of lacustrine rather than marine Quaternary settings [11]. Thus, Cretaceous through Paleocene marine sediments may reflect alkenone contributions from both Isochrysidaceae and Noelaerhabdaceae following their Early Cretaceous divergence [12], recording a broader suite of alkenone biosynthetic pathways than marine haptophytes now express. The occurrence of C<sub>37</sub> and C<sub>38</sub> alkatrienones in Paleocene samples revises the timing of their first appearance and refutes the hypothesis [13] that this biosynthetic innovation in haptophytes reflects a post-EEO response to ocean cooling. Similarities in alkenone profiles for Maastrichtian and Danian samples suggests conformity in their biosynthetic pathways across the Cretaceous/Paleogene (~66 Ma) boundary, attesting to survival and subsequent recovery of their source haptophytes after the extinction event.

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