

The impact of diagenesis and sequestration on long-chain diols and derived marine proxies

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The impact of diagenesis and sequestration on long-chain diols and derived marine proxies

De gevolgen van diagenese en sequestratie op langkettige diolen en bijbehorende mariene proxies

(met een samenvatting in het Nederlands)

Der Einfluss von Diagenese und Sequestration auf langkettige Diole und zugehörige marine Proxies

(mit einer Zusammenfassung in deutscher Sprache)

Proefschrift

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*To the Sea, to the Sea! The white gulls are crying,
the wind is blowing, and the white foam is flying.*

J.R.R. Tolkien

Für meine Großeltern.

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Summary

The current global warming and climate change is widely discussed in science, politics and media. To predict climate development in the future, we depend on climate models that are validated and tested on appropriate climate archives of Earth's history, e.g. marine sediments, by application of climate proxies. These climate proxies are preserved material standing in for direct measurements and provide information about past environmental and climate conditions. Their application helps in understanding past and present climate change and in turn, they give insight into potential future climate changes. A variety of inorganic and organic chemical proxies are used to reconstruct paleoenvironmental parameters that include sea surface temperature (SST), concentration of CO₂ in the atmosphere, ocean salinity or marine primary productivity. Organic proxies are based on preserved sedimentary lipids, termed biomarkers and serve as indicators for the past presence of specific groups of organism or biogeochemical processes as well as environmental parameters. For example, two frequently applied organic SST proxies are the U^K₃₇ index, based on the degree of unsaturation of long chain alkenones produced by haptophyte algae, and the TEX₈₆ index, based on the number of cyclopentane moieties in glycerol dialkyl glycerol tetraether lipids (GDGTs) biosynthesized by Thaumarchaeota.

Recently, long-chain alkyl diols (LCDs) have been developed to reconstruct SST, productivity and upwelling conditions as well as terrestrial input into marine environments. LCDs, consisting of long alkyl chains with carbon numbers commonly ranging from C₂₈ to C₃₂ and alcohol groups at the first carbon position and in a mid-chain position at carbon position 12 to 15, have been recognized in a wide variety of marine and lacustrine environments. While saturated and mono-unsaturated 1,14-diols have been shown to be produced by *Proboscia* diatoms in marine environments, 1,13- and 1,15-diols have been identified in cultures of freshwater/ brackish Eustigmatophyceae algae, e.g. *Nannochloropsis salina* or *Nannochloropsis oculata*, however their marine producers remain unknown. The 1,13- and 1,15-diols commonly are present as extractable-bound lipids in sediments but only occur in small abundances as freely extractable lipids in algal biomass. They have been shown to form the building blocks of the highly aliphatic and insoluble biomacromolecules termed algaenan, which are built up of mainly 1,15-diols linked by ether bonds and are part of the cell-wall. While several proxies, i.e. the Long-chain Diol Index (LDI), to reconstruct SST, and the Diol Index, to study productivity and upwelling conditions, have been defined and are increasingly used in paleo-reconstructions, many uncertainties remain. Marine organic proxies are impacted by a large number of chemical and physical changes from the time the proxy signal is generated by the producer, during the transport and settling of the signal through the water column and the subsequent burial in the sediment. An impact of these diagenetic changes, i.e. by oxic degradation or mode of occurrence, has been shown for a variety of organic proxies, but has not been studied yet in detail for LCDs and their derived proxies. Therefore, the focus of thesis was to assess potential biases that may arise when LCD proxies are impacted by diagenetic changes.

In order to constrain the impact of oxygen exposure on LCDs and the LDI, *Nannochloropsis oculata* biomass was incubated aerobically for 271 days and the concentrations of free-extractable, extractable-bound, residually ester-bound and residually-bound glycosidic ether- or amide-bound fatty acids and LCDs were determined. Both saturated and polyunsaturated fatty acids were significantly impacted, showing substantially decreasing concentrations with increasing oxic incubation time, irrespective of their mode of binding. In contrast, LCDs showed a large-scale increase in concentration from incubation day 125 until the end of incubation at day 271, in particular for residually ester-bound LCDs which showed a 30-fold

increase in concentration. This increase in concentration likely reflects a release of LCDs from the biopolymer algaenan upon oxic degradation. In contrast to this large-scale increase in LCD concentration, LDI values remained stable throughout the degradation experiment, although LDI values were dependent on their mode of occurrence. Overall these experiments indicated a likely negligible impact of oxygen exposure on the LDI.

To ascertain if prolonged oxygen exposure indeed led to a release of LCDs from the biopolymer algaenan, residues left after extraction and saponification and hydrolysis of the aerobically degraded biomass of *Nannochloropsis oculata*, were analyzed by step-wise pyrolysis. The pyrolyzate of the algaenan was characterized by a series of *n*-alkenes/ *n*-alkanes with chain lengths ranging from C₁₀ to C₃₃ with two maxima at C₁₅ to C₁₇ and C₂₆ to C₂₈. Mid-chain ketones were present with chain lengths of C₂₈ to C₃₄ with a maximum at C₃₁ and a mid-chain keto group at position 14. Additionally, a pronounced C₁₉ *n*-alkan-2-one was present. These results point to a structural composition of the algaenan built-up of C₃₂ 1,15-diol units, linked by ether bonds, which is in agreement with previously published studies. Analysis of residues after extraction at different time points throughout the oxic degradation experiment, however, suggests that the algaenan structure was not largely altered, indicating that oxygen exposure does not have a substantial impact on the structure of the algaenan.

Since LCDs are thought to be constituents of the algaenan and have been reported to be released after hydrolysis of algal extracts, the identification of potential macro-molecular sources of LCDs in *Nannochloropsis sp.* biomass was attempted. However, application of ultra-high pressure liquid chromatography combined with high resolution mass spectrometry (UHPLC/HRMS) and semi-preparative HPLC did not lead to the identification of potential macromolecular precursor lipids. Additionally, only freely extractable and not extractable-bound lipids could be identified, which is in contrast to previously published studies. Subsequent analysis of three different strains of *Nannochloropsis spp.* showed that LCDs mainly occur as freely extractable lipids and only minor increases in concentration were detected after application of acid hydrolysis on the biomass extracts. These results therefore call into question the common thought that LCDs occur mainly as extractable-bound lipids.

A potential bias in the application of organic biomarkers may arise, when residually-bound lipids in marine sediments are released during diagenesis and may therefore yield different distributions and abundances than freely occurring lipids. Therefore, the impact of the mode of occurrence on LCDs and structurally related mid-chain keto-ols (keto-ols) was analyzed in the S5 sapropel of the Eastern Mediterranean. Some LCDs were shown to occur partly bound to the organic sedimentary matrix; extractable-bound 1,14-diols showed a 1.5-fold increase in abundance when compared to their freely extractable counterparts. While freely extractable LCDs dominate in the sapropel, LCDs occur predominantly as residually ester-bound and residually glycosidic ether- or amide-bound lipids in sediments outside of the sapropel. However, irrespective of the mode of occurrence of the LCDs, LCD indices showed similar trends over sapropel depth. Keto-ols, which may be formed by post-depositional oxidation of LCDs, were only detected in the sapropel and not in the sediments below or above it, in agreement with the hypothesis that they reflect intermediate LCD oxidation products. Their relative abundance in the sapropel, quantified by the Diol Oxidation Index (DOXI), did not show significant changes.

Additionally, the impact of post-depositional oxidation on LCDs and keto-ols was studied in the f-turbidite of the Madeira Abyssal Plain. Comparison of lipids in the upper, oxidized part of the turbidite, which has been impacted by post-depositional oxidation, with the anoxic lower part of the turbidite, shows a higher preservation of the C₃₀ 1,15-diol and C₃₀ 1,15-keto-ol compared to the other LCDs and keto-ols. While the LDI showed increased values, the Diol Index and FC₃₂ 1,15-diol, a proxy for the riverine input into marine environments, showed

decreased values with increasing oxygen exposure. Therefore, all three indices were impacted by post-depositional oxidation, likely resulting from a preferential preservation of the C₃₀ 1,15- and C₃₂ 1,15-diol over 1,13- and 1,14-diols. However, compared to GDGTs and GDGT-based organic proxies (BIT and TEX₈₆ index), measured in the same turbidite, LCDs and keto-ols seem to be better preserved upon post-depositional oxidation. Changes in oxygen exposure time in sediments may therefore cause different changes in organic terrestrial and temperature proxies and care has to be taken when applying these different indices in severely oxidized sediments.

Overall, the results of this thesis represent a first step towards constraining the impact of diagenesis and sequestration on LCDs and LCD proxies. A variable impact of both oxic degradation and mode of occurrence could be shown for LCDs and LCD proxies, both in algae biomass as well as in marine sediments, which points to potential biases that have to be taken into account when applying these proxies in paleoenvironmental reconstructions.

Zusammenfassung

Der Klimawandel und die damit verbundene globale Erwärmung sind Themen, die aus aktuellem Anlass in der Forschung, Politik und den Medien diskutiert werden. Um die zukünftige Klimaentwicklung vorhersagen zu können, werden verschiedene Klimamodelle verwendet, welche anhand geeigneter Klimaarchive der Erdgeschichte durch Anwendung von Klima-Proxies getestet und validiert werden. Klima-Proxies sind dabei indirekte Anzeiger, durch welche Informationen über die Umwelt- und Klimabedingungen, unter denen z. B. marine Sedimente abgelagert wurden, ermittelt werden können. Ihre Anwendung hilft beim Verständnis vergangener und gegenwärtiger Klimaveränderungen und erlaubt einen Einblick in mögliche zukünftige Klimaveränderungen. Verschiedene anorganisch- und organisch-geochemischen Proxies werden verwendet, um Paläo-Umweltparameter zu rekonstruieren. Beispiele für Paläo-Umweltparameter sind unter anderem Meeresoberflächentemperaturen, die CO₂-Konzentrationen in der Atmosphäre, der Salzgehalt der Ozeane oder die Primärproduktivität des Meeres. In organisch-geochemischen Proxies werden dabei Biomarker angewendet, welche charakteristisch sind für bestimmte Gruppen von Organismen, biogeochemische Prozesse oder Paläo-Umweltparameter. Häufig angewendete organisch-geochemische Proxies sind unter anderem der U^K₃₇-Index, basierend auf von Haptophyten produzierten ungesättigten langkettigen Alkenonen, und der TEX₈₆-Index, welcher die von Thaumarchaeota biosynthetisierten Membranlipide (GDGTs) berücksichtigt. In den letzten Jahren werden vermehrt langkettige Alkyldiole (Diole) genutzt um Meeresoberflächentemperaturen, Primärproduktivität, Upwelling und die Analyse des Eintrages von terrigenem organischem Material in marine Ablagerungsmilieus zu rekonstruieren. Diese Diole, welche in einer Vielzahl von marinen und lakustrinen Ablagerungsmilieus identifiziert wurden, sind charakterisiert durch Alkylketten mit Längen von C₂₈ bis C₃₂ und Hydroxygruppen an der ersten Kohlenstoffposition sowie an der Kohlenstoffposition 12 bis 15. Gesättigte und einfach-ungesättigte 1,14-Diole werden von marinen *Proboscia*-Diatomeen produziert. 1,13- und 1,15-Diole wurden in Kulturen von Eustigmatophyceae-Algen, z.B. *Nannochloropsis salina* oder *Nannochloropsis oculata*, nachgewiesen, jedoch ist der marine Produzent unbekannt. Üblicherweise, liegen 1,13- und 1,15-Diole als extrahierbar-gebundene Lipide vor und kommen nur in geringen Mengen als frei-extrahierbare Lipide in der Algenbiomasse vor. Weiterhin, sind sie Bausteine des aliphatischen und unlöslichen Biomakromoleküls Algaenan, welches einen Teil der Zellwand darstellt und hauptsächlich aus durch Ether-Verbindungen vernetzte 1,15-Diole besteht. Mehrere Proxies, basierend auf Diolen, wurden bisher definiert, u.a. der Long-chain Diol Index (LDI) zur Rekonstruktion von Meeresoberflächentemperaturen, sowie der Diol Index zur Untersuchung der Primärproduktivität und Upwelling-Konditionen. Während diese zunehmend in Paläo-Rekonstruktionen verwendet werden, ist ihre Nutzung mit Unsicherheiten verbunden. Marine organisch-geochemische Proxies sind von einer Anzahl chemischer und physikalischer Veränderungen betroffen. Diese beeinflussen das Proxy-Signal sobald es vom marinen Produzenten erzeugt wird, während es durch die marine Wassersäule transportiert und schließlich in marinen Sedimenten abgelagert wird. Diese diagenetischen Veränderungen beinhalten unter anderem den Einfluss von oxidativer Degradation sowie die Sequestrierung von Lipiden in der Sedimentmatrix und der Algenbiomasse. Daher liegt der Fokus dieser Doktorarbeit auf der Untersuchung von verschiedenen diagenetischen Einflüssen auf Diole und der ihnen zugrunde liegenden Proxies, welche bisher noch nicht ausreichend von der Forschung betrachtet wurden.

Um den Einfluss der oxidativen Degradation auf Diole und den LDI zu analysieren, wurde Biomasse der Alge *Nannochloropsis oculata* über 271 Tage aerob inkubiert und die

Konzentration von frei-extrahierbaren, gebunden-extrahierbaren, residual Ester-gebundenen und residual glykosidisch Ether- oder Amid-gebundene Fettsäuren und Diolen bestimmt. Gesättigte und mehrfach ungesättigte Fettsäuren zeigen stark abnehmende Konzentrationen über die aerobe Inkubationszeit, unabhängig von der Bindungsart, in welcher sie vorlagen. Im Gegensatz dazu, steigt die Konzentration der Diole ab dem Inkubationstag 125 bis zum Ende der aeroben Inkubation stark an, welches insbesondere bei den residual Ester-gebundenen Diolen zu einem 30-fachen Konzentrationsanstieg führt. Dieser Konzentrationsanstieg spiegelt dabei wahrscheinlich eine Freisetzung von Diolen durch oxidative Degradation aus dem Biopolymer Algaenan wider. Trotz des starken Anstiegs der Diol-Konzentration, bleiben die errechneten LDI-Werte stabil, sind jedoch abhängig davon wie die zur Berechnung herangezogenen Diole in der Biomasse gebunden sind. Ein vernachlässigbarer Einfluss von oxidativer Degradation auf den LDI ist daher wahrscheinlich.

Um zu überprüfen, ob eine länger andauernde Sauerstoffexposition tatsächlich zu einer Freisetzung von Diolen aus dem Biopolymer Algaenan führt, wurden die Rückstände, welche nach der Extraktion, sowie basischen und sauren Hydrolyse der aerob degradierten Biomasse von *Nannochloropsis oculata* zurückblieben, durch schrittweise Pyrolyse analysiert. Dabei zeigt sich, dass die Pyrolysate des Algaenan durch eine Serie von *n*-Alkenen/ *n*-Alkanen mit Kettenlängen von C₁₀ bis C₃₃ und zwei Maxima bei C₁₅ bis C₁₇ und C₂₆ bis C₂₈ charakterisiert sind. Mittelkettige Ketone liegen mit Kettenlängen von C₂₈ bis C₃₄ und einem Maximum bei C₃₁ mit einer mittelkettigen Ketogruppe an Position 14 vor. Besonders auffällig ist ein C₁₉ *n*-Alkan-2-on. Diese Ergebnisse deuten darauf hin, dass das Algaenan durch C₃₂-1,15-Diol-Einheiten aufgebaut ist, wie in bereits veröffentlichten Studien diskutiert. Die Analyse von Rückständen zu verschiedenen Zeitpunkten der aeroben Inkubation legt nahe, dass die Algaenan-Struktur nicht durch die oxidative Degradation verändert wurde. Somit ist es wahrscheinlich, dass eine Sauerstoffexposition über die hier analysierten Zeiträume, keinen wesentlichen Einfluss auf die Struktur des Algaenans hat.

Da davon ausgegangen werden kann, dass Diole Strukturbestandteile des Algaenan sind und durch Hydrolyse von Algenextrakten freigesetzt werden, wurde versucht potentielle makromolekulare Quellen für Diole in der Biomasse von *Nannochloropsis* sp. zu identifizieren. Mittels Ultrahochdruck-Flüssigkeitschromatographie in Kombination mit hochauflösender Massenspektrometrie (UHPLC/HRMS) und semi-präparativer HPLC, können jedoch keine potenziellen makromolekulare Vorläuferlipide nachgewiesen werden. Zusätzlich können nur frei-extrahierbare und nicht-extrahierbare gebundene Lipide identifiziert werden, welches im Gegensatz zu zuvor veröffentlichten Studien steht. Eine darauf aufbauende Analyse von drei verschiedenen Spezies von *Nannochloropsis* spp. zeigt, dass Diole hauptsächlich als frei-extrahierbare Lipide vorliegen, mit einer nur geringfügigen Konzentrationserhöhung nach Anwendung von saurer Hydrolyse auf die Biomasseextrakte. Diese Ergebnisse stehen daher im Gegensatz zu früheren Veröffentlichungen und stellen die allgemeine Auffassung, dass Diole als hauptsächlich extrahierbare-gebundene Lipide vorliegen, in Frage.

Weitere systematische Fehler in der Anwendung von organischen Biomarkern können auftreten, wenn residual-gebundene Lipide in marinen Sedimenten während der Diagenese freigesetzt werden und, im Gegensatz zu frei-vorliegenden Lipiden, zu veränderten Verteilungen und Häufigkeiten von Diolen führen. Daher wurde der Einfluss des Bindungsmodus des Auftretens von Lipiden für Diole und strukturell verwandte mittelkettige Ketole (Ketole) in Sedimenten eines S5 Sapropels des östlichen Mittelmeers analysiert. Dabei konnte gezeigt werden, dass einige Diolisomere an die organische Sedimentmatrix gebunden sind, da extrahierbare-gebundene 1,14-Diole im Vergleich zu frei-extrahierbaren Diolen eine 1,5-fache Zunahme der Konzentrationen zeigen. Frei-extrahierbare Diole dominieren innerhalb des Sapropels, während Diole außerhalb des Sapropels als residual Ester-gebundene und residual glykosidische Ether- oder Amid-gebundene Lipide vorliegen. Unabhängig vom

Bindungsmodus des Auftretens von Diolen zeigen verschiedene Diol-Indizes ähnliche Trends über die Tiefe des Sapropels. Ketole können nur innerhalb des Sapropels nachgewiesen werden. Ihre relative Häufigkeit, ausgedrückt durch den Diol Oxidation Index (DOXI), welcher zur Untersuchung von Auswirkungen der post-depositionalen Oxidation herangezogen wird, zeigt jedoch keine signifikanten Änderungen.

Der Einfluss der post-depositionaler Oxidation auf Diole und Ketole wurde zusätzlich im f-Turbidit der Tiefsee-Ebene von Madeira untersucht. Eine höhere Präservierung des C₃₀ 1,15-Diols und des C₃₀ 1,15-Ketols im Vergleich zu anderen Diolen und Ketolen liegt vor, wenn der obere oxidierte Teil des Turbidits, welcher durch post-depositionale Oxidation beeinflusst wurde, mit dem unteren anoxischen Teil verglichen wird. Während der LDI mit zunehmender Sauerstoffexposition erhöhte Werte zeigt, zeigen der Diol-Index und der FC₃₂ 1,15-Diol-Index, ein Proxy für den terrigenen Eintrag in marine Ablagerungsmilieus, verringerte Werte. Daher ist wahrscheinlich, dass alle drei Indizes durch post-depositionale Oxidation beeinflusst werden, was auf eine präferentielle Konservierung der C₃₀ 1,15- und C₃₂ 1,15-Diole im Vergleich zu 1,13- und 1,14-Diolen zurückzuführen ist. Im Vergleich zu anderen organischen Proxies und Biomarkern, im Besonderen GDGTs, der BIT-Index und der TEX₈₆-Index, weisen Diole und Ketole eine höhere Präservierung auf, wenn sie von post-depositionaler Oxidation beeinflusst werden. Bei veränderlichen Sauerstoffexpositionszeiten, kann es daher zu unterschiedlichen Beeinflussungen von organisch-terrestrischen und Meeresoberflächentemperatur-Proxies kommen, welche bei der Anwendung von organisch-geochemischen Proxies in stark oxidierten Sedimenten berücksichtigt werden müssen.

Die Ergebnisse dieser Arbeit stellen eine erste tiefgründigere Analyse dar, wie Diole und Diol-Proxies durch Diagenese und Sequestrierung beeinflusst werden. Es konnte gezeigt werden, dass die oxidative Degradation und der Bindungsmodus des Auftretens von Lipiden zu unterschiedlichen Veränderungen von Diolen und Diol-Proxies sowohl in Algenbiomasse als auch marinen Sedimenten führt. Diese systematischen Fehler müssen daher in der Anwendung dieser Proxies in Paläo-Rekonstruktionen berücksichtigt werden.

Samenvatting

De huidige klimaatverandering en de daarmee geassocieerde opwarming van de aarde worden veel besproken in de wetenschap, politiek en media. Om de toekomstige klimaatontwikkeling adequaat te kunnen voorspellen zijn we afhankelijk van klimaatmodellen die getest zijn op het accuraat reconstrueren van het vroegere klimaatcondities. Hiertoe dienen de klimaatarchieven van de geschiedenis van de aarde, opgeslagen in bijvoorbeeld mariene sedimenten, uitgelezen te worden door toepassing van klimaatproxies. Deze klimaatproxies zijn gebaseerd op bewaard gebleven materiaal in o.a. mariene sedimenten dat de toenmalige omgevings- en klimaatomstandigheden kan reconstrueren. Toepassing van klimaatproxies helpt bij het begrijpen van vroegere en huidige klimaatveranderingen en op hun beurt geven ze inzicht in mogelijke toekomstige klimaatveranderingen. Een verscheidenheid aan anorganische en organische chemische proxies wordt gebruikt om paleomilieuparameters te reconstrueren, waaronder de oppervlaktezeewatertemperatuur (OZT), de concentratie van CO₂ in de atmosfeer, het zoutgehalte in de oceaan en de mariene primaire productiviteit. Organische proxies zijn gebaseerd op gepreserveerde sedimentaire lipiden, biomarkers genoemd, en dienen als indicatoren voor de vroegere aanwezigheid van specifieke groepen organismen, biogeochemische processen, en omgevingsparameters. Twee vaak toegepaste organische OZT-proxies zijn de U^K₃₇-index, die gebaseerd is op de mate van onverzadiging van alkenonen met lange ketens geproduceerd door algen vallend in de groep van de haptofyten, en de TEX₈₆-index, die gebaseerd is op het aantal cyclopentane-groepen in glycerol dialkyl glycerol tetraether lipiden (GDGTs), membraanlipiden van Thaumarchaeota, een belangrijke groep van oerbacteriën in de oceaan.

Onlangs zijn nieuwe proxies op basis van langketige diolen (LKDs) ontwikkeld om OZT, productiviteit, mate van opwelling en de aanlevering van terrestrische materiaal in mariene milieu's te reconstrueren. LKDs bestaan uit lange lineaire alkylketens met een aantal koolstofatomen dat gewoonlijk varieert van C₂₈ tot C₃₂ en twee alcoholgroepen die op het eerste koolstofpositie en het midden van de koolstofketen (op positie C-12 tot C-15) gepositioneerd zijn. Deze componenten worden aangetroffen in een grote verscheidenheid aan mariene en lacustriene milieu's. Verzadigde en enkelvoudig onverzadigde 1,14-diolen worden geproduceerd door *Proboscia* diatomeeën in kustzeeën en de oceaan. De 1,13- en 1,15-diolen zijn aangetroffen in culturen van Eustigmatofyten algen, b.v. *Nannochloropsis salina* of *Nannochloropsis oculata*, maar in een andere distributie dan in het mariene milieu. Hun mariene producenten zijn dan ook onbekend. De 1,13- en 1,15-diolen komen gewoonlijk voor als extraheerbare gebonden lipiden en slechts in kleine hoeveelheden als vrij extraheerbare lipiden in algenbiomassa. Er is aangetoond dat zij de bouwstenen vormen van de zeer alifatische en onoplosbare biomacromoleculen die algenaan worden genoemd en die deel uitmaken van de celwand. Verschillende op LKD gebaseerde proxies, zoals de Langketige Diol Index (LDI) om OZT te reconstrueren, en de Diol Index om productiviteit en opwelling te bestuderen, zijn ontwikkeld en worden in toenemende mate gebruikt bij paleoreconstructies. Er blijven echter nog veel onzekerheden bestaan over hun toepassing. Mariene organische proxies worden beïnvloed door een groot aantal chemische en fysische processen vanaf het moment dat de biomarkers worden geproduceerd in het oppervlaktewater van de oceaan, tijdens het transport en de bezinking van de componenten door de waterkolom en vervolgens tijdens de begraving in het sediment. De invloed van deze diagenetische veranderingen, zoals bijvoorbeeld door oxidatieve degradatie of de manier van voorkomen, is aangetoond voor verschillende organische proxies, maar is nog niet in detail bestudeerd voor LKDs en diol-proxies. Het onderwerp van dit proefschrift is het ontstaan van mogelijke complicaties die kunnen ontstaan wanneer diol-proxies worden beïnvloed door diagenetische veranderingen.

Om de invloed van zuurstofblootstelling op de concentratie en distributie van LKDs en de LDI vast te stellen, werd gekweekte biomassa van *Nannochloropsis oculata* gedurende 271 dagen aëroob geïncubeerd. De concentraties werden bepaald van LKDs en vetzuren in verschillende verschijningsvormen: vrij extraheerbaar, extraheerbaar-gebonden, residuaal ester-gebonden en residuaal gebonden via glycosidisch ether of amide bindingen. De concentratie aan verzadigde en meervoudig onverzadigde vetzuren daalde aanzienlijk met toenemende incubatietijd, ongeacht de manier waarop zij gebonden waren. LKDs vertoonden daarentegen een toename in concentratie vanaf incubatiedag 125 tot het einde van de incubatie op dag 271, in het bijzonder voor residueel ester-gebonden LKDs, die maar liefst een 30-voudige concentratieverhoging vertoonden. Deze toename in concentratie weerspiegelt waarschijnlijk het vrijkomen van LKDs uit het biopolymeer algenaan na partiële afbraak door zuurstof. In tegenstelling tot de grootschalige toename van de LKD concentratie, bleven de LDI waarden stabiel gedurende het afbraakexperiment. De blootstelling aan zuurstof heeft dus waarschijnlijk verwaarloosbare invloed op de LDI waarden, hoewel deze wel afhankelijk zijn van de manier waarop de LKDs in de biomassa waren gebonden.

Om vast te stellen of langdurige blootstelling aan zuurstof effect heeft op het biopolymeer algenaan, werden residuen die overbleven na extractie en verzeeping en hydrolyse van de aëroob afgebroken biomassa van *Nannochloropsis oculata*, geanalyseerd met behulp van stapsgewijze pyrolyse. Het pyrolysaat van het algenaan werd gekenmerkt door een reeks *n*-alkenen / *n*-alkanen met een ketenlengte van C₁₀ tot C₃₃ en twee maxima bij respectievelijk C₁₅ tot C₁₇ en C₂₆ tot C₂₈. Ketonen waren aanwezig met ketenlengtes van C₂₈ tot C₃₄ met een maximum op C₃₁ en hadden de ketogroep in het midden van de keten op positie 14. Bovendien werd er relatief veel van de C₁₉ *n*-alkaan-2-on gevonden. Deze resultaten wijzen er op dat het algenaan opgebouwd is uit C₃₂ 1,15-diol-eenheden, verbonden via etherbindingen. Dit is in overeenstemming met eerder gepubliceerde studies. Analyse van residuen op verschillende tijdstippen gedurende de aërobe afbraak suggereert dat de algenaan-structuur niet substantieel veranderde. Dit geeft aan dat blootstelling aan zuurstof geen grote invloed heeft op de algenaanstructuur.

Aangezien LKDs een bestanddeel vormen van het algenaan en zij pas vrijkomen bij hydrolyse van de algenextracten, is geprobeerd potentiële macromoleculaire bronnen van LKDs in *Nannochloropsis* sp. biomassa te identificeren. De toepassing van ultrahoge druk vloeistofchromatografie in combinatie met hoge resolutie massaspectrometrie (UHPLC/HRMS) en semi-preparatieve HPLC leidde echter niet tot de identificatie van potentiële macromoleculaire precursorlipiden. Bovendien konden alleen vrij extraheerbare en niet extraheerbare gebonden LKDs worden geïdentificeerd, in tegenstelling tot eerder gepubliceerd onderzoek. Analyse van drie andere *Nannochloropsis* soorten toonde aan dat LKDs ook hier voornamelijk voorkomen als vrij extraheerbare lipiden en dat kleine verhogingen van de concentratie pas gedetecteerd werden na zure hydrolyse van de biomassa-extracten. Deze resultaten trekken daarom de gangbare gedachte dat LKDs voornamelijk voorkomen als extraheerbare gebonden lipiden in twijfel.

Een mogelijke complicatie bij de toepassing van organische biomarkers kan optreden wanneer residueel gebonden lipiden in mariene sedimenten vrijkomen tijdens diagenese en daarmee de concentraties en distributies van de veel meer bestudeerde vrij-voorkomende lipiden beïnvloeden. Daarom werd de invloed van de manier van voorkomen op LKDs en structureel gerelateerde keto-olen met de keto-groep in de middenpositie geanalyseerd in mariene sedimenten van het oostelijke Middellandse Zeegebied waar de S5-sapropeel onderdeel van uit maakte. Sommige LKDs bleken in belangrijke mate gebonden te zijn aan de organische sedimentaire matrix, aangezien extraheerbaar-gebonden voorkomende 1,14-diolen een anderhalfvoudige toename in concentratie vertoonden in vergelijking met vrij extraheerbare LKDs. Terwijl vrij extraheerbare LKDs domineren in de sapropeel, komen buiten de sapropeel

LKDs alleen residuaal voor (zowel ester-gebonden als glycosidisch ether- of amide-gebonden). De verschillende LKD-indices vertoonden vergelijkbare trends met sapropeeldiepte, ongeacht de manier waarop de LKDs gebonden zijn. Keto-olen waren alleen aanwezig in de sapropeel en niet in de sedimenten onder of boven de sapropeel. Keto-olen worden wel eens gezien als intermediären in de oxidatie van LKDs maar hun relatieve concentratie, gekwantificeerd m.b.v. de Diol Oxidation Index (DOXI), vertoonde geen significante verandering in de sapropeel.

Daarnaast werd de invloed van oxidatie na afzetting op LKDs en keto-ols bestudeerd in de f-turbidiet van de Abyssale vlakte van Madeira. Vergelijking van lipiden in het bovenste geoxideerde deel van het turbidiet, beïnvloedt door oxidatie na afzetting, met het anoxische onderste gedeelte van de turbidiet, toonde een hogere preservatie van het C₃₀ 1,15-diol en C₃₀ 1,15-keto-ol vergeleken met de andere LKDs en keto-olen. Terwijl de LDI verhoogd was vertoonden de Diol Index en FC₃₂ 1,15-diol, een proxy voor de rivierinvoer in mariene omgevingen, verlaagde waarden bij toenemende zuurstofblootstelling. Alle drie indices werden dus beïnvloed door oxidatie na afzetting, waarschijnlijk als gevolg van een preferentiële preservatie van de C₃₀ 1,15- en C₃₂ 1,15-diol t.o.v. de 1,13- en 1,14-diolen. LKDs en keto-olen blijven beter bewaard na oxidatie na afzetting dan GDGTs met implicaties van op GDGTs gebaseerde organische proxies, d.w.z. de BIT en TEX₈₆-index. Veranderingen in de zuurstofblootstellingstijd in sedimenten kunnen daarom veranderingen veroorzaken in terrestrische en OZT organische proxies en daarom is voorzichtigheid is geboden bij hun toepassing in sterk geoxideerde sedimenten.

De resultaten van dit proefschrift vormen een eerste stap om de invloed van diagenese en manier van voorkomen op LKDs en LKD-proxies te bepalen. Zowel oxische degradatie als de wijze van voorkomen van LKDs, zowel in algenbiomassa als in mariene sedimenten, is aangetoond en heeft invloed op de waarden van LKD-proxies. Dit kan mogelijk tot complicatie in hun toepassing leiden en hier zal rekening mee moeten worden gehouden in hun toepassing in paleomilieu-reconstructies.

Chapter 1

Introduction

1.1. Climate reconstruction and temperature proxies

Over the last decades, the current climate change, especially the global warming induced by an anthropogenic input of greenhouse gases, is an important topic widely discussed in science (IPCC, 2014), politics and media. We depend on climate models to understand the current climate as well as to predict future climate development. These climate models are validated and tested on appropriate climate archives of Earth's history. Examples of such archives are tree rings, ice cores, speleothems as well as terrestrial and lacustrine sediments, but especially marine sediments as they provide the longest and continuous climate records of Earth's history. To extract useful information from these archives, climate records and climate proxies are applied, which are preserved environmental parameters standing in for direct measurements and which provide information about the environmental and climate conditions under which sediments were deposited. The application of these proxies therefore helps in understanding past and present climate changes, which in turn gives a potential insight into future climate development.

A variety of inorganic and organic chemical proxies are used to reconstruct paleoenvironmental parameters such as sea surface temperature (SST), the concentration of CO₂ in the atmosphere (pCO₂) as well as ocean salinity and marine primary productivity. For example, the oxygen isotopic ratio $\delta^{18}\text{O}$ and Mg/Ca ratio of foraminiferal tests are common inorganic proxies to reconstruct past SST (Urey, 1947; Epstein et al., 1953; Emiliani, 1954; Shackleton, 1974; Nürnberg et al., 1996; Rosenthal et al., 1997; Bemis et al., 1998). Organic proxies are based on preserved sedimentary lipids, termed biomarkers. These biomarkers serve as indicators for the past presence of certain (groups of) organisms and sometimes biogeochemical processes as well as certain environmental parameters (Peters et al., 2005). One of the earliest developed proxies was an SST proxy based on the degree of unsaturation of long chain alkenones produced by haptophyte algae, i.e. the U^K₃₇ index (de Leeuw et al., 1980; Volkman et al., 1980; Brassell et al., 1986; Prahl and Wakeham, 1987). Other proxies for SST reconstruction include the TEX₈₆ index (Schouten et al., 2002, 2013), based on the number of cyclopentane moieties in glycerol dialkyl glycerol tetraether lipids (GDGTs) biosynthesized by Thaumarchaeota. All these inorganic and organic proxies have, however, their own limitations and can be impacted by a variety of factors, e.g. effects of diagenesis, lateral transport, seasonality and productivity (Hoefs et al., 1998a; Gong and Hollander, 1999; Ohkouchi et al., 2002; Schouten et al., 2004; Weijers et al., 2006, 2009; Mollenhauer et al., 2008; Kim et al., 2009a, 2009b; Rontani et al., 2013). This is why a multiproxy approach is usually preferred for the reconstruction of past climate and why new proxies are continuously being developed. Among the newer proxies are the long-chain diols (LCDs) which are used for reconstructions of SST, productivity and terrestrial input.

1.2. Distribution and sources of long-chain diols and mid-chain keto-ols

LCDs consist of long alkyl chains with carbon numbers ranging from C₂₆ to C₃₆ with two alcohol groups at the first carbon position and in a mid-chain position at carbon position 12 to 16 (de Leeuw et al., 1981; Volkman et al., 1992; Versteegh et al., 1997). Most frequently, LCDs are detected as compounds with carbon chain lengths of C₂₈ to C₃₂ and alcohol moieties at 1,13, 1,14 and 1,15 (Figs. 1A, B). After their initial discovery in Black Sea sediments (de Leeuw et al., 1981), LCDs have been recognized in a large variety of marine (Smith et al., 1983; Ten Haven and Rullkötter, 1991; Ten Haven et al., 1992; Yamamoto et al., 1996; Versteegh, 1997, 2000; Pearson et al., 2001; Rampen et al., 2007; 2008, 2012; Van Soelen et al., 2010; Willmott et al., 2010; Naafs et al., 2012) and lacustrine environments (Robinson et al., 1986; Cranwell et

al., 1987; Xu et al., 2007; Shimokawara et al., 2010; Castañeda and Schouten, 2011; Romero-Viana et al., 2012; Villanueva et al., 2014).

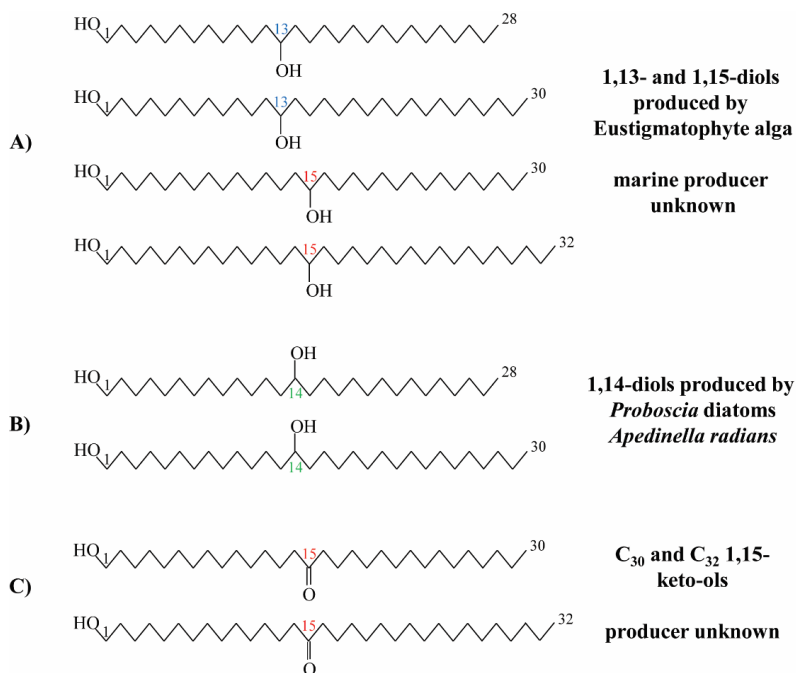


Fig. 1: Chemical structure of A) 1,13- and 1,15-diols which are produced by *Eustigmatophyte* alga although the marine producer remains unknown, B) 1,14-diols which are produced by *Proboscia* diatoms and C) C₃₀ and C₃₂ 1,15-keto-ols for which the producers are unknown.

LCDs were first proposed to be produced by cyanobacterial blooms of *Aphanizomenon flos-aquae* (Morris and Brassell, 1988), which could not be confirmed in laboratory studies of both *A. flos-aquae* or other cyanobacteria cultures (de Leeuw et al., 1992). LCDs were, however, identified in cultures of Eustigmatophyceae algae, e.g. *Nannochloropsis salina*, *Nannochloropsis oculata*, *Eustigmatos vischeri* or *Vischeria helvetica* (Volkman et al., 1992; Gelin et al., 1996; 1997a, 1997b; Rampen et al., 2014a), often showing a dominance of C₃₂ 1,15-diols (Volkman et al., 1992; Méjanelle et al., 2003; Rampen et al., 2014a). This predominance of C₃₂ 1,15-diols differs from the distribution of LCDs in marine surface sediments, which are generally characterized by a dominance of the C₃₀ 1,13- and 1,15-diols (Versteegh et al., 1997, 2000; Rampen et al., 2012, 2014a). Therefore, the role of Eustigmatophyceae as producers of LCDs in marine environments remains uncertain. Contrastingly, LCDs produced by cultured freshwater Eustigmatophyceae are in general agreement with the LCD distribution in lacustrine environments (Volkman et al., 1999; Shimokawara et al., 2010; Rampen et al., 2014a; Villanueva et al., 2014) suggesting that they are the predominant producers of LCDs in lacustrine systems. This was confirmed by a comparison of the identity and abundance of Eustigmatophyceae 18S rRNA genes and the distribution of LCDs in the water column of Lake Challa (East Africa) (Villanueva et al., 2014).

The 1,13- and 1,15-diols (Fig. 1A) have been shown to occur as mostly extractable-bound and only in minor abundances as freely extractable lipids in algal biomass (Volkman et al., 1992) and can be released from the biomass by saponification. The LCDs have also been shown to form the building blocks of the highly aliphatic and insoluble biomacromolecules termed algaenans (Derenne et al., 1989; Tegelaar et al., 1989b; de Leeuw and Largeau, 1993; Gelin et al., 1996, 1997b; Blokker et al., 1998b, 1999, 2006; Obeid et al., 2014). These algaenans are thought to be important contributors to kerogens by selective preservation (Largeau et al., 1986; Goth et al., 1988; Tegelaar et al., 1989b). Algaenan has been analyzed by both thermal (Derenne et al., 1992a; de Leeuw and Largeau, 1993; Gelin et al., 1996, 1997b, 1999) and chemical degradation (Blokker et al., 1998b, 2000, 2006) methods and is thought to be built up of mainly 1,15-diols in Eustigmatophyte algae, linked by ether bonds. It is part of the cell wall, which has been characterized to be bi-layered, comprising both an algaenan and a cellulose-like layer. The algaenan layer has been shown to contain carboxyl, aldehyde, ketone and ester functional moieties (Scholz et al., 2014).

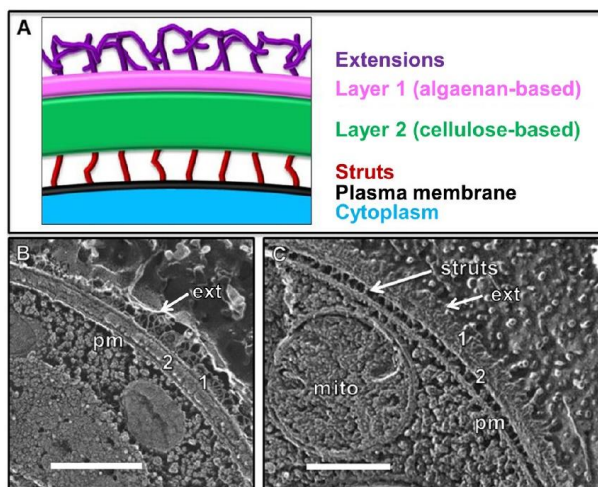


Fig. 2: A) Model of cell wall showing a bi-layer structure comprising a cellulose-based and algaenan-based layer. B, C) Quick-freeze deep-etch electron microscopy images of *Nannochloropsis gaditana* cell wall (Scholz et al., 2014).

In contrast to 1,13- and 1,15-diols, saturated and mono-unsaturated 1,14-diols have been shown to be produced by *Proboscia* diatoms (Fig. 1B), i.e. *Proboscia alata*, *Proboscia indica* and *Proboscia inermis* (Sinninghe Damsté et al., 2003; Rampen et al., 2007). They are believed to be the likely producers of 1,14-diols in marine environments based on e.g. sediment trap studies in the Arabian Sea (Rampen et al., 2007). The C₂₈, C₃₀ and C₃₂ 1,14-diols have also been shown to be produced by the Dictyochophyceae *Apedinella radians*, however its role in the marine environments is unclear (Rampen et al., 2011).

Along with LCDs, mid-chain keto-ols (keto-ols, Fig. 1C) are often encountered in a wide range of marine and lacustrine environments and represent compounds that are structurally very similar to LCDs (Versteegh et al., 1997, 2000; Ferreira et al., 2001; Xu et al., 2007; Bogus et al., 2012). Keto-ols are usually occurring with the same carbon chain lengths and position of

functional group when compared to LCDs (Fig. 1C). In contrast to LCDs, however, producers of keto-ols have not been identified and only one study has been able to detect keto-ols in trace amounts in *Nannochloropsis gaditana* (Méjanelle et al., 2003). This has led to the idea that they are produced by oxidation in the water column and sediment (Ferreira et al., 2001; Sinnighe Damsté et al., 2003).

1.3. Paleoenvironmental proxies based on the distribution of LCDs

Two important proxies using LCD distributions have been proposed, i.e. the Long-chain Diol Index (LDI) and the Diol Index, and are discussed in detail below.

The LDI uses the fractional abundances of the C₂₈ 1,13-, C₃₀ 1,13- and C₃₀ 1,15-diols to determine past SSTs due to a strong positive relationship of these LCDs with SST (Rampen et al., 2012). It is calculated as follows:

$$\text{LDI} = [\text{C}_{30} \text{ 1,15-diol}] / \{[\text{C}_{28} \text{ 1,13-diol}] + [\text{C}_{30} \text{ 1,13-diol}] + [\text{C}_{30} \text{ 1,15-diol}]\} \quad [1]$$

The LDI was calibrated by evaluation of the distribution of 1,13- and 1,15-diols in 162 marine surface sediments, that were globally distributed, and covered an SST range of -3 to 27 °C (Fig. 3). This yielded the following correlation:

$$\text{with SST} = (\text{LDI} - 0.095) / 0.033 \quad (R^2 = 0.97; \text{RSME} = 2^\circ\text{C}) \quad [2]$$

The first downcore applications in marine environments have been shown to be promising as the trends of the LDI records agree well with those of other temperature proxies (Naafs et al., 2012; Rampen et al., 2012; Lopes dos Santos et al., 2013; Smith et al., 2013; Rodrigo-Gámiz et al., 2014). So far the LDI has been applied to sediments as old as the early Pleistocene (Naafs et al., 2012). However, a potential application further back in time might be possible due to the occurrence of 1,13- and 1,15-diols, used for the calculation of the LDI, in sediments from the Miocene (Ahmed et al., 2000) and the Cretaceous-Tertiary boundary (Yamamoto et al., 1996). The LDI was also tested in lacustrine environments by analyzing the LCDs in lake surface sediments (Rampen et al., 2014a). However, only a weak correlation was observed between the LDI and annual mean air temperature, a proxy for lake surface temperature, which was probably due to different LCD producers in the different lakes. Therefore, this proxy is probably restricted to marine environments.

The 1,14-diols have been proposed to be used for the reconstruction of upwelling and productivity conditions (Rampen et al., 2008) since 1,14-diols have been shown to be produced by *Proboscia* diatoms, which are predominant in upwelling regions with high nutrient conditions. Based on this, the Diol Index was introduced (Rampen et al., 2008):

$$\text{Diol Index I} = [\text{C}_{28} + \text{C}_{30} \text{ 1,14-diol}] / \{[\text{C}_{28} + \text{C}_{30} \text{ 1,14-diol}] + [\text{C}_{30} \text{ 1,15-diol}]\} \quad [3]$$

High values of the Diol Index I were shown in association with the Southwest monsoon at the onset of upwelling, when applied to the Arabian Sea, and the Diol Index I followed the same trends of other proxies reconstructing upwelling intensities, i.e. organic carbon content, barium/aluminum ratios and stable isotopic composition of foraminifera (Rampen et al., 2007, 2008). The Diol Index I was later redefined by Willmott et al. (2010) for the Western Bransfield Basin in Antarctica as the C₃₀ 1,15-diol was present in very low concentration, in contrast to the C₃₀ 1,13-diol which was more abundant. It is calculated as follows:

$$\text{Diol Index II} = [\text{C}_{28} + \text{C}_{30} \text{ 1,14-diol}] / \{[\text{C}_{28} + \text{C}_{30} \text{ 1,14-diol}] + [\text{C}_{28} \text{ 1,13-diol}] + [\text{C}_{30} \text{ 1,13-diol}]\} \quad [4]$$

Both indices have since been applied in a variety of study locations but uncertainties with respect to their validity remain. For example, *Proboscia* diatoms have been shown to not only bloom during periods of upwelling, but also during pre-upwelling, and even in stratified water column conditions (Contreras et al., 2010). It is therefore likely that Diol Index I and II do not represent upwelling or high nutrient conditions as such but rather are affected by various factors which influence *Proboscia* ecology.

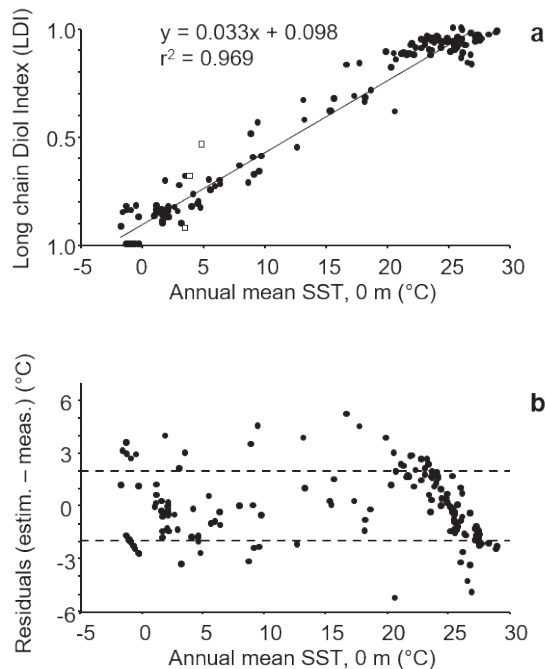


Fig. 3: a) LDI core-top calibration, correlation of LDI values of marine surface sediments with annual mean sea surface temperatures. b) Correlation of annual mean sea surface temperatures versus residual temperature errors. (Rampen et al., 2012).

1.4. Sequestration and diagenesis of long-chain diols

Although the above described LCD indices have shown some promise in paleo-reconstructions, many uncertainties remain. Marine organic proxies are impacted by a large number of physical and chemical changes from the time the proxy signal is generated by the producer. These changes may take place during their transport and settling through the water column and their subsequent burial in the sediment. These diagenetic changes can lead to changes in the composition and distribution of the organic proxies, e.g. by selective degradation of one compound over the other. Here, two important diagenetic processes, i.e. oxic degradation and matrix sequestration, are discussed.

1.4.1 Oxidic and anoxic degradation

A large number of studies have analyzed the role of oxygen in the preservation of marine organic matter, in particular oxygen exposure time (Hedges et al., 1999) as well the structure and packaging of sediments (Sun and Wakeham, 1994; Canuel and Martens, 1996). A large variability in degradation rates of lipids have been shown and different lipid biomarker classes may behave significantly different with increasing levels of oxygen exposure (Hoefs et al., 2002; Sinninghe Damsté et al., 2002a). While these degradation processes have been studied in laboratory experiments by oxidation of biomass of microbes producing biomarker lipids (Harvey and Macko, 1997a, 1997b; Teece et al., 1998; Grossi et al., 2001), a number of studies have analyzed the impact of oxidic degradation in natural settings. Commonly, the latter is done by comparing the abundance and distribution of lipid biomarkers in anoxic and oxidic marine and lacustrine environments, i.e. in the water column and suspended particulate matter (Kim et al., 2009a; Rontani et al., 2009; Schouten et al., 2012), at the interface of water column and surface sediments (Canuel and Martens, 1996; Sun and Wakeham, 1998; Lengger et al., 2014) or in sediment cores (Cranwell, 1981; Sun and Wakeham, 1994; Hartnett et al., 1998; Hoefs et al., 1998a, 1998b, 2002; Gong and Hollander, 1999; Prahl et al., 2003; Huguet et al., 2009; Bogus et al., 2012; Lengger et al., 2012, 2013). Several locations and regions have been studied extensively due to their unique setting and formation history. For example, Mediterranean sapropels have been studied to assess post-depositional oxidation resulting from oxidation of organic matter by downward moving oxygen fronts induced by the change of redox state of the bottom water (i.e. from anoxic during sapropel deposition to oxidic after sapropel deposition) (Ferreira et al., 2001; Menzel et al., 2003). Post-depositional oxidation has furthermore also been studied in organic matter-rich turbidites of the Madeira Abyssal Plain, which are influenced by downward diffusing oxygen from oxygen rich abyssal waters (Hoefs et al., 1998a, 1998b, 2002; Huguet et al., 2009; Lengger et al., 2013). Sediments were initially deposited under low oxygen conditions on the shelf and subsequently transported as turbidites and deposited in oxygen-rich bottom waters of the Madeira Abyssal Plain, allowing oxygen to diffuse into the turbidite.

Up to now, only a few studies investigated the impact of oxidic and anoxic degradation on LCD abundance and distribution. Sun and Wakeham (1994) reported LCDs to be highly resistant to anoxic degradation and showed an almost 100% burial rate in Black Sea sediments. Relatively high preservation efficiencies of LCDs upon oxygen exposure compared to other biomarkers were shown in sediment cores from the Arabian Sea by Sinninghe Damsté et al. (2002), while Hoefs et al. (2002) showed a higher preservation of bound LCDs in comparison to freely extractable LCDs in turbidites of the Madeira Abyssal Plain. Furthermore, the impact of post-depositional oxidation on LCDs in S1 sapropel sediments of the Mediterranean Sea was studied and a diagenetic link between keto-ols and LCDs suggested, based on the idea that keto-ols are likely intermediate products formed by post-depositional oxidation of LCDs (Ferreira et al., 2001). To study the impact of post-depositional oxidation and paleo-oxicity, the Diol Oxidation Index (DOXI) was thus proposed by Ferreira et al. (2001) and is calculated as follows:

$$\text{DOXI} = [\text{keto-ol}] / \{[\text{keto-ol}] + [\text{diol}]\} \quad [5]$$

Grossi et al. (2001) incubated *Nannochloropsis salina* anaerobically for 442 days, showing small increases in LCD concentration over shorter timescales, believed to be related to a release of LCDs from the algaenan, and suggesting only low degradation rates for LCDs under anaerobic conditions.

LCDs seems thus to be relatively well preserved when impacted by oxidic and anoxic degradation, however the impact of these processes on LCD indices remains unclear.

1.4.2. Matrix sequestration and mode of occurrence of LCDs

Several organic biomarkers are known to not only occur as freely extractable compounds but also bound to the sedimentary matrix by e.g. ester, glycosidic, amide or ether bonds (Goossens et al., 1986, 1989b; Gelin et al., 1997b).

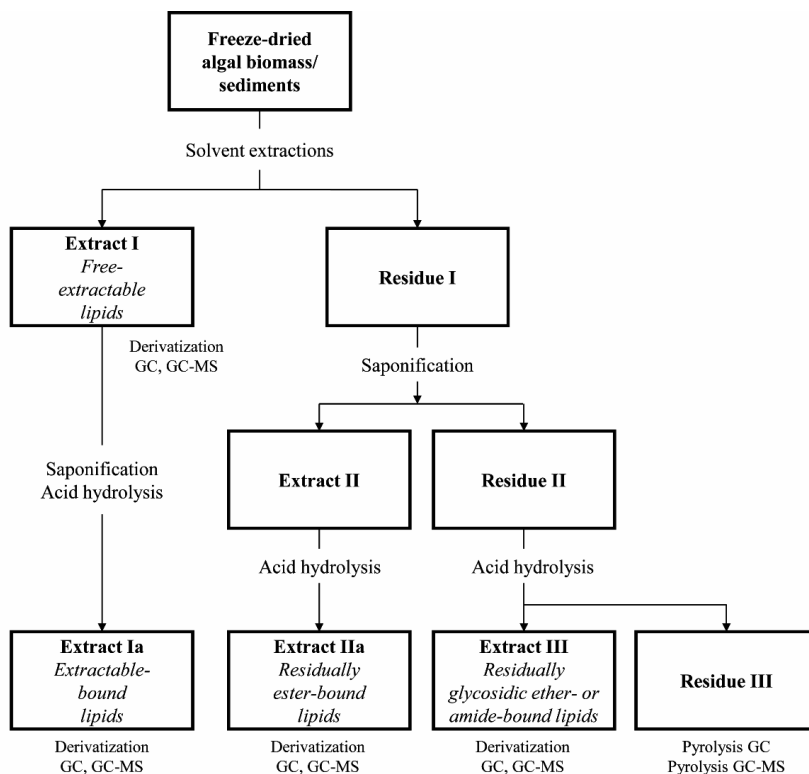


Fig. 4: Analytical scheme for analysis of mode of occurrence in algal biomass and sediments. Modified from Goossens et al. (1989b) and Gelin et al. (1997a).

These compounds are released from the sedimentary matrix by saponification and acid hydrolysis of the residue left after extraction (Fig. 4) and sometimes yield different distributions and abundances than freely occurring biomarker. If these residually-bound biomarkers are released during diagenesis, they may represent a potential bias in the application of the relevant proxies (Schouten et al., 2004; Pancost et al., 2008).

While LCDs have so far been mostly analyzed as freely extractable compounds in sediments, several studies have shown that they do occur bound to sedimentary organic matter by ester or glycosidic-ether bonds and can be released by chemical methods, i.e. saponification and acid hydrolysis. Residually ester-bound C_{30} - and C_{32} 1,15-diols and keto-ols were identified in sediments from the Monterrey Formation (Ahmed et al., 2000), while LCDs ether-linked to long-chain compounds and small amounts of ester-bound LCDs accounted for 0 – 30% of total LCDs in Lake Baikal sediments (Shimokawara et al., 2010). Additionally, Shimokawara et al. (2010) showed that the C_{32} 1,15-diol was the most abundant freely occurring LCD, while, in

contrast, the C₃₀ 1,13-diol was the most abundant ester- and ether-bound LCD. However, it is unclear how much of the LCDs are occurring bound to the sedimentary matrix and whether they impact the application of LCD proxies.

1.5. Objectives and outline of the thesis

As outlined above, LCDs occur ubiquitously in a variety of environments and proxies based on these compounds show great promise when applied to reconstruct environmental conditions. However, it is apparent that uncertainties remain, which may impact our understanding of LCDs and LCD proxies. While LCDs are increasingly used in paleo-reconstructive studies, few studies as of yet have analyzed how LCDs and related proxies are impacted by diagenesis and sequestration. Therefore, this thesis aims to evaluate the impact of diagenesis, especially of oxic degradation and mode of occurrence, on LCDs and LCD proxies. To this end, LCDs were studied in aerobically degraded biomass of *Nannochloropsis oculata* in order to constrain the impact of oxic degradation on the distribution and abundance of LCDs, the LDI and the algaenan. Furthermore, the identification of possible macromolecular sources of LCDs was attempted. Finally, the impact of mode of occurrence of differently bound LCDs and the impact of post-depositional oxidation were studied in sediments from a sapropel from the Eastern Mediterranean and from a turbidite from the Madeira Abyssal Plain, respectively.

Chapter 2 presents a large-scale aerobic incubation of *Nannochloropsis oculata* biomass over 271 days in an effort to constrain the impact of oxygen exposure on LCDs and the LDI. For this, the concentrations of free-extractable, extractable-bound, residually ester-bound and residually-bound glycosidic ether- or amide-bound fatty acids and LCDs were determined. A significant impact of oxygen exposure was determined for saturated and polyunsaturated fatty acids, showing substantially decreasing concentrations over oxic incubation time, irrespective of their mode of binding. In contrast, LCDs showed a large-scale increase in concentration from incubation day 125 until the end of incubation at day 271, in particular for residually ester-bound LCDs which showed a 30-fold increase in concentration. This increase in concentration upon oxygen exposure is thought to reflect a release of LCDs from the highly aliphatic, non-hydrolysable biopolymer algaenan. LDI values remained stable throughout the degradation experiment, irrespective of the large increases in LCDs, albeit showing differences in values dependent on how the LCDs were bound to the biomass. Therefore, the impact of oxygen exposure on the LDI is likely negligible.

Chapter 3 represents a follow up on chapter 2, studying the impact of oxic degradation on the algaenan of *Nannochloropsis oculata*. Residues left after extraction and saponification and hydrolysis of aerobically degraded biomass (chapter 2) were analyzed by step-wise pyrolysis. The pyrolyzate of the algaenan was characterized by a series of *n*-alkenes/ *n*-alkanes with chain lengths of C₁₀ to C₃₃ and two maxima at C₁₅ to C₁₇ and C₂₆ to C₂₈. Mid-chain ketones were present with chain lengths of C₂₈ to C₃₄ with a maximum at C₃₁ and a mid-chain keto group at position 14. Additionally, a pronounced C₁₉ *n*-alkan-2-one could be shown. These results point to an algaenan built-up of C₃₂ 1,15-diol units, linked by ether bonds, which is in agreement with previously published studies. Analysis of residues at different time points throughout oxic degradation suggested that there were no large alterations of the algaenan structure, indicating that oxygen exposure has a no substantial impact on the structure of the algaenan.

In **Chapter 4** we attempted to identify macro-molecular sources of LCDs in *Nannochloropsis sp.* biomass since LCDs in algal biomass have been reported to mainly be released after hydrolysis of the extract and are thought to be a constituent of the biopolymer algaenan. However, application of ultra-high pressure liquid chromatography combined with high resolution mass spectrometry (UHPLC/HRMS) and semi-preparative HPLC did not lead to the

identification of potential macromolecular precursor lipids. Additionally, only freely extractable and not extractable-bound lipids, as commonly reported, could be identified. We subsequently analyzed 3 different strains of *Nannochloropsis spp.* and could show that LCDs mainly occur as freely extractable lipids and minor increases in concentration were only detected after application of acid hydrolysis on the biomass extracts. These results therefore call into question the idea that LCDs occur mainly as extractable-bound lipids.

In **Chapter 5** we analyzed the impact of the mode of occurrence on LCDs and keto-ols in sediments. For this, we analyzed LCDs and keto-ols in sediments of an S5 sapropel of the Eastern Mediterranean by Bligh Dyer extraction followed by base and acid hydrolysis of the extract and the residual sediment, to release additional lipids from the sedimentary matrix. Extractable-bound 1,14-diols showed a 1.5-fold increase in abundance, when compared to freely extractable LCDs, indicating some LCDs are bound to the organic sedimentary matrix. While freely extractable LCDs dominate in the sapropel, LCDs occur as residually ester-bound and residually glycosidic ether- or amide-bound lipids outside of the sapropel. LCD indices showed similar trends irrespective of the mode of occurrence of the LCDs. Keto-ols were only detected in the sapropel and not below or above it, however their relative abundance, quantified by the DOXI index, did not show significant changes in trends.

In **Chapter 6** the impact of post-depositional oxidation on LCDs and keto-ols was studied in the f-turbidite of the Madeira Abyssal Plain. Comparison of lipids in the upper oxidized part of the turbidite, impacted by post-depositional oxidation, with the anoxic lower part of the turbidite, shows a higher preservation of the C₃₀ 1,15-diol and C₃₀ 1,15-keto-ol compared to the other LCDs and keto-ols. While the LDI showed increased values, the Diol Index and FC₃₂ 1,15-diol, a proxy for the riverine input into marine environments, showed decreased values with increasing oxygen exposure. Therefore, all three indices were impacted by post-depositional oxidation, likely resulting from a preferential preservation of the C₃₀ 1,15- and C₃₂ 1,15-diol over 1,13- and 1,14-diols. When compared to GDGTs and GDGT proxies, measured for the same turbidite, LCDs and keto-ols seem to be better preserved upon post-depositional oxidation. Additionally, while the BIT index showed increasing values with increasing oxygen exposure time, an impact of the TEX₈₆ index by post-depositional oxidation was negligible. Therefore, changes in oxygen exposure time in sediments may cause different changes in organic terrestrial and temperature proxies and care has to be taken when applying these different indices in severely oxidized sediments.

To summarize, the work described in this thesis showed that LCDs and LCD proxies are variably impacted by diagenesis and sequestration. An impact of oxic degradation was shown for incubated algal biomass, resulting in a significant release of LCDs. The source of these LCDs however, remains unclear, since the structure of the algaenan was not impacted during incubation time. However, while a significant increase in LCD concentration occurred in algal biomass when impacted by oxic degradation, the LDI was not affected by this. Intriguingly, LCDs mainly occur as freely extractable lipids, in contrast to previous studies. In sediments, LCDs are bound in different ways to the sedimentary organic matrix and are not only available as freely extractable lipids, as commonly assumed in different paleo-environmental studies. While this mode of occurrence did not really impact the application of LCD proxies, it was shown that applying LCD indices in severely oxidized sediments may lead to a preferential preservation of certain LCDs.

The results of this thesis represent a first step towards constraining the impact of diagenesis and sequestration on LCDs and LCD proxies. Further research should aim at resolving the contrasting observation on the impact of oxic degradation on the LDI as well as the role of the biopolymer algaenan as a factor impacting the distribution and abundance of 1,13- and 1,15-LCDs in marine sediments.

Chapter 2

The impact of oxygen exposure on long-chain alkyl diols and the long chain diol index (LDI) – a long-term incubation study

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Abstract

In recent years, long chain alkyl diols (LCDs) have been increasingly used to study and reconstruct past sea surface temperatures using the long chain diol index (LDI), which is based on changes in the distribution of 1,15-LCDs. However, the impact of diagenesis on LCDs and the LDI is still poorly constrained. Here we studied the impact of oxygen exposure on LCDs and the LDI, by aerobically incubating biomass of the LCD-producing alga *Nannochloropsis oculata* for 271 days. The concentrations of extractable free- and bound, residually ester-bound and residually-bound glycosidic ether- or amide-bound saturated fatty acids and LCDs were determined. A significant impact of oxygen exposure was observed for C₁₄, C₁₆ and C₁₈ saturated fatty acids and the C_{20:5} polyunsaturated fatty acid as their concentration decreased significantly over time, irrespective of their mode of binding. LCDs, in contrast, increased significantly in concentration over incubation time, e.g. up to a 30-fold increase in concentrations for residually ester-bound LCDs at day 125 compared to concentrations at the beginning of the experiment. This increase in concentration most likely represents a release of LCDs from the insoluble biopolymer algaenan due to the impact of oxygen exposure. Values of the LDI differed strongly depending on the mode of occurrence of LCDs in the biomass. However, despite the large changes in concentration of LCDs in response to oxygen exposure, the LDI remained relatively stable after prolonged degradation. This suggests that oxygen exposure may not have a substantial impact on the LDI of extractable LCDs used for its determination.

Keywords: long chain diols; long chain diol index; aerobic degradation; *Nannochloropsis oculata*; algaenan

1. Introduction

Long chain alkyl diols (LCDs) were first discovered in sediments of the Black Sea (de Leeuw et al., 1981) and since then have been studied in a large variety of marine and freshwater environments (Versteegh et al., 1997; Sinninghe Damsté et al., 2003; Rampen et al., 2007, 2008, 2014a, 2014b; Zhang et al., 2011; Romero-Viana et al., 2012; de Bar et al., 2016; Lattaud et al., 2017a). They are composed of long n-alkyl chains with chain lengths of mainly C₂₈ to C₃₂ and contain alcohol groups at C₁ and at a mid-chain position, most notably between positions C₁₂ to C₁₅ (Versteegh et al., 1997). Saturated and unsaturated C₂₈ and C₃₀ 1,14-LCDs have been shown to be produced by *Proboscia* diatoms (Sinninghe Damsté et al., 2003; Rampen et al., 2007) and C₂₈-C₃₂ 1,14-LCDs have also been identified in *Apedinella radians* (Rampen et al., 2011). C₂₈ to C₃₂ 1,13- and 1,15-LCDs have been identified in cultures of freshwater and marine Eustigmatophyte algae (Volkman et al., 1992; Gelin et al., 1997b, 1999; Méjanelle et al., 2003; Shimokawara et al., 2010; Rampen et al., 2014a; Balzano et al., 2017; Zhang and Volkman, 2017), but the diol distributions in marine environments are dissimilar to those of cultured algae (Volkman et al., 1992; Versteegh et al., 1997; Rampen et al., 2012, 2014a), suggesting unknown marine sources. LCDs in algal cell material of Eustigmatophytes and in the sediment have been shown to occur in different forms, i.e. as free-extractable, extractable-bound (released by hydrolysis of the extract) and residually-bound lipids (released by hydrolysis of the residual biomass left after extraction) (Gelin et al., 1997b; Ahmed et al., 2000; Grossi et al., 2001; Hoefs et al., 2002; Shimokawara et al., 2010). Additionally, they are also thought to be the building blocks of the insoluble biopolymer algaenan which occurs in the cell wall of Eustigmatophytes (de Leeuw et al., 1981; Tegelaar et al., 1989a; Gelin et al., 1997b; Volkman et al., 1998; Scholz et al., 2014; Zhang and Volkman, 2017).

Several indices based on LCD distributions have been proposed to reconstruct past conditions such as upwelling, oxic degradation, riverine input as well as sea surface temperatures (SST).

The long chain diol index (LDI) is based on the fractional abundances of C₂₈ 1,13-, C₃₀ 1,13- and C₃₀ 1,15-LCDs and correlates strongly with SST (Rampen et al., 2012). Upwelling indices use the relative abundances of C₂₈ and C₃₀ 1,14- to C₂₈/C₃₀ 1,13- or C₃₀ 1,15-LCDs (Rampen et al., 2008; Willmott et al., 2010). The diol oxidation index (DOXI), calculated as the ratio of 1,15-keto-ols versus LCDs, is used to study oxic degradation in sediments thereby assuming that keto-ols are oxidative products of LCDs (Ferreira et al., 2001). Finally, the %C₃₂ 1,15 of all LCDs (de Bar et al., 2016; Lattaud et al., 2017a, 2017b) is used to study riverine inputs into the marine environment.

These indices have since shown promise in paleo-environmental reconstruction studies and have been applied in different marine environments (Pancost et al., 2009; Willmott et al., 2010; Rampen et al., 2012, 2014b; Seki et al., 2012; Lopes dos Santos et al., 2013; Rodrigo-Gámiz et al., 2014, 2015, 2016; Nieto-Moreno et al., 2015; Plancq et al., 2015).

Certain uncertainties in the application of LCD indices still remain to be addressed. Only a few studies have considered the impact of diagenesis on the abundance and distribution of LCDs (Ferreira et al., 2001; Versteegh et al., 2010; Bogus et al., 2012). As shown by various studies, biomarker distributions can be altered substantially by oxygen exposure (Sun and Wakeham, 1994; Hoefs et al., 1998b; Prahl et al., 2003; Rontani et al., 2009, 2013), but only Sinninghe Damsté et al. (2002), Hoefs et al. (2002) and Rodrigo-Gámiz et al. (2016) analyzed the impact of oxic degradation on LCD distributions. Sinninghe Damsté et al. (2002) analyzed sediment cores from the northern Arabian Sea studying the impact of oxygen exposure time on the preservation potential of various biomarkers, finding relatively high preservation efficiencies of LCDs. Hoefs et al. (2002) studied free and bound lipid biomarkers in oxidized and unoxidized parts of Madeira Abyssal Plain turbidites, finding higher preservation of bound LCDs compared to free LCDs. Analysis of Arabian Sea surface sediments deposited under contrasting redox conditions by Rodrigo-Gámiz et al. (2016) showed a strong decrease in concentration of LCDs with increasing oxygen exposure as well as a slightly higher degradation rate of 1,15-LCDs compared to 1,14- and 1,13-LCDs. This resulted in a drop of the LDI from 0.94 to 0.85 with increasing water depth. Additionally, the DOXI ratios showed no distinctive trends over water depth and only slight increases in value with increasing oxygen concentration were observed. The impact of anoxic degradation on LCDs has been constraint by a few studies, most notably by Sun and Wakeham (1994) and Grossi et al. (2001). Sun and Wakeham (1994) reported a high level of resistance to anoxic degradation during early diagenesis and an almost 100% burial efficiency of LCDs in Black Sea sediments. Grossi et al. (2001) incubated *Nannochloropsis salina* anaerobically over 442 days, revealing short-term increases (degradation days 50 to 100) in LCD concentrations, which were proposed to be caused by a release of LCDs from the algaenan. Following this, concentrations of LCDs decreased over time, albeit to a lesser extent than other associated compounds, such as alkenes, phytol, sterols and fatty acids. Both studies thus show relatively low degradation rates for LCDs under anoxic conditions.

Here we present a laboratory study analyzing the impact of oxygen exposure on LCD concentrations and distributions in *Nannochloropsis oculata* biomass, which was aerobically incubated for 271 days. The concentrations of free and bound LCDs and major other lipids were monitored over degradation time. Furthermore, the impact of oxic degradation on the LDI was constrained.

2. Material and methods

2.1. Set up of aerobic biomass incubations

A large batch of the LCD-producing alga *Nannochloropsis oculata*, strain CCMP525, was ordered from Reed Mariculture Inc and kept frozen at 20 °C until the set-up of incubation. 20 g of defrosted biomass were transferred to Erlenmeyer flasks each containing 10 g of sediment and 210 ml of sea water, both collected from the Dutch Wadden Sea in March 2015 and acclimatized after collecting for 24h at 25°C. 42 Erlenmeyer flasks were prepared and placed into a shaker table in the dark, at 25 °C and stirred at 100 rpm for 271 days under aerobic conditions. Sea water levels were monitored weekly and bi-distilled water was added when required to keep the water volume constant and to counteract evaporation effects.

2.2. Lipid analysis

At predetermined sampling days, three Erlenmeyer flasks were randomly selected. For each flask, the water layer was separated from the particulate matter by centrifugation at 3000 rpm for 120 min. The particulate matter was subsequently freeze dried and processed using the analytical scheme outlined in Fig. 1.

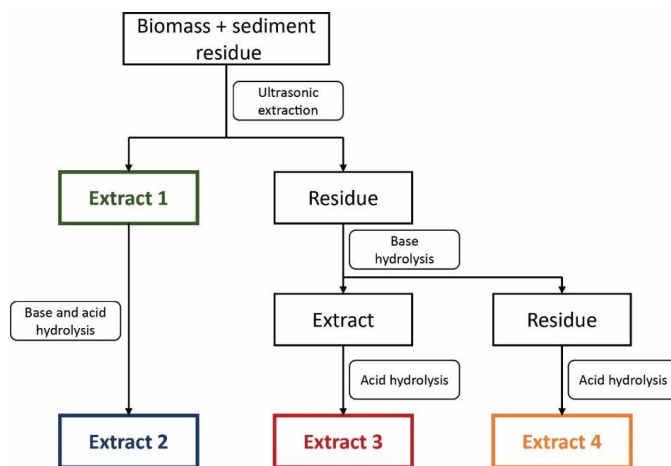


Fig.1: Analytical scheme representing the work up steps of *Nannochloropsis oculata* biomass + sediment, as described in 2.2.

The particulate matter was ultrasonically extracted with methanol/dichloromethane (MeOH/DCM) (2:1, v/v, 4x). DCM phases were combined and dried using rotary evaporation and drying under N₂. The extract resulting from this work-up step was labelled Extract 1 (E1).

An aliquot of E1 was subsequently saponified by refluxing with 1 M KOH in MeOH for 1 h. The DCM phase was separated from the H₂O/MeOH phase and dried under N₂. This fraction was subsequently acid hydrolyzed in a last step by refluxing with 1.5 M HCl in MeOH for 3 h. After cooling, the mixture was neutralized using 2 N KOH in MeOH and extracted with DCM/H₂O (1:1, v/v, 3x). The DCM phases were separated from the H₂O/MeOH phase and

combined to be dried under N₂. The base and acid hydrolyzed E1 extract is termed Extract 2 (E2).

The residue obtained after ultrasonic extraction of the particulate phase of the incubation experiment was dried under N₂ and saponified by refluxing with 1 M KOH in MeOH for 1 h. The mixture was centrifuged and the liquid phase was collected into a separatory funnel and acidified to a pH of 3 using 2 N HCl in MeOH. The residue was subsequently extracted using H₂O/MeOH (1:1, v/v), MeOH (2x) and DCM (3x) and all solvents were combined into the separatory funnel. The DCM layer was separated from the MeOH/H₂O layer and collected, the MeOH/H₂O layer was extracted with DCM (3x) and the combined DCM phases were dried under N₂. This extract was subsequently acid hydrolyzed using the conditions described above and yielded Extract 3 (E3). In a last step, the saponified residues were acid hydrolyzed by refluxing with 1.5 M HCl in MeOH for 3 h. Solvents were collected into a separatory funnel after cooling and neutralized using 2 N KOH in MeOH. Residues were subsequently extracted using H₂O/MeOH (1:1, v/v), MeOH (2x) and DCM (3x) and solvents were collected into the separatory funnel. The DCM layer was separated from the MeOH/H₂O layer and collected, the MeOH/H₂O layer was extracted with DCM (3x) and the combined DCM phases were dried under N₂. The extracted residue after base hydrolysis was subsequently also acid hydrolyzed following the procedures described above yielding Extract 4 (E4).

E1-E4 were eluted with DCM over a small pasteur pipette containing Na₂SO₄ to remove remaining salts. An internal standard C₂₂ 7,16 - diol (MolPort SIA, Latvia) was added to aliquots of E1-E4 which were subsequently methylated with diazomethane in diethylether. Very polar components were removed by eluting the methylated E1-E4 with ethyl acetate over a small pasteur pipette containing silica gel.

Prior to analysis by gas chromatography (GC) and gas chromatography/ mass spectrometry (GC/MS), E1-E4 were silylated by adding BSTFA [N,O-bis(trimethyl-silyl)trifluoroacetamine] and pyridine and heated up to 60 °C for 20 min. Subsequently, samples were dissolved in ethyl acetate.

2.3. GC-MS analysis of LCDs

Samples were analyzed for LCDs on an Agilent 7890B gas chromatograph coupled to an Agilent 5977A mass spectrometer, equipped with a fused silica capillary column (25 m x 0.32 mm) coated with CP Sil-5 (film thickness 0.12 µm). Samples were injected at 70 °C oven temperature, increasing to 130 °C at 20 °C/min and then 320 °C at 4°/min which was held for 25 min with a helium flow of 2 mL/min. The mass spectrometer operated with an ionization energy of 70 eV. LCDs were quantified in selective ion monitoring (SIM) of m/z 187, 313, 339 and 341 characteristic for the different LCDs (Versteegh et al., 1997; Rodrigo-Gámiz et al., 2015). Additional compounds, e.g. fatty acids, polyunsaturated fatty acids, phytol and sterols, were identified on the same GCMS in full scan (m/z 50-800) and analyzed on an Agilent 7890B gas chromatograph with helium as the carrier gas at a flow of 2 ml per minute using the same temperature program as that of the GCMS. Fatty acids were identified by comparison with published mass spectra and quantified by integration of respective peak areas of fatty acids and the internal standard. The calculated concentrations given in this paper were normalized on dry weight of the biomass, in mg/g or µg/g.

The LDI was calculated as follows (Rampen et al., 2012):

$$\text{Long chain Diol Index (LDI)} = \frac{F_{C_{30}1,15\text{-diol}}}{F_{C_{28}1,13\text{-diol}} + F_{C_{30}1,13\text{-diol}} + F_{C_{30}1,15\text{-diol}}} \quad [1]$$

3. Results

Biomass of the LCD-producing alga *Nannochloropsis oculata* was incubated in the presence of sediment and water from the North Sea under aerobic conditions for a period of 271 days. Four extracts (E1-E4), each representing lipids occurring in the biomass in a different way, were obtained (Fig. 1) following the work-up procedures of e.g. Gelin et al. (1997) or Goossens et al. (1986). E1 contains freely occurring and –extractable lipids, while E2 contains free-extractable lipids from E1 as well as extractable lipids which are ester-bound or bound by glycosidic ether- or amide bonds. E3 represents residually ester-bound lipids while E4 contains residually-bound glycosidic ether- or amide-bound lipids (Goossens et al., 1986, 1989a). LCDs were detected in all fractions. Both E1 and E2 contained saturated and polyunsaturated fatty acids, phytol and sterols, predominantly cholesterol and minor amounts of $C_{29}\Delta^{5,24}$ sterols. E3 contained saturated and polyunsaturated fatty acids, α -hydroxy and β -hydroxy fatty acids with chain lengths of C_{24} to C_{26} and C_{12} to C_{18} , respectively, and to a lesser extent sterols, predominantly cholesterol. E4, was mostly comprised of α - and β -hydroxy fatty acids with chain lengths of C_{24} to C_{26} and C_{12} to C_{18} , respectively, and to a lesser extent, of saturated fatty acids. Long chain keto-ols were not detected and mid-chain hydroxy methyl alkanates were only present in trace amounts. This lipid composition is in good agreement with previous studies of the lipids of *Nannochloropsis* species (Volkman et al., 1992, 1993; Olofsson et al., 2012; Mitra et al., 2015; Balzano et al., 2017; Zhang and Volkman, 2017). Analysis of the sediment and sea water from the Dutch Wadden Sea used, did not show the presence of LCDs. Saturated fatty acids and the $C_{20:5}$ PUFA were present in low abundances in the Wadden Sea sediment and water and are ignored here. Additionally, the pure biomass of strain CCMP525 was analyzed to confirm that the reported LCDs in this study were solely derived from this alga.

In this study, we will mainly focus on LCDs, as well as the dominant saturated fatty acids and the $C_{20:5}$ polyunsaturated fatty acid (PUFA), to contrast their different responses to oxygen exposure.

3.1 Fatty acids

C_{14} , C_{16} and C_{18} saturated fatty acids were identified in E1-E4. The concentrations of summed saturated fatty acids ($C_{14} + C_{16} + C_{18}$) were highest in E1 and E3 with 7.9 ± 1.4 and 10.1 ± 1.6 mg/g respectively (Fig. 2A). The concentration of free-extractable saturated fatty acids (E1) decreased by 98% during the first 35 days of incubation and stayed low at 0.2 ± 0.1 mg/g throughout the remainder of the experiment. Free- and extractable-bound (E2) saturated fatty acids initially increased during the first 3 days of incubation up to 30.7 ± 8.3 mg/g and subsequently showed a continuous slow decrease until reaching a plateau of 6.1 ± 0.7 mg/g at 169 incubation days for the remainder of the experiment. The concentration of residually ester-bound (E3) saturated fatty acids decreased slowly, reaching concentrations of 2.9 ± 0.1 mg/g by the end of the experiment. Residually-bound glycosidic ether- or amide-bound (E4) fatty acids stayed low in concentration at 0.3 ± 0.1 mg/g throughout the experiment.

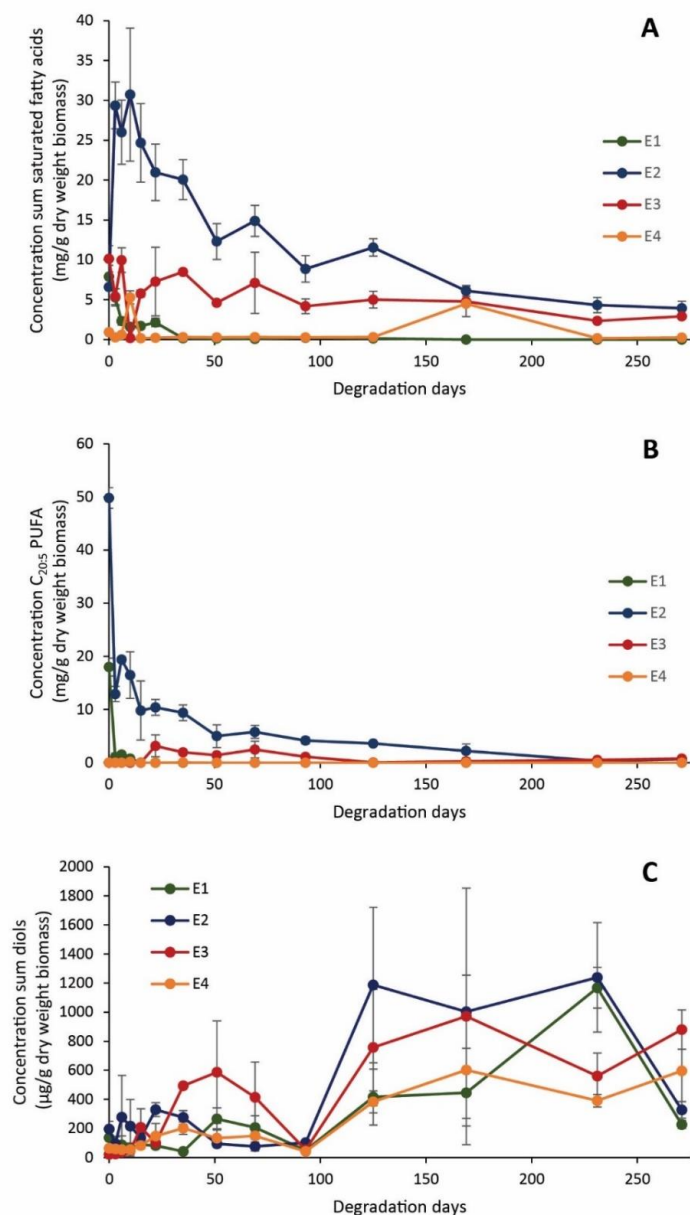


Fig. 2: Concentrations of lipids over time in an incubation experiment where biomass of the Eustigmatophyte alga *Nannochloropsis oculata* is exposed to oxygen in the presence of sediment and sea water: A) concentration of the sum of saturated fatty acids, B) concentration of C_{20:5} PUFA, C) concentration of the sum of LCDs. Concentrations are given over degradation time for free-extractable (E1), free- and extractable-bound (E2), residually ester-bound (E3) and residually-bound glycosidic ether- or amide-bound compounds (E4). Error bars represent standard deviations calculated from triplicate incubations.

The C_{20:5} PUFA is also a prominent lipid in addition to saturated fatty acids in *N. oculata* biomass. In this incubation study it occurred in the free-extractable (E1) and free- and extractable-bound (E2) extracts, but was not identified in E3 and E4 (Fig. 2B). Its concentration in the free-extractable (E1) form decreased rapidly during the first 15 days of incubation from initial values of 18.0 ± 1.7 mg/g to below detection limits. The concentration of free- and extractable-bound (E2) C_{20:5} PUFA decreased rapidly from 49.8 ± 2.0 mg/g to 12.9 ± 1.4 mg/g during the first 3 days and stayed approximately constant until 35 days. Subsequently, values slowly decreased until the end of incubation time to concentrations of 0.7 ± 0.1 mg/g. The residually ester-bound C_{20:5} PUFA was first detected at day 22 of incubation at 3.2 ± 2.1 mg/g, but subsequently decreased and was not detected from 125 days onwards.

3.2 Long chain alkyl diols

Four major LCDs were identified in this incubation study at the beginning of incubation, i.e. the C₂₈ 1,13-, C₃₀ 1,13-, C₃₀ 1,15- and C₃₂ 1,15-LCD. The C₂₈ 1,13-LCD (Fig. 3A) showed concentrations of 0.7 ± 0.1 µg/g (E1), 1.1 ± 0.2 µg/g (E2), 0.4 ± 0.2 µg/g (E3) and 0.3 ± 0.0 µg/g (E4) and was the least abundant of the four major LCDs. The concentration of the C₃₀ 1,13-LCD (Fig. 3B) was 7.5 ± 4.7 µg/g (E1), 19.7 ± 4.7 µg/g (E2), 2.9 ± 1.6 µg/g (E3) and 4.4 ± 1.7 µg/g (E4), while the C₃₀ 1,15-LCD (Fig. 3C) showed concentrations of 5.2 ± 2.3 µg/g (E1), 11.0 ± 2.9 µg/g (E2), 1.4 ± 0.7 µg/g (E3) and 4.2 ± 1.3 µg/g (E4). The C₃₂ 1,15-LCD (Fig. 3D) was the most dominant LCD with 124 ± 61 µg/g (E1), 163 ± 47 µg/g (E2), 17.4 ± 8.7 µg/g (E3) and 55.0 ± 26.7 µg/g (E4). Additionally, a low abundance of the C_{32:1} 1,15-LCD was detected.

The concentration trends of LCDs over time developed in two phases and individual LCDs show similar concentration patterns. In the first phase, spanning the first 93 incubation days, summed LCD concentrations increased and then decreased to a minimum at 93 degradation days (Fig. 2C). Free-extractable (E1) and free- and extractable-bound (E2) LCDs decreased in concentration during the first 3 days of incubation by 42% and 52% and subsequently increased to reach maximum concentration at days 51 with 270 ± 80 µg/g for free-extractable (E1) LCDs and maximum concentration at day 22 with 330 ± 50 µg/g for free- and extractable-bound (E2) LCDs. Residually ester-bound (E3) LCDs showed the highest maximum concentrations in this first phase at day 51 with 590 ± 350 µg/g, while residually-bound glycosidic ether- or amide-bound (E4) LCDs stayed overall lower and showed highest concentration at day 35 with 200 ± 40 µg/g.

In the second phase, summed LCD concentrations reached even higher concentrations. At day 125, concentrations were substantially higher compared to initial values at day 0 of incubation time: double in concentration to 420 ± 190 µg/g in the E1 fraction, a five-fold increase to 1190 ± 540 µg/g in fraction E2, a thirty-fold increase to 760 ± 400 µg/g in fraction E3 and a five-fold increase to 380 ± 80 µg/g in fraction E4. After this maximum, concentrations stayed approximately constant until the end of the experiment for residually ester-bound (E3) and glycosidic ether- or amide-bound (E4) LCDs, while both free-extractable (E1) and free- and extractable-bound (E2) LCDs decreased sharply at the end of the experiment to concentrations of 230 ± 30 and 330 ± 60 µg/g respectively.

In two instances, at incubation days 51 and 69, the concentrations of LCDs in E2 were below concentrations of E1, which is unexpected as hydrolysis should potentially result in additional release of LCDs. However, in most cases the majority of LCDs are present as free-extractable LCDs and the amount released by base and acid hydrolysis is relatively low. Possibly the lower amount of LCDs after the hydrolysis steps are due to work up losses during consecutive hydrolyses or artifact formation (unpublished data).

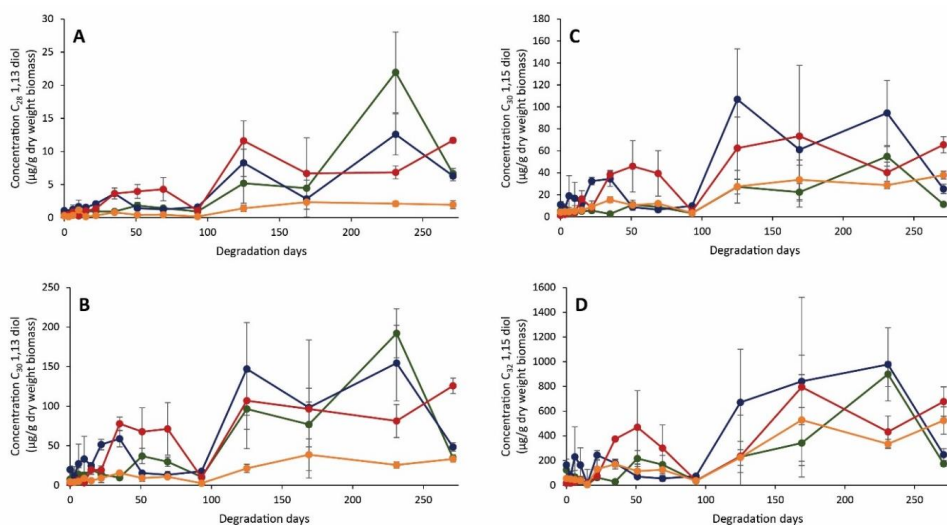


Fig. 3: Concentrations of LCDs over time in an incubation experiment where biomass of the Eustigmatophyte alga *Nannochloropsis oculata* is exposed to oxygen in the presence of sediment and sea water: C_{28} 1,13- (A), C_{30} 1,13- (B), C_{30} 1,15- (C) and C_{32} 1,15-LCDs (D) as free-extractable compounds (E1 in green), free- and extractable-bound compounds (E2 in blue), residually ester-bound compounds (E3 in red) and residually-bound glycosidic ether- or amide-bound compounds (E4 in yellow). Error bars represent standard deviations calculated from triplicate incubations.

The LDI showed some variation over the course of the experiment, each fraction having distinct values (Fig. 4). The LDI calculated from residually-bound glycosidic ether- or amide-bound (E4) LCDs ranged from 0.34 ± 0.03 (day 10) to 0.55 ± 0.03 (day 125), free- and extractable-bound (E2) and residually ester-bound (E3) ranged from 0.31 ± 0.02 (day 15) to 0.40 ± 0.01 (day 125) and 0.28 ± 0.01 (day 22) to 0.52 ± 0.04 (day 10), respectively. The LDI calculated from free-extractable (E1) LCDs had the lowest values and ranged from 0.19 ± 0.03 (day 15) to 0.43 ± 0.25 (day 0). The largest variability was observed during the first 35 incubation days. The LDI for the free-extractable (E1) LCDs dropped from 0.43 ± 0.25 at the beginning of the experiment to 0.25 ± 0.02 at day 3, thereafter remaining approximately constant. The LDI for the residually ester-bound (E3) LCDs showed a maximum at day 10 of 0.52 ± 0.04 and slowly dropped to constant values by day 35 of 0.32 ± 0.00 , approximately remaining at that level for the remainder of the experiment, while the LDI for the residually-bound glycosidic ether- or amide-bound LCDs decreased from an initial value of 0.48 ± 0.01 to 0.34 ± 0.03 by day 10, recovering to initial values by degradation day 15 and staying constant throughout the rest of the experiment.

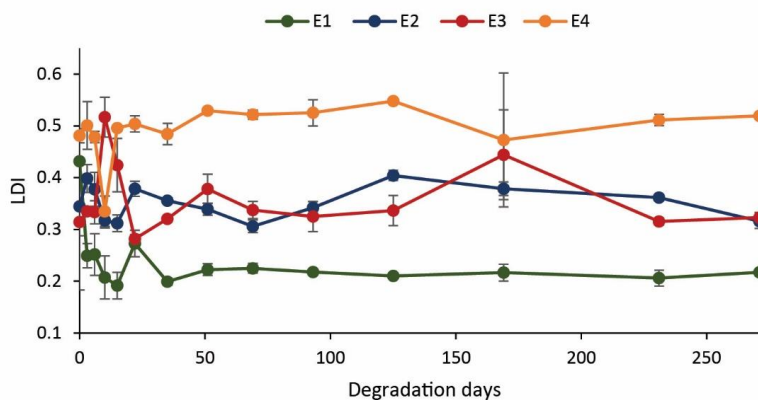


Fig. 4: LDI in an incubation experiment where biomass of the Eustigmatophyte alga *Nannochloropsis oculata* is exposed to oxygen in the presence of sediment and sea water: calculated from free-extractable (E1), free- and extractable-bound (E2), residually ester-bound (E3) and glycosidic ether- or amide-bound LCDs (E4). Error bars represent standard deviations calculated from triplicate incubations.

4. Discussion

4.1 Impact of oxygen exposure on the abundance of fatty acids and LCDs

The occurrence of C_{14} , C_{16} and C_{18} saturated fatty acids and the $C_{20:5}$ PUFA in *Nannochloropsis oculata* is consistent with earlier studies of the lipid composition of *Nannochloropsis* species (Volkman et al., 1993; Olofsson et al., 2012; Xiao et al., 2013; Mitra et al., 2015; Balzano et al., 2017). These lipids showed significant decreases in concentration when exposed to aerobic conditions during 271 days of incubation, to the point of non-detection in the case of the $C_{20:5}$ PUFA. This was independent of the mode of occurrence of these lipids (i.e. as free-extractable, free- and bound-extractable and residually ester-bound or residually-bound glycosidic ether- or amide-bound). Thus, a significant impact of oxygen exposure was observed, especially for the $C_{20:5}$ PUFA. This mirrors previous studies, which showed the relatively rapid degradation of fatty acids and PUFAs in both incubation studies and in sediment cores under oxic as well as anoxic conditions (Farrington and Quinn, 1973, 1977; Sun and Wakeham, 1994, 1997; Canuel and Martens, 1996; Harvey and Macko, 1997a; Grossi et al., 2001; Hoefs et al., 2002).

In contrast to the fatty acids, the impact of oxygen exposure on LCDs resulted in substantially different concentration trends. Following a first minor phase of increase in concentration of LCDs, which was most pronounced in the residually ester-bound fraction (Fig. 2C), a second more significant phase of increasing concentrations was observed. During this second phase from 93 days onwards, LCD concentrations increased discernibly in all four fractions and subsequently plateaued from 125 days onwards, indicating a large-scale release of LCDs from a source outside of our analytical window. The most likely source for these LCDs is the insoluble biopolymer algaenan, which is produced in substantial amounts by Eustigmatophyte algae. Based on chemical degradation methods such as RuO_4 and/or HI or thermal degradation

methods such as pyrolysis GC-MS (Gelin et al., 1997b, 1999; Blokker et al., 1998a, 2000, 2006) this algaenan is thought to be composed of LCDs linked by ether bonds (Zeliber et al., 1988; Tegelaar et al., 1989b; de Leeuw and Largeau, 1993; Gelin et al., 1997b; Blokker et al., 1998b, 2000). If so, it is difficult to envisage how oxidation of ether-linked diols would result in the release of LCDs themselves, i.e. it would be expected that oxidized products of LCDs such as dicarboxylic acids (Blokker et al., 2000) would be detected. However, Scholz et al. (2014) extended the commonly accepted algaenan structure by identifying a bi-layer cell wall composed of a cellulose layer and an algaenan layer while in addition to ether bonds, the algaenan was found to comprise carboxyl, aldehyde, ketone and ester functional groups. Possibly, the impact of oxygen exposure on the algaenan of *N. oculata* may have led to the release of LCDs bound intra-polymeric by ester bonds as opposed to ether bonds, possibly during the initial stage of break-down of the cell and cell wall containing the algaenan. This would mean that the algaenan has remained relatively intact during oxygen exposure but that weakly bonded LCDs attached to the algaenan may have been released and entered our analytical window. It is unclear why this would happen in two phases but possibly is the result of different steps in the breakdown of algal cell and cell wall. Further research examining the integrity of structure of the algaenan during oxygen exposure may shed more light on this.

At 271 degradation days, the time span our incubation was set up for, a substantial decrease of LCD concentrations can be observed in the free-extractable (E1) and free- and extractable-bound fractions (E2), while the residually ester-bound (E3) and glycosidic ether- or amide-bound (E4) fractions showed an increase in concentration. If a constant supply by release of LCDs from the algaenan and a simultaneous degradation of released LCDs by oxic degradation is assumed, free-extractable and free- and extractable-bound LCDs consequently show a higher degradation rate compared to the rate of release from the algaenan, at the end of incubation time.

In a similar incubation study, set up under anaerobic conditions, complete degradation of LCDs in *Nannochloropsis salina* was also not observed (Grossi et al., 2001) but in contrast to our results, only a transient minor increase in LCD concentrations was observed. This is possibly due to the anaerobic nature of the incubation experiment of Grossi et al. (2001), which may have led to a limited release of LCDs from the algaenan over the studied time span. A further comparison is complicated by the fact that Grossi et al. (2001) only considered LCDs that are ester-bound to the biomass and did not take into account the individual patterns of degradation of free-extractable or residually-bound glycosidic ether- or amide-bound LCDs.

The results of this incubation study potentially have implications for our understanding of the fate of LCDs in the context of organic matter settling through the water column and the subsequent burial into the sediment. On short time scales i.e. the settling of organic matter through an oxic water column which is in the order of days to weeks, oxygen exposure is not expected to exert a substantial effect on the LCD concentrations. Only on longer time scales, i.e. in the context of organic matter burial into the sediment which may take multiple years to decades, does oxic as well as anoxic degradation lead to decreasing concentrations of LCDs in the sediment, as previously shown by Hoefs et al. (2002) and Rodrigo et al. (2016).

Keto-ols were not detected in this study, indicating that they were not formed by oxidation of LCDs in the timespan that this incubation study covered. Possibly, the time frame of our study was too short to form these products or the hypothesis that keto-ols are formed by oxic degradation of LCDs is not correct.

4.2. Impact of oxidic degradation on the LDI

To examine the impact of oxidic degradation of LCD distributions we focused on the LDI although we realize that the alga species used in our study is not the main producer of LCDs in the marine environment and thus LDI values cannot be converted to temperature values following Rampen et al. (2012). Despite the large changes in concentrations of LCDs, the calculated LDI showed only some variability at the initial stages of oxygen exposure, while after 35 days, no substantial changes in the LDI were observed anymore. Values of the LDI, however, differed strongly depending on how the respective LCDs were present in the algal biomass, i.e. the LDI calculated from free-extractable LCDs (E1) showed much lower values compared to e.g. the LDI calculated from residually-bound glycosidic ether- or amide-bound LCDs (E4) (Fig. 4). Since the values for the LDI remained distinctly different between these differently bound LCDs throughout the degradation experiment, it suggests that LCDs were not transferred between the differently bound compound pools. Furthermore, a strong preferential degradation of one diol over another diol was not observed. This consistent difference in LCD distributions between the differently bound compound pools is difficult to reconcile with the hypothesis that LCDs are released from the algaenan upon oxygen exposure as it might be expected that these released LCDs would have a rather uniform distribution. Possibly, these results suggest that LCDs are bound to the algaenan in certain ratios and are also released in these ratios upon oxygen exposure.

The fact that the LDI is not substantially affected by oxidic degradation is in contrast to the study of Rodrigo et al. (2016), which showed a preferential preservation of 1,13- over 1,15-LCDs, which consequently impacted the LDI. However, the studies differ in time scales of oxidic degradation on LCDs, i.e. while our study represents a time window of less than a year, Rodrigo et al. (2016) analyzed the long term (years to decades) impact of oxidic exposure on LCDs during burial in sediments. Thus, short term exposure to oxygen, for instance during the settling of LCDs through an oxidic water column, is unlikely to alter the LDI.

5. Conclusion

Nannochloropsis oculata biomass was aerobically degraded for 271 days. While saturated fatty acids and the polyunsaturated fatty acid C_{20:5} were degraded to a large extent, the concentrations of LCDs increased over time and were probably released from the biopolymer algaenan. A substantial impact of oxidic exposure on the LDI was not observed.

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Chapter 3

The impact of oxic degradation on the algaenan of *Nannochloropsis oculata* (Eustigmatophyceae)

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Abstract

Certain freshwater and marine alga, such as Chlorophyceae and Eustigmatophyceae, have been shown to form highly aliphatic and insoluble biopolymers termed algaenan located in the cell wall, which might be preserved in sediments. However, the exact fate of these biopolymers during diagenesis is not well established. Here, in an effort to study the impact of oxic degradation on algaenans, insoluble residues left after hydrolysis of biomass of the Eustigmatophyte *Nannochloropsis oculata* (*N. oculata*) degraded under oxic conditions for up to 271 days, were analyzed by stepwise pyrolysis gas chromatography mass spectrometry. These pyrolyzates were characterized by *n*-alkene/ *n*-alkane doublets with chain lengths from C₁₀ to C₃₃ and two maxima at C₁₅ to C₁₇ and C₂₆ to C₂₈. Additionally, ketones, with a mid-chain keto group at position 14, were identified with chain lengths of C₂₈ to C₃₄ with a maximum at C₃₁. A further pronounced abundance of C₁₉ *n*-alkan-2-one was observed. These results are in line with an algaenan structure composed of C₃₂ 1,15-diol units linked by ether bonds, as proposed previously. Importantly, analysis of residues at four different time points in the oxic degradation experiment did not show a difference in the algaenan chemical composition. The structure of the algaenan of *N. oculata* is therefore relatively resistant to oxic degradation over time scales of at least 270 days.

1. Introduction

There is a long-standing debate on the composition and sources of marine sedimentary organic matter, in particular the insoluble part termed kerogen. While kerogen forms the majority of organic matter in marine sediments and has been analyzed by using a large variety of physical and chemical methods, its formation remains largely disputed (Vandenbroucke and Largeau, 2007). Research in the Seventies and Eighties favored formation of kerogen by biodegradation of proteins and carbohydrates of organic matter into amino acids and sugars and subsequent abiotic recondensation into complex insoluble geopolymers (Durand, 1980; Tissot and Welte, 1984). Later research focused on the preservation and enrichment of highly aliphatic and insoluble biomacromolecules produced by a variety of higher plants and freshwater algae (Largeau et al., 1986; Goth et al., 1988; Tegelaar et al., 1989b). Among the latter, several microalgae from the classes Eustigmatophyceae, Chlorophyceae and Dinophyceae are known to contain these selectively preserved highly aliphatic biopolymers, which are located in the cell wall and are termed algaenan (Derenne et al., 1989; de Leeuw and Largeau, 1993; Gelin et al., 1996, 1997b, 1999; Blokker et al., 1998b, 2006; Kodner et al., 2009; Obeid et al., 2014). Algaenans mostly occur in freshwater microalgae (Largeau et al., 1986; Goth et al., 1988; Zelibor et al., 1988; Tegelaar et al., 1989b; Kodner et al., 2009), e.g. *Botryococcus braunii* or *Tetraedron minimum*, and have only been detected in some marine alga, i.e. *Nannochloropsis oculata* (*N. oculata*) and other *Nannochloropsis* species (Volkman et al., 1992; Gelin et al., 1997b, 1999). While the algaenan only represents a small percentage of living cell material, its resistance against diagenesis is thought to lead to selective preservation, eventually forming the main part of kerogen (Hutton, 1987; Goth et al., 1988; Tegelaar et al., 1989b; de Leeuw and Largeau, 1993; Largeau and de Leeuw, 1995; Blokker et al., 2000; van Bergen et al., 2004). However, to the best of our knowledge, this resistance has not been tested extensively, e.g. by degradation experiments of algaenans.

Due to their insoluble nature, algaenans are often analyzed by thermally degradative methods, e.g. pyrolysis gas chromatography mass spectrometry (pyrolysis GC-MS). Analysis of algaenan by pyrolysis GC-MS gives rise to a suite of characteristic *n*-alkene/ *n*-alkane doublets (Derenne et al., 1992b; de Leeuw and Largeau, 1993; Gelin et al., 1996, 1999; Zhang and Volkman, 2017). The distribution of *n*-alkene/ *n*-alkane doublets of algaenan pyrolyzates of various

Eustigmatophytes showed maxima at C₁₅ to C₁₇ and C₂₆ to C₂₈ for *Nannochloropsis salina* (*N. salina*) and *Nannochloropsis sp.* (Gelin et al., 1997b) and maxima at C₁₀ and C₂₇ for *N. oculata* and *Nannochloropsis granulata* (*N. granulata*) (Gelin et al., 1999). Additionally, mid-chain ketones, indicating that ether cross bridges occur in the algaenan structure, are present with chain lengths varying from C₂₅ to C₃₆ in e.g. *N. salina* and *Nannochloropsis sp.* (Gelin et al., 1996, 1997b, 1999). Small amounts of prist-1-ene, toluene and traces of aromatic, isoprenoid and branched hydrocarbons were detected as well (Gelin et al., 1997b). Further information has been gained from the application of chemical degradation, i.e. RuO₄ and/ or HI/LiAlH₄ (Blokker et al., 1998b, 2000, 2006), which showed that the algaenan of Eustigmatophytes is likely composed mainly of long-chain 1,15-diols (LCDs) linked by ether bonds (Zelibor et al., 1988; Tegelaar et al., 1989b; de Leeuw and Largeau, 1993; Gelin et al., 1997b; Blokker et al., 1998b, 2000). Additionally, Scholz et al. (2014), identified a bi-layer cell wall composed of an outer algaenan and an inner cellulose layer in *Nannochloropsis gaditana* (*N. gaditana*) and used Fourier transform infra-red spectroscopy to show that the algaenan layer comprised carboxyl, aldehyde, ketone and ester functional moieties.

More recently, Zhang and Volkman (2017) analyzed the algaenan of *Nannochloropsis oceanica* (*N. oceanica*), formerly *N. oculata* (Fawley et al., 2015; Zhang and Volkman, 2017, 2018), by stepwise pyrolysis GC-MS, enabling a more in-depth analysis of the algaenan by generation of products at different temperatures. Importantly, they found a series of mid-chain ketones from C₂₉ to C₃₃, the most abundant being the C₃₁ ketone, as well as an abundant C₁₇ *n*-alkan-2-one. These results confirmed LCDs as important building blocks of the algaenan, described earlier by Gelin et al. (1997), albeit with 1,17-diols, as the building blocks of the algaenan, rather than 1,15-diols. This is in contrast with lipid analyses which showed a dominance of 1,15-diols and only trace amounts of 1,17-diols in this alga (Volkman et al., 1992).

In a recent study, we analyzed the impact of oxygen exposure on lipids of *N. oculata* by aerobic incubation of its biomass for over 271 days (Reiche et al., 2018). During degradation, the concentrations of freely-extractable, but also bound-extractable and residually bound LCDs increased over oxygen exposure time, which was hypothesized to result from a potential break-up of the algaenan and therefore a release of these differently bound LCDs. This would suggest that, in contrast to previous hypotheses, the algaenan is not stable upon oxic degradation and therefore not selectively preserved.

Here, in an effort to study the fate of algaenan during oxic degradation, we analyzed extracted and hydrolyzed biomass of *N. oculata* from the oxic degradation incubations of Reiche et al. (2018) using stepwise pyrolysis GC-MS following the approach outlined by Zhang et al. (2016) and Zhang and Volkman (2017). The results give insights into potential chemical alterations of algaenan during oxic degradation.

2. Material and methods

2.1. Oxic degradation incubations

The oxic degradation incubation of *N. oculata* was previously described in Reiche et al. (2018). Briefly, biomass of *N. oculata* strain CCMP525 (20g) was incubated aerobically with sediment (10 g) and water from the Dutch Wadden Sea (210 ml) for 271 days at 25 °C in the dark and stirred at 100 rpm on a shaker table. Forty-two slurries were prepared and at fourteen predetermined sampling days, the water layer of the biomass slurries was separated from the particulate matter by centrifugation and the obtained particulate matter was freeze dried.

2.2. Work-up procedure

Four time points, i.e. at the start of the oxic degradation experiment (t_0), at the conclusion of a first minor phase of increase in LCD concentrations (t_{93}), at the beginning of a large scale second increase in LCD concentration (t_{125}) and the end of the incubation experiment (t_{271}), were selected for the work-up (see Reiche et al. 2018 for details). The work-up procedure followed a modified approach by Gelin et al (1997) and Goossens et al. (1986).

In a first step, the particulate matter was ultrasonically extracted with methanol/dichloromethane (MeOH/ DCM) (2:1, v/v, 4x). The resulting residue was dried under N_2 and in a second step saponified by refluxing with 1 M KOH in MeOH for 1 h. The residue was subsequently extracted using H_2O / MeOH (1:1, v/v), MeOH (2x) and DCM (3x) and dried under N_2 . In a last step, the saponified residues were acid hydrolyzed by refluxing with 1.5 M HCl in MeOH for 3 h and subsequently extracted using H_2O / MeOH (1:1, v/v), MeOH (2x) and DCM (3x) and dried under N_2 . The dried, extracted, saponified and acid hydrolyzed residues were homogenized using a mortar and pestle. A blank, consisting of only sediment and sea water was worked up in the same way and prepared for analysis as outlined above.

2.3. Stepwise pyrolysis GC-MS

The residues and blanks were analyzed by sequential stepwise pyrolysis GC-MS as outlined in Zhang et al. (2016) and Zhang and Volkman (2017). Briefly, ca. 1 mg of the respective residue was placed into a deactivated stainless-steel sample cup, which was subsequently pushed into a pyrolysis quartz tube of a Frontier PY-3030D pyrolyzer and maintained for 20 s to accomplish flash pyrolysis at the pre-set temperature. After flash pyrolysis, the residue was pulled from the pyrolysis zone and a blank run was carried out at 700 °C in preparation for the subsequent higher temperature step. The sample was pyrolyzed in 4 steps at temperatures from 360 - 660 °C in 100 °C intervals. In each step, the formed pyrolyzate was directly swept into the GC injector with He as the carrier gas and analyzed using an Agilent 7890B-5977 A GC-MS system with a J&W DB-5MS column (60 m x 0.25 mm i.d., 0.25 μ m film thickness) and an ionization voltage of 70 eV in full scan mode (m/z 50-800). Peak assignment was accomplished by comparison with literature data (Volkman et al., 1992) and the library spectra.

3. Results

As described in Reiche et al. (2018), biomass of *N. oculata* with sediment and sea water was incubated in the dark at 25 °C for 271 days. Particulate matter (i.e. biomass and sediment) from four time points in the degradation experiment (t_0 , t_{93} , t_{125} and t_{271}) was collected. The time points at t_{93} and t_{125} were chosen because they represent time points where large increases in LCD concentrations were observed (Fi. 1). The particulate matter was extracted, saponified and acid hydrolyzed and analyzed by stepwise pyrolysis GC-MS at different temperatures, e.g. 360 °C to 660 °C in 100 °C steps.

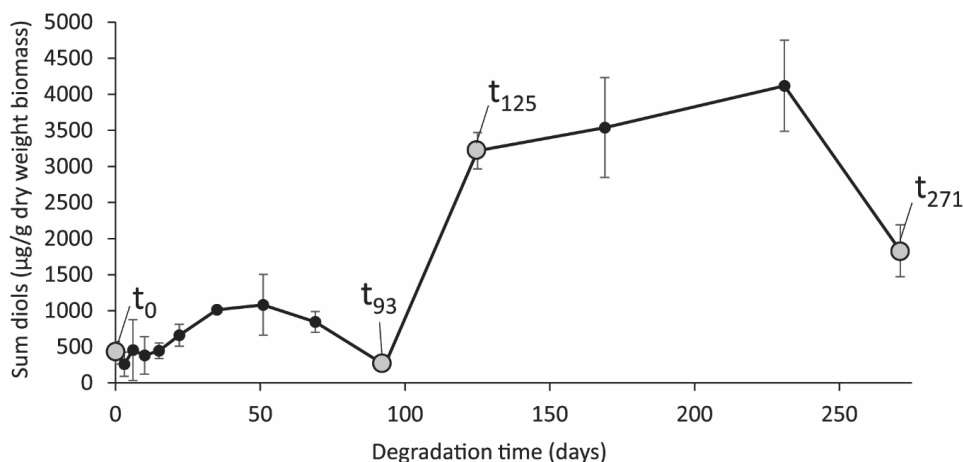


Fig. 1: Summed concentrations of LCDs over time in an incubation experiment where biomass of the Eustigmatophyte alga *N. oculata* was exposed to oxygen in the presence of sediment and water (modified from Reiche et al., 2018). T₀, t₉₃, t₁₂₅ and t₂₇₁ represent sampling points for analysis of the residue by pyrolysis GC-MS methods. Error bars represent standard deviations calculated from triplicate incubations.

For the residue at t₀, at 360 °C, C₁₆ fatty acids and a series of C₂₄ to C₃₀ fatty acids were detected. Furthermore, tocopherol and small amounts of aromatic compounds including indole and benzenepropanenitrile, could be identified. At 460 °C, a series of C₂₈ to C₃₄ mid-chain ketones with a maximum at C₃₁ and a series of *n*-alkenes/ *n*-alkanes with chain lengths from C₂₄ to C₃₀ predominated. The *n*-alkenes/ *n*-alkanes were most pronounced at 560 °C with chain lengths of C₁₀ to C₃₃ and two maxima at C₁₅ to C₁₇ and C₂₆ to C₂₈ (Fig. 2). Mid-chain ketones were again identified with chain lengths of C₂₈ to C₃₄ with a maximum at C₃₁. GC-MS analysis showed that for the C₃₁ mid-chain ketones, the keto group is predominantly at position 14, based on the major fragment ions of *m/z* 211 and *m/z* 267 resulting from cleavage either side of the carbonyl group (Fig. 3). Furthermore, an abundant C₁₉ *n*-alkan-2-one was detected (Fig. 4). At 660 °C, only minor amounts of products were detected in the pyrolyzate. A blank, which consisted of only sediment and sea water and worked up in the same way as the other samples, did not generate any significant products when analyzed by stepwise pyrolysis GC-MS, indicating that all detected pyrolysis products were derived from *N. oculata* biomass.

Comparison of the chemical composition of the pyrolyzates of t₀, t₉₃, t₁₂₅ and t₂₇₁, shows similar distributions and products for the different pyrolysis temperatures, although a few subtle differences are distinguishable (Fig. 2-4). While a slight maximum at C₁₅ *n*-alkane is visible for *n*-alkene/ *n*-alkanes at t₉₃, in contrast, t₀, t₁₂₅ and t₂₇₁ show a maximum at C₁₆ *n*-alkane in the pyrolyzate at 560 °C. Additionally, a more pronounced C₂₇ *n*-alkane maximum can be observed for t₀, whereas t₉₃, t₁₂₅ and t₂₇₁ show more equal maxima between C₂₆ to C₂₈ *n*-alkanes (Fig. 2). While for t₀ and t₉₃ mid-chain ketones showed a maximum at C₃₁ in the 560 °C pyrolyzate, a slight shift to a maximum of C₃₂ is visible at t₁₂₅ and t₂₇₁ (Fig. 3). In contrast, the C₁₉ *n*-alkan-2-one was pronounced in all incubation time points (Fig. 4).

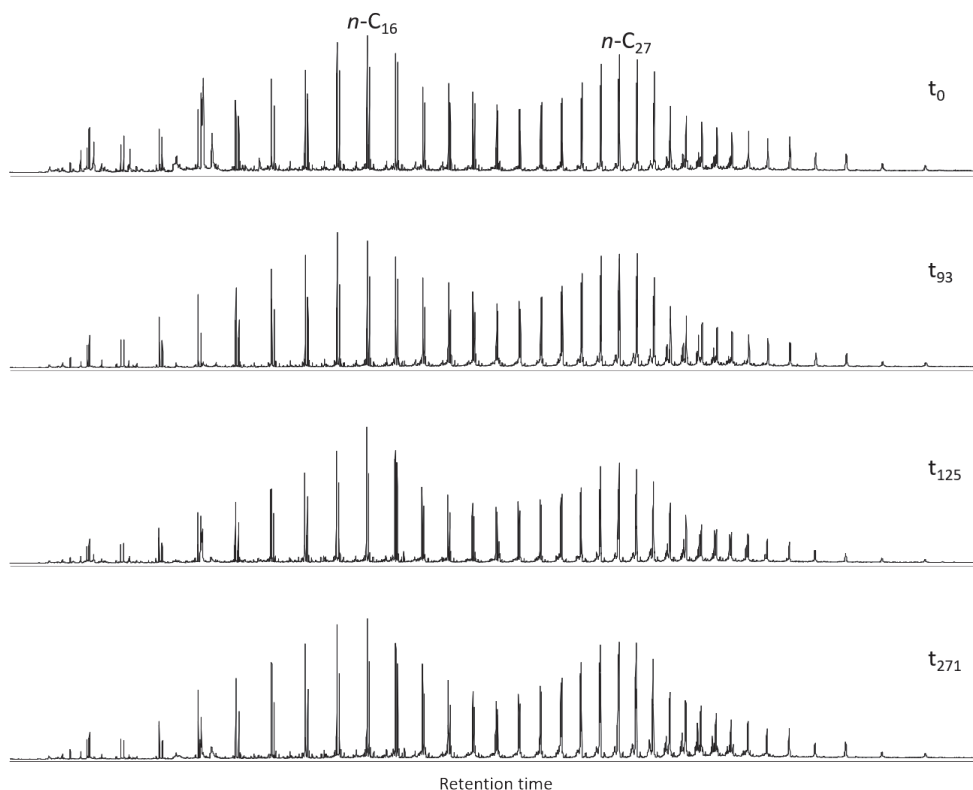


Fig. 2: Summed mass chromatograms of m/z 83 + 85 showing the distributions of n -alkene/ n -alkane pairs with chain lengths of C_{10} to C_{33} and two maxima at C_{15} to C_{17} and C_{26} to C_{28} in the pyrolyzates obtained at 560 °C at different timepoints in oxic incubation experiment.

4. Discussion

4.1. Stepwise pyrolysis results of *Nannochloropsis* biomass

We first compared the results of stepwise pyrolysis at t_0 , i.e. the start of degradation, with those obtained by Zhang and Volkman (2017) who studied the stepwise pyrolysis of biomass of *N. oceanica*. At lower temperatures (360 °C) both pyrolyzates were relatively clean, showing only low amounts of aromatic compounds, such as indole and benzenepropanenitrile, which have been interpreted to reflect pyrolysis products of proteins (Valdés et al., 2013; Biller and Ross, 2014). In contrast to Zhang et al. (2017), only small amounts of fatty acids, which represent freely extractable compounds in *Nannochloropsis* species (Reiche et al., 2018), could be detected. This is likely due to differences in the initial material that was analyzed. Zhang and Volkman (2017) analyzed freeze dried biomass of *N. oceanica* while the material used here was extracted, saponified and acid hydrolyzed resulting in a removal of most of the carbohydrates, proteins and ester- and amide-bound fatty acids. N -alkene/ n -alkane doublets were first identified at 460 °C and more clearly visible at 560 °C in our study, which is somewhat different to Zhang and Volkman (2017). That study showed the clear presence of n -alkene/ n -alkane doublets at 460 °C and 510 °C, but could not detect higher molecular weight compounds at

temperatures > 510 °C. However, in our study we analyzed the residues in temperature steps of 100 °C, i.e. 460 °C and 560 °C, while Zhang and Volkman (2017) used temperature steps of 50 °C and thus also generated pyrolysates at 510 °C. The distribution of the series of *n*-alkenes/ *n*-alkanes in our experiments (Fig. 2) is characterized by a carbon chain length range of C_{10} to C_{33} and two maxima at C_{15} to C_{17} and C_{26} to C_{28} . This is similar to the results of *N. oceanica* (Zhang and Volkman, 2017) and is representative for algaenan pyrolysates of different species of Eustigmatophytes, i.e. *Nannochloropsis sp.* or *N. granulata* (Gelin et al., 1997b).

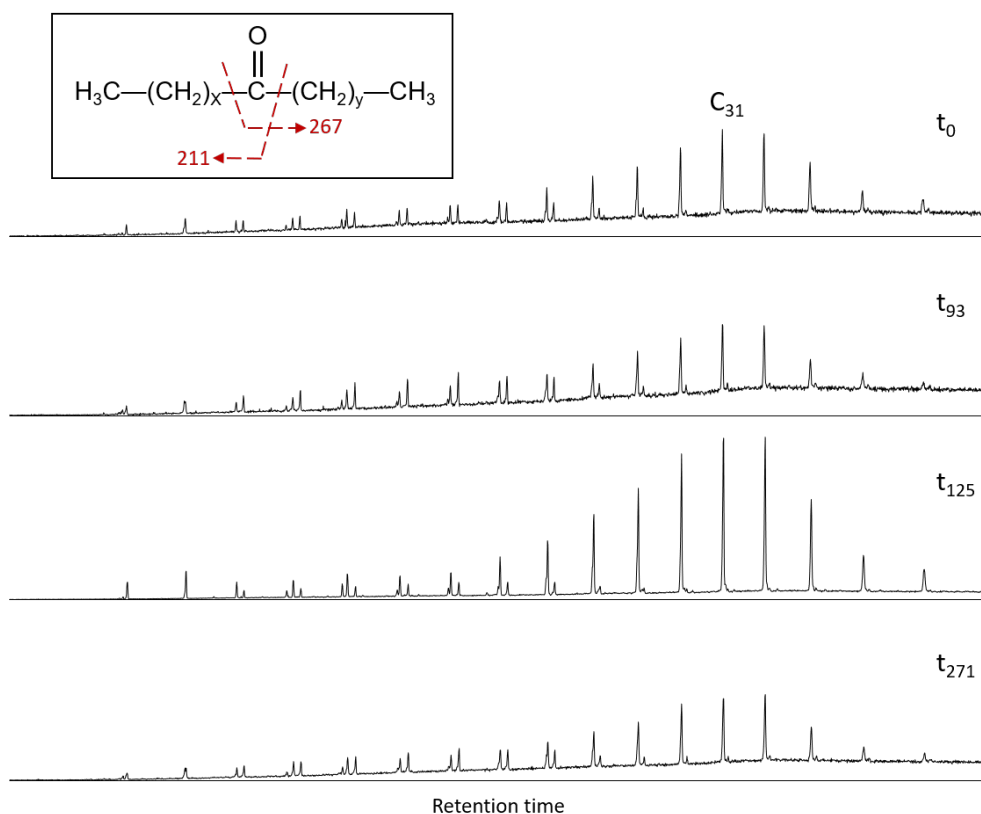


Fig. 3: Mass chromatograms of m/z 267 showing the predominance of $C_{24} - C_{35}$ mid-chain ket-14-ones (peaking at C_{31}) in the pyrolysates obtained at 560 °C at different timepoints in the oxic incubation experiment. The structural formula ($x = 16$, $y = 12$) shows the C_{31} 14 mid-chain ketone and depicts the formation of the observed major fragment ions at m/z 211 and 267 by cleavage at both sides of the carbonyl group.

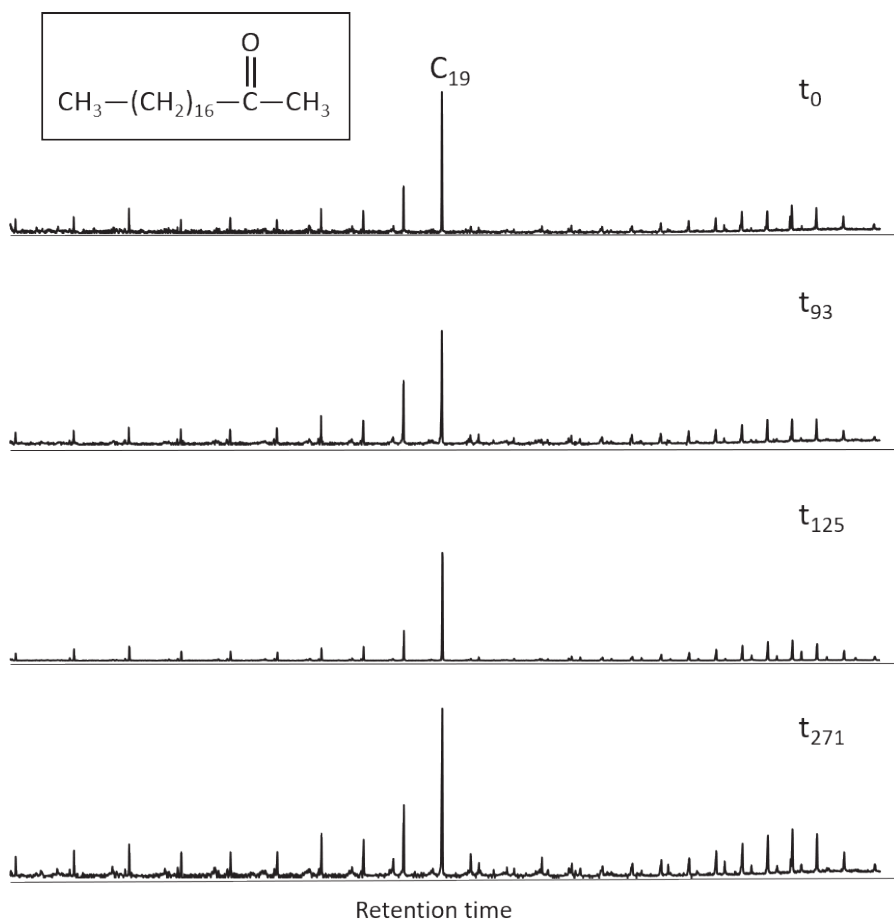


Fig. 4: Mass chromatograms of m/z 59 showing the predominance of the C_{19} n -alkan-2-one in the pyrolyzates obtained at 560 °C at different timepoints in the oxic incubation experiment. The chemical structure shown depicts the C_{19} n -alkan-2-one in the pyrolyzates obtained at 560 °C at different timepoints in the oxic incubation experiment.

Mid-chain ketones were detected first at 460 °C and were abundant at 560 °C in our study. This is different from the study of *N. oceanica* by Zhang and Volkman (2017), which showed the highest abundances of mid-chain ketones at 460 °C, but did not detect them at temperatures >460 °C. Possibly, this may be due to the different residues that were analyzed, i.e. freeze dried biomass versus extracted, saponified and acid hydrolyzed biomass. Mid-chain ketones were present with chain lengths of C_{28} to C_{34} with a maximum at C_{31} and a keto-group at position 14 in our analyzed residues. In combination with a pronounced C_{19} n -alkan-2-one detected at 560 °C, this strongly suggests that the C_{32} 1,15-diol comprised the main building block of the algaenan, bound by ether bonds. This is consistent with the results of lipid analyses of the biomass of *N. oculata* (Reiche et al., 2018), in which the C_{32} 1,15-diol was the predominant LCD, as a freely extractable, bound extractable and residually bound compound. It contrasts the results of Zhang and Volkman (2017), who found a predominance of the C_{17} n -alkan-2-one

and mid-chain ketones with the keto group at position 16 which would indicate an algaenan built up of 1,17-diols. These differences might be a result of the different species that were analyzed, e.g. *N. oculata* here and *N. oceanica* for the study of Zhang and Volkman (2017). Interestingly, *N. oceanica* was also previously analyzed by Volkman et al. (1992), who reported only small amounts of 1,17-diol isomers and the predominance of 1,15-diols in the extractable biomass. This is similar to our work on the *N. oculata* strain (Reiche et al., unpublished data 2018). Possibly, this difference may again be explained by the fact that residues were analyzed here, in contrast to Zhang and Volkman (2017) who analyzed the whole, freeze-dried biomass. Indeed, the C₁₉ *n*-alkan-2-one detected here occurred in the pyrolyzates at the temperature step of 560 °C, while the C₁₇ *n*-alkan-2-one of Zhang and Volkman (2017) was predominant at 460 °C and was not detected at temperatures higher than 510 °C, suggesting it was less well bound. Therefore, the *n*-alkan-2-one in their case may have not been generated from the algaenan, but from another hydrolysable macromolecule, which is removable by saponification and acid hydrolysis. However, most LCDs do not seem to be bound to soluble macromolecules (Reiche et al. unpublished results, chapter 4). Alternatively, these contrasting results represent real differences between different strains of *Nannochloropsis* species. Indeed, for flash pyrolyzates of *Nannochloropsis sp.* mid-chain ketones with chain lengths of C₂₂ to C₃₂, a maximum at C₂₆ and keto groups at position 17 and 18 were observed (Gelin et al., 1996). Additionally, Gelin et al. (1996) indicated the presence of *n*-alkan-2-ones with chain lengths of C₇ to C₂₅ and a maximum at C₁₉ for flash pyrolyzates of *Nannochloropsis sp.* Therefore, there may be potential differences between the basic building blocks of algaenan of different *Nannochloropsis* species and further research is needed, e.g. by application of chemical degradation.

4.2. Impact of oxidic degradation on the algaenan of *N. oculata*

In our incubation experiments, we previously showed a large-scale increase in LCD concentration when biomass of the Eustigmatophyte *N. oculata* was aerobically degraded (Fig. 1). We interpreted this large-scale increase in LCD concentration as a release of LCDs from a source outside of our analytical window, most likely the algaenan, as an impact of oxygen exposure. However, it remains chemically unclear how oxidation of ether-bound LCDs in the algaenan would lead to a release of LCDs and why the release of LCDs would happen in two phases (Fig. 1). We, therefore, chose, besides the residues at the beginning (t_0) and the end of incubation (t_{271}), residues from a first minor phase of increase in LCD concentrations (t_{93}) and the large scale second increase in LCD concentration (t_{125}) (Fig. 1) as subjects for study. When comparing the different pyrolyzates for the four different timepoints in degradation, it is however apparent that there are no major changes in the distributions of the *n*-alkenes/ *n*-alkanes (Fig. 2), mid-chain ketones (Fig. 3) and *n*-alkan-2-ones (Fig. 4). Their distribution remained approximately constant throughout the various stages of *N. oculata* biomass degradation indicating only minimal alterations in the structure of the algaenan over the course of the experiment. Since we did not determine the absolute abundance of the pyrolysis products per g of biomass, we cannot say if the algaenan decreased in abundance. However, the relative intensity of the signal of pyrolysis products remained relatively similar (Fig. 2-4). Interestingly, no additional compounds that may have been formed by oxidation of ether-bonds or other parts of the algaenan, e.g. peroxides or dicarboxylic acids (Blokker et al., 1998b, 2000), were detected when analyzed by GC-MS. Overall, our results thus suggest that oxidic degradation did not alter the composition of the algaenan itself substantially, making it unlikely that the observed releases of LCDs in the degradation experiment are derived from the core of the algaenan. Alternatively, the release of LCDs may reflect the release of weakly-bonded LCDs of the outer parts of the algaenan, which perhaps were becoming more sensitive to release during degradation of the outer structure of the algaenan. Scholz et al. (2014) have shown that the algaenan is comprised of a two-member algaenan outer layer together with a cellulose-like

layer. If during degradation, the more labile cellulose layer was degraded, parts of the algaenan may have become exposed and accessible for extraction and hydrolysis. The reason that the increase in LCD concentration was happening in two phases may reflect releases of LCDs from different parts of the algaenan. In any case our results suggest that algaenans are relatively resistant against oxic degradation as their structure is not substantially altered after >270 days of exposure to oxygen. They are therefore expected to preserve well during settling in the water column and sediment burial.

5. Conclusion

In order to constrain the impact of oxygen exposure on the chemical composition of the algaenan of *N. oculata*, residues of ultrasonically extracted, saponified and acid hydrolyzed biomass, subjected to oxic degradation, were analyzed by stepwise pyrolysis. Homologous series of *n*-alkenes/ *n*-alkanes, characteristic pyrolysis products of the algaenan, were detected with chain lengths ranging from C₁₀ to C₃₃ with two maxima at C₁₅ to C₁₇ and C₂₆ to C₂₈. Mid-chain ketones were identified with chain lengths of C₂₈ to C₃₄ with a maximum at C₃₁ and a mid-chain keto group at position 14. Additionally, an abundant C₁₉ *n*-alkan-2-one was present. This suggests that the building block of the algaenan of *N. oculata* comprised the C₃₂ 1,15-diol, bound by ether bonds into a macromolecular network. This is different from a previous study of *N. oceanica*, in which the building block of the algaenan was predominantly the C₃₂ 1,17-diol, suggesting different building blocks of the algaenan within Eustigmatophytes. Analysis of the different time points in oxic degradation did not reveal a pronounced impact of oxygen exposure on the chemical structure of the algaenan. This suggests that the algaenan is relatively resistant to oxic degradation on timescales of at least 270 days and well preserved during settling in the water column and subsequent burial in the sediment.

Acknowledgements

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Chapter 4

A search for macromolecular sources of long-chain alkyl diols

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Abstract

C₂₈ to C₃₂ 1,13- and 1,15- long chain diols (LCDs) have been shown to occur in cultured freshwater and marine Eustigmatophytes and are thought to be the building blocks of the insoluble biopolymer algaenan, which forms a constituent of their cell wall. Commonly, LCDs are reported to occur as bound, yet extractable lipids in biomass, which can be released by base hydrolysis of the extract or biomass, while only minor amounts of LCDs occur freely. Here, we attempted to identify potential (macro-)molecular sources of LCDs in extracts of cell material of *Nannochloropsis* sp. by application of ultra-high pressure liquid chromatography combined with high resolution mass spectrometry (UHPLC/HRMS) and semi-preparative HPLC. However, no LCD precursor lipids could be identified by UHPLC-HRMS, while semi-preparative HPLC led to the identification of freely occurring, but not of bound-extractable LCDs, in contrast to previous studies. Subsequent analysis of biomass of three different strains of *Nannochloropsis* spp. showed that LCDs largely occur as free-extractable lipids with only minor increases in LCD concentration after application of acid hydrolysis on the extract. These results therefore call the common hypothesis that LCDs occur as bound-extractable lipids in alga biomass into question.

1. Introduction

Long chain alkyl diols (LCDs) occur in a variety of environments, i.e. freshwater, soil or marine environments. In the latter, they are used to reconstruct sea surface temperatures (Rampen et al., 2012), upwelling and productivity conditions (Rampen et al., 2008; Willmott et al., 2010) and riverine input into the marine environment (de Bar et al., 2016; Lattaud et al., 2017a, 2017b; He et al., 2020). LCDs are composed of long, even carbon-numbered n-alkyl carbon chains with chain lengths of C₂₈ to C₃₆ with alcohol groups at the first carbon and at a varying mid-chain position at C₁₂ to C₁₅ (Versteegh et al., 1997). Saturated and unsaturated 1,14-diols with chain lengths of C₂₈ to C₃₂ have been shown to be produced by *Proboscia* diatoms and have also been identified in *Apedinella radians* (Sinninghe Damsté et al., 2003; Rampen et al., 2007, 2011). C₂₈ to C₃₂ 1,13- and 1,15-diols have been shown to be produced by cultured freshwater and marine Eustigmatophytes, i.e. *Nannochloropsis oculata* and *Nannochloropsis oceanica* (Volkman et al., 1992; Gelin et al., 1997b, 1999; Méjanelle et al., 2003; Shimokawara et al., 2010; Rampen et al., 2014a; Balzano et al., 2017; Zhang and Volkman, 2017), however the distributions of LCDs in marine environments and cultures differ, suggesting unknown marine sources (Volkman et al., 1992; Versteegh et al., 1997; Rampen et al., 2012, 2014a).

The 1,13- and 1,15-diols have been postulated as the building blocks of the insoluble biopolymer algaenan occurring in the cell wall of Eustigmatophytes (de Leeuw et al., 1981; Tegelaar et al., 1989b; Gelin et al., 1997b; Volkman et al., 1998; Scholz et al., 2014). Several studies have characterized the algaenan using chemical degradation methods, e.g. RuO₄ or HI (Blokker et al., 1998b, 2000, 2006), or thermal degradation methods such as pyrolysis GC-MS (Gelin et al., 1997b, 1999). These studies led to the conclusion that the algaenan is mainly composed of ether-bound LCDs (Zeliber et al., 1988; Tegelaar et al., 1989b; de Leeuw and Largeau, 1993; Gelin et al., 1997b; Blokker et al., 1998b, 2000). A more recent characterization by Scholz et al. (2014) has extended the commonly accepted algaenan structure, by identifying carboxyl, aldehyde, ketone and ester moieties in a bi-layer cell wall composed of a cellulose and an algaenan layer. Besides being part of the insoluble algaenan, LCDs also occur in extracts obtained by ultrasonic or Bligh Dyer extraction of algal biomass (Volkman et al., 1992; Gelin et al., 1997b) and as bound lipids which can be released by base and acid hydrolysis of the extract or residue left after extraction (Gelin et al., 1997b; Grossi et al., 2001; Shimokawara et al., 2010; Reiche et al., 2018). Since LCDs are commonly detected in e.g. *Nannochloropsis*

oculata and *Nannochloropsis salina* as bound-extractable lipids and are not detected as free-extractable lipids or only in minor quantities (Volkman et al., 1992; Gelin et al., 1997b), LCDs seem to be bound to an as of yet unknown polar (macro)molecule. Holguin et al. (2013) analyzed the intact polar lipids of different *Nannochloropsis* spp. and also could detect LCDs only after hydrolysis and not in their free-extractable form. Analysis of a *Nannochloropsis salina* enrichment culture using FT-ICR mass spectrometry, suggested that long-chain sulfate lipids as the source of LCDs released by base and acid hydrolysis. Additionally, Shimokawara et al. (2010) identified LCDs in Eustigmatophyte alga as free-extractable lipids and as lipid bound by ester bonds to more polar compounds, e.g. hydroxy fatty acids bound to amino acids and saccharides, or by ether bonds to macromolecules.

However, none of these studies were successful in unambiguously identifying the polar (macro-)molecules that represent the precursors from which LCDs are released by base and acid hydrolysis.

Here, we attempted to identify the polar (macro-)molecular sources of LCDs in extracts of *Nannochloropsis* biomass by application of ultra-high pressure liquid chromatography high resolution mass spectrometry (UHPLC-HRMS) in combination with semi-preparative HPLC. Furthermore, we quantified the amount of free and bound-extractable LCDs in extracts of different *Nannochloropsis* sp. strains.

2. Methods

2.1. Cultivation of alga strains

In this study, three strains of *Nannochloropsis* sp. were analyzed, e.g. *Nannochloropsis oculata* CCMP525 (hereafter *N. oculata* CCMP525), *Nannochloropsis oculata* CCMP2195 (hereafter *N. oculata* CCMP2195) and *Nannochloropsis oceanica* CCMP1779 (hereafter *N. oceanica* CCMP1779). *N. oculata* CCMP525 was originally ordered and grown by Mariculture Inc and kept frozen at -20°C until analysis.

N. oculata CCMP2195 and *N. oceanica* CCMP1779 were cultured according to Balzano et al. (2017). In short, both strains were cultured in f/2 medium (Guillard, 1975) and maintained under a 16: 8 light: dark cycle at 15°C for *N. oculata* CCMP2195 and 20°C for *N. oceanica* CCMP1779. Both strains were grown in 0.25 L volumes within 0.5 L Erlenmeyer flasks until reaching late exponential phase at 10 days as described by Balzano et al. (2017). The number of cells per culture were counted by flow cytometry (Marie et al., 1999) and subsequently the batch culture was harvested by filtration through 0.7 µm GF/F filters (Whatman, Maidstone, UK).

2.2. Lipid analysis

Frozen filters were freeze dried using a LyoQuest (Telstart, Life Sciences) prior to analysis and were subsequently ultrasonically extracted with methanol/ dichloromethane (MeOH/ DCM) (2:1, v/v, 4x). DCM phases were combined and dried using rotary evaporation and subsequent drying under N₂.

An aliquot of the extract was subsequently saponified by refluxing with 1 M KOH in MeOH for 1 h. The DCM phase was separated from the H₂O/ MeOH phase and dried under N₂. Finally, aliquots of both the extract and the saponified extract were acid hydrolyzed by refluxing with 1.5 M HCl in MeOH for 3 h. After cooling, the mixture was neutralized using 2 N KOH in

MeOH and extracted with DCM/ H₂O (1:1, v/v, 3x). The DCM phases were separated from the H₂O/MeOH phase and combined to be dried under N₂.

The different extracts were eluted with DCM over a small pasteur pipette containing Na₂SO₄ to remove remaining salts. A known amount of the internal standard C₂₂ 7,16- diol (MolPort SIA, Latvia) was added to the extracts. Prior to analysis by gas chromatography (GC), extracts were silylated by adding BSTFA [N,O-bis(trimethyl-silyl)trifluoroacetamine] and pyridine and heated up to 60 °C for 20 min. Subsequently, samples were dissolved in ethyl acetate.

Samples were analyzed for LCDs on an Agilent 7890B gas chromatograph equipped with a fused silica capillary column (25 m x 0.32 mm) coated with CP Sil-5 (film thickness 0.12 µm). Samples were injected at 70 °C oven temperature, increasing to 130 °C at 20 °C/min and then 320 °C at 4°/min which was held for 25 min with a helium flow of 2 mL/min. LCDs were quantified by integration of respective peak areas of LCDs and the internal standard.

4

2.3. Ultra-High Pressure Liquid Chromatography-Electro Spray Ionization/High Resolution Mass Spectrometry (UHPLC-ESI/HRMS) analysis

An aliquot of extract obtained from *N. oculata* CCMP525 biomass was analyzed by normal phase high performance liquid chromatography on an Ultimate 3000 RS UHPLC, equipped with thermostatted auto-injector and column oven, coupled to a Q Exactive Orbitrap MS with Ion Max source with heated electrospray ionization (HESI) probe (Thermo Fisher Scientific, Waltham, MA). Separation was achieved on a YMC-Triart Diol-HILIC column (250 x 2.0 mm, 1.9 µm particles, pore size 12 nm; YMC Co., Ltd, Kyoto, Japan) maintained at 30 °C. The following elution program was used with a flow rate of 0.2 mL min⁻¹: 100% A for 5 min, followed by a linear gradient to 66% A: 34% B in 20 min, maintained for 15 min, followed by a linear gradient to 40% A: 60% B in 15 min, followed by a linear gradient to 30%A:70%B in 10 min, where A = hexane/2-propanol (80:20 [volume in volume, v/v]) and B = 2-propanol/water (90:10 [v/v]). Total run time was extended to 100 min to allow detection of potentially late eluting lipids. HESI settings were as follows: T 50 °C, sheath gas (N₂) pressure 35 (arbitrary units), auxiliary gas (N₂) pressure 10 (arbitrary units), sweep gas (N₂) pressure 5 (arbitrary units), spray voltage 4.0 kV for positive ion ESI and -5,5 kV for negative ion ESI, capillary temperature 275 °C, S-Lens 70 V. Lipids were analyzed in both positive and negative ion mode with a mass range of *m/z* 375 to 2000 and *m/z* 1000-4000 (resolution 70,000), followed by data dependent MS² (resolution 17,500), in which the ten most abundant masses in the mass spectrum (with the exclusion of isotope peaks) were fragmented successively (stepped normalized collision energy 15, 22.5, 30; isolation window 1.0 *m/z*). A dynamic exclusion window, with a mass tolerance of 3 ppm, of 6 sec was applied.

2.4. Normal and reverse phase semi-preparative high-performance liquid chromatography (HPLC)

In an attempt to isolate macromolecular sources of LCDs, normal phase semi-preparative HPLC following the method described by Bauersachs et al. (2009), and reverse phase semi-preparative HPLC following the method described by Zhu et al. (2013) and Wörmer et al. (2013) , was applied to an extract of *N. oculata* biomass CCMP525.

In short, for normal phase semi-preparative HPLC using a 1260 Infinity Binary HPLC (Agilent, San Jose, CA), a filtered ultrasonically extracted *N. oculata* CCMP525 biomass extract was injected onto a LiChrospher DIOL column (250mm x 10 mm, 5µm, Alltech Deerfield, IL, USA) following the gradients and mobile phase described in 2.3, at a flow rate of 3 mL min⁻¹.

For reverse phase semi-preparative HPLC, the same *N. oculata* CCMP525 extract was injected onto a C18 column (SymmetryPrep C18, 150 mm x 7.8 mm, 5 μ m, Waters Corporation, USA). Lipids were eluted at 3 mL min⁻¹ using a linear gradient: 100% eluent A for 2 min, increase to 15% eluent B at 2.10 min, 43,5 % eluent B at 15 min and holding to 16 min, washing with 100% eluent B from 16 min to 24 min before returning to 100% eluent A for 5 min. Eluent A was methanol/ water (85:15, v/v) and eluent B was methanol/ 2-propanol (50:50, v/v).

For both normal and reverse phase semi-preparative HPLC, eluents were collected in 1 min fractions. For initial screening purposes, aliquots of the 1 min fractions were combined into 10 min fractions. Aliquots of the 10 min fractions were subsequently saponified according to the procedure described above. The C₂₂ 7,16- diol internal standard was added to both the HPLC fractions as well as the saponified fractions, which were subsequently silylated and analyzed using gas chromatography as described above.

3. Results and discussion

3.1. UHPLC-ESI/HRMS analysis of an extract of *N. oculata* CCMP525

In a first attempt to identify sources of bound LCDs in extracts of Eustigmatophyte alga, we analyzed an extract of *N. oculata* CCMP525 biomass by UHPLC-ESI/HRMS. With ESI in positive mode, phosphatidylcholines, diacylglyceryl-trimethylhomoserine (DGTS), diacylglyceryl-hydroxymethyltrimethylalanine (DGTA), dihexoses, sulfoquinovosyl-diacylglycerol (SQDG), inositolphosphoceramide (IPC) as well as lyso forms of these intact polar lipids (IPLs) were detected (Fig. 1A). With ESI in negative mode, SQDGs and lyso SQDGs as well as C₃₂ sulfate lipids, i.e. with elemental formula C₃₂H₆₃O₄S (based on accurate mass) and a C₃₂ hydroxyl sulfate lipids, i.e. C₃₂H₆₅O₅S, were identified (Fig. 1B). Common esterified fatty acid moieties of the identified IPLs included saturated and unsaturated C₁₆ and C₁₈ fatty acids and C_{20:5} polyunsaturated fatty acids. This composition agrees with previous studies of IPLs of *Nannochloropsis* species (He et al., 2011; Holguin and Schaub, 2013). In order to find polar lipids potentially containing LCDs, we analyzed the MS² data in search for diagnostic mass fragments or mass losses related to LCDs, e.g. *m/z* 18 or *m/z* 57, 71, 85 indicating the loss of a hydroxyl group or alkyl chains, respectively. However, this did not lead to any tentative identification of lipids containing LCD moieties. While C₃₂ (hydroxy) sulfate lipids were identified, in agreement with earlier findings of Holguin and Schaub (2013), we could not replicate the hypothesis that sulfate lipids serve as a source of LCDs upon release by hydrolysis.

3.2. Semi-preparative HPLC of extract of *N. oculata* CCMP525

The large abundances of IPLs in the biomass extract dominated the base peak chromatograms and, in combination with the unknown nature (and hence unknown mass spectra) of the precursors of the bound LCDs, severely complicated their identification. Therefore, we took an alternative approach. An extract of *N. oculata* CCMP525 obtained by ultrasonication was subjected to semi-preparative HPLC and the eluent was collected in 10 min fractions which were analyzed directly by GC as well as saponified and analyzed for LCDs in order to identify the HPLC retention time of free and bound LCDs, respectively.

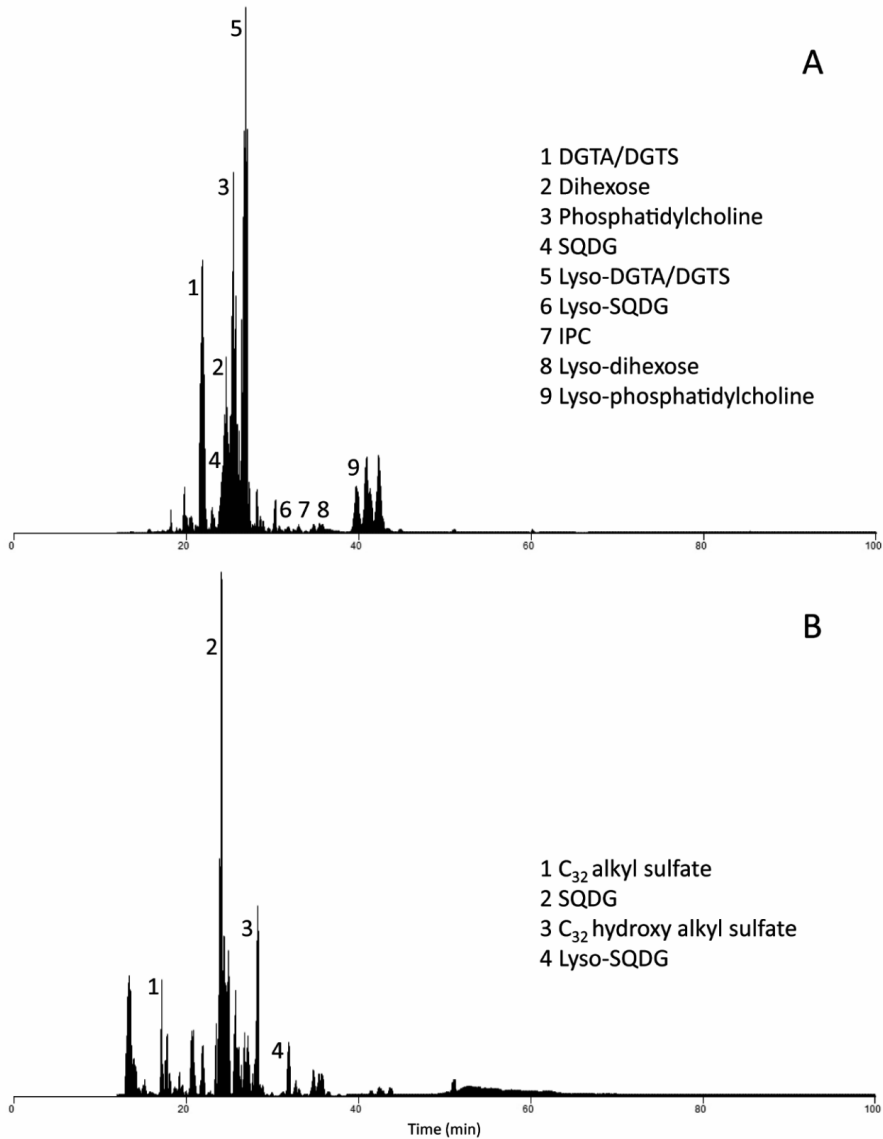


Fig. 1: UHPLC base peak chromatograms of *N. oculata* CCMP525 biomass with identified IPLs in A: positive ion mode and B: negative ion mode. Identified IPLs include among others diacylglyceryl-trimethylhomoserine (DGTS), diacylglyceryl-hydroxymethyltrimethylalanine (DGTA), sulfoquinovosyldiacylglycerol (SQDG) and inositolphosphoceramide (IPC).

We first attempted normal phase semi preparative HPLC (Bauersachs et al., 2009) which resulted in the detection of freely occurring LCDs in the first 10 min fraction, with summed up LCD concentrations of 0.98 $\mu\text{g/g}$ dry weight biomass (Fig. 2A). Very small abundances of freely occurring LCDs were also detected in the second to fourth 10 min fractions. Surprisingly,

saponification of all collected fractions (in the time window 0-70 min) did not lead to the detection of any additional LCDs suggesting that extractable, polar macromolecular sources of bound LCDs did not elute from the column under normal phase HPLC conditions. Therefore, we repeated the experiment using the reversed phase semi-preparative HPLC methods described by Wörmer et al. (2013) and Zhu et al. (2013) as any compound that elutes very late or not at all using normal phase liquid chromatography should elute early when using a reverse phase column. Applying reverse phase semi-preparative HPLC resulted again in the detection of freely occurring LCDs, but now in the second fraction (20-30 min), with summed up LCD concentrations of 0.67 $\mu\text{g/g}$ dry weight biomass (Fig. 2B). Small amounts of LCDs were also detected in the third fraction (30-40 min). Again, however, application of saponification on the collected 7 fractions did not lead to the detection of bound LCDs.

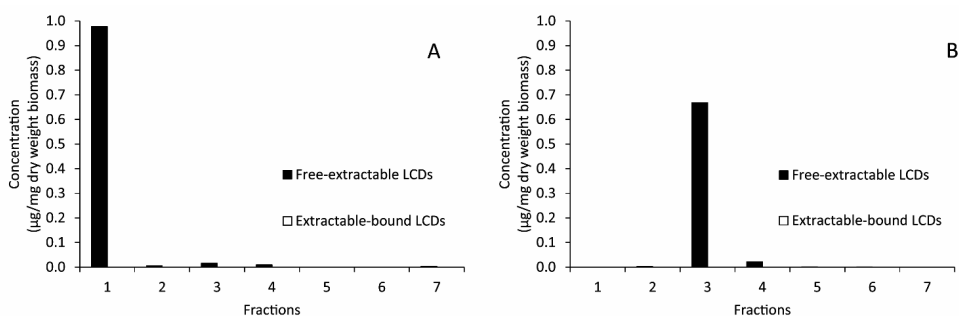


Fig. 2: Yields of free- and bound extractable LCDs after semi-preparative HPLC. A: yields of summed up LCD concentrations following normal phase semi-preparative HPLC. B: yields of summed up LCD concentrations following reverse phase semi-preparative HPLC. Fractions 1-7 represent 10 min intervals of eluent collection.

3.3. Occurrence of LCDs in *Nannochloropsis* species

The relatively high abundances of freely occurring LCDs and the lack of bound LCDs released by base hydrolysis in the fractions obtained by both normal and reverse phase semi-preparative HPLC are in contrast to the common hypothesis that LCDs occur as bound-extractable LCDs and that relatively minor amounts of freely occurring LCDs can be detected (Volkman et al., 1992). However, only few studies have studied free-extractable LCDs, instead analyzing LCDs in biomass cultures by either immediate base hydrolysis of the extract (Gelin et al., 1997b; Rampen et al., 2007) or of the biomass directly (Balzano et al., 2017). Therefore, we re-examined the mode of occurrence of LCDs in extracts by analyzing three different strains of *Nannochloropsis* species, e.g. *N. oculata* CCMP2195, *N. oceanica* CCMP1779 and *N. oculata* CCMP525. The different strains were ultrasonically extracted and subsequently subjected to base hydrolysis, acid hydrolysis and consecutive base and acid hydrolysis following the methodological approaches presented by Volkman et al. (1992), Gelin et al. (1997) and Reiche et al. (2018).

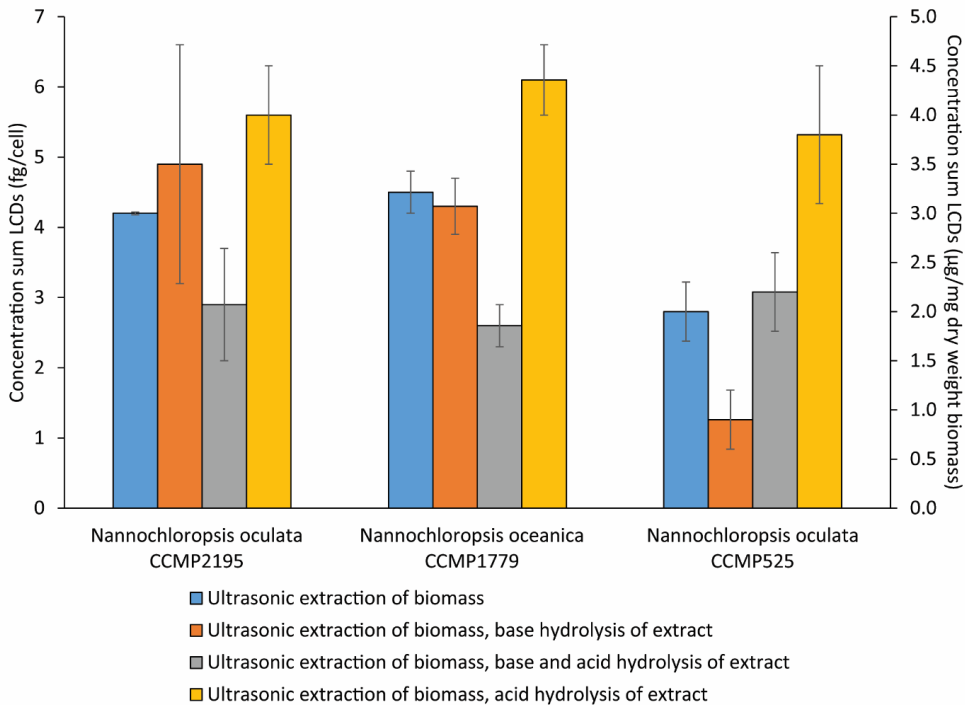


Fig. 3: Yields of summed LCDs in three different *Nannochloropsis* strains following the application of ultrasonic extraction and subsequent base hydrolysis of the extract, base and acid hydrolysis of the extract and acid hydrolysis of the extracts. Yields for *N. oculata* CCMP2195 and *N. oceanica* CCMP1779 given in fg/ cell. Yields for *N. oculata* CCMP525 given in µg/ mg dry weight biomass. Standard deviations calculated from duplicate work-ups.

Both *N. oculata* CCMP2195 and *N. oceanica* CCMP1779 showed similar concentrations of freely occurring summed LCDs with 4.2 ± 0.02 and 4.5 ± 0.3 fg/ cell respectively (Fig. 3), with *N. oculata* CCMP525 showing concentrations of 2.0 ± 0.3 µg/ mg dry weight biomass. The LCDs in base hydrolyzed extracts, which represent both freely extractable and LCDs released by base hydrolysis, showed a slight increase in concentration to 4.9 ± 1.7 fg/ cell for *N. oculata* CCMP2195 and a decrease to 4.3 ± 0.4 fg/ cell for *N. oceanica* CCMP1779. *N. oculata* CCMP525 showed a larger decrease in concentration to 0.9 ± 0.3 µg/mg dry weight biomass. Subsequent acid hydrolysis of the base hydrolyzed extract, thus representing freely extractable LCDs and LCDs released by base and acid hydrolysis, showed decreasing concentrations for summed LCDs for *N. oculata* CCMP2195 and *N. oceanica* CCMP1779 to 2.9 ± 0.8 fg/ cell and 2.6 ± 0.3 fg/ cell respectively. In contrast, the concentration of summed LCDs for *N. oculata* CCMP525 increased to 2.2 ± 0.4 µg/mg dry weight biomass. Finally, when direct acid hydrolysis to the extract was applied, thus representing freely extractable LCDs and LCDs released by acid hydrolysis, all three strains showed an increase in LCD concentration compared to the freely occurring LCDs, increasing to 5.6 ± 0.7 fg/ cell for *N. oculata* CCMP2195, 6.1 ± 0.5 fg/ cell for *N. oceanica* CCMP1779 and 3.8 ± 0.7 µg/mg dry weight biomass for *N. oculata* CCMP525.

Our results clearly show that applying base hydrolysis does not lead to a substantial increase in concentration due to a release of bound LCDs, but instead often led to a decrease in concentration. In principle, an extract treated by base hydrolysis should at least contain the same amount of LCDs as detected directly in the extract. This decrease might be explained by losses through artifact formation as also described in the context of algaenan isolation (Allard et al., 1997, 1998). Harsh consecutive hydrolyses in alkaline or acidic conditions may form melanoidin-like polymers due to the recondensation of polar molecules present in the cell material and our work-up, albeit of less harsh nature, may have led to the formation of similar artifacts binding LCDs to other compounds. Alternatively, work up losses during the hydrolysis procedures, e.g. from shaking against water and subsequent dissolution or micelle formation, may have led to a decrease. Furthermore, weak hydrolysis might be induced when the solvents used for ultrasonic extraction are stored over longer timescales, e.g. leading to the generation of phytadienes from chlorophylls. However, if that would be the case, common work-up products by base and acid hydrolyses of biomass (Grossi et al., 1996) would be found as freely extractable lipids in our *Nannochloropsis* spp biomass, which was not the case. Applying a straight acid hydrolysis on the extract did result in a significant increase in LCD concentration in all three strains and therefore to a release of bound LCDs from unknown sources. Since base, in contrast to acid, hydrolysis did not release any LCDs, these LCDs might have been ether-bound to a glycosidic moiety or amide bound (Goossens et al., 1986, 1989a). Nevertheless, the amount of LCDs released after acid hydrolysis is still only around 30-40% of freely occurring LCDs and so the large majority of LCDs occur as free- extractable lipids in Eustigmatophyte algae, contrasting previous studies. Thus, studies analyzing LCD-containing biomass and not taking free-extractable LCDs into account, might have overlooked a large pool of lipids leading to potential misinterpretations, i.e. creating a bias in distribution and an underestimation of LCD occurrences.

4. Conclusions

In an attempt to identify and characterize macromolecular sources of LCDs, LCD-containing biomass was analyzed by various HPLC methods. Identification of potential polar macromolecules containing LCDs by UHPLC-ESI/HRMS proved to be unsuccessful. Subsequently, an isolation of macromolecular sources by semi-preparative HPLC was attempted. However, both in normal and reverse phase semi-preparative HPLC no ester-bound LCDs and only free-extractable LCDs were detected. Analysis of three strains of *Nannochloropsis* species showed high abundances of free-extractable LCDs and lower amounts of ester-bound LCDs, partially explaining the unsuccessful attempt to isolate macromolecular sources by HPLC methods. Our results thus call into question the current hypothesis that extractable LCDs occur mainly as bound lipids in extracts of biomass of Eustigmatophyte alga.

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Chapter 5

The impact of mode of occurrence of long chain diols and keto-ols and related indices in an S5 sapropel from the Eastern Mediterranean

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Abstract

Long chain diols (LCDs), mid-chain keto-ols (keto-ols) and associated indices are increasingly used as proxies in paleoenvironmental reconstructions, i.e. to determine sea surface temperatures, upwelling conditions, riverine inputs into marine environments and post-depositional oxidation. However, few studies consider that LCDs have different modes of occurrence, i.e. they occur not only as free-extractable compounds but also bound to organic matter by ester- or glycosidic ether- or amide-bonds which may result in an underestimation of LCD abundances and possible differences in LCD indices in the sedimentary record. Here, we studied the effect of matrix sequestration on LCDs, keto-ols and associated indices in a well-defined S5 sapropel from the Eastern Mediterranean by analyzing freely occurring extractable lipids but also LCDs and keto-ols released by base and acid hydrolysis of the extract as well as the residue after sediment extraction. Our results show that the concentrations of 1,14-diols are higher after base and acid hydrolysis, indicating that some LCDs are bound to organic matter and the sedimentary matrix. While freely occurring LCDs dominate the sapropel, LCDs are only detected as residually ester-bound and residually glycosidic ether- or amide-bound lipids above or below the sapropel. However, LCD indices in the S5 sapropel are quite similar to each other irrespective of the mode of occurrence, suggesting that matrix sequestration does not have a large impact on the application of LCD indices in paleoenvironmental reconstructions. Finally, keto-ols were only identified in the sapropel and their relative amount, quantified using the DOXI index, did not show significant trends throughout the sapropel.

1. Introduction

In order to reconstruct past environmental conditions, several organic proxies have been established. Among them, long chain diols (LCDs) have been described in a large number of marine and freshwater environments (Versteegh et al., 1997; Sinninghe Damsté et al., 2003; Rampen et al., 2007, 2008, 2014a; 2014b; Zhang et al., 2011; Romero-Viana et al., 2012; de Bar et al., 2016; Lattaud et al., 2017a) and are increasingly used in paleo-environmental reconstructions. LCDs are commonly composed of a long n-alkyl chain of C₂₈ to C₃₂ carbon atoms with alcohol groups at the first carbon atom and at a mid-chain position, commonly at position C₁₂ to C₁₅ (Versteegh et al., 1997) and were first described in Black Sea sediments (de Leeuw et al., 1981). C₂₈ and C₃₀ 1,14- saturated and unsaturated diols have been shown to be produced by *Proboscia* diatoms (Sinninghe Damsté et al., 2003; Rampen et al., 2007) while C₂₈ to C₃₂ 1,14-diols (Rampen et al., 2011) have also been identified in *Apedinella radians*. C₂₈ to C₃₂ 1,13- and 1,15-diols are produced by Eustigmatophyte algae (Volkman et al., 1992; Gelin et al., 1997b, 1999; Rampen et al., 2012, 2014a; Balzano et al., 2017; Zhang and Volkman, 2017), although LCD distributions produced by these algae are dissimilar to those in the marine environment (Volkman et al., 1992; Versteegh et al., 1997; Rampen et al., 2012, 2014a). LCDs produced by Eustigmatophytes have been shown to occur as part of the insoluble, non-hydrolysable biopolymer algaenan (de Leeuw et al., 1981; Tegelaar et al., 1989b; Gelin et al., 1997b; Volkman et al., 1998; Scholz et al., 2014; Zhang and Volkman, 2017).

Several indices based on LCD distribution have been proposed for use in paleo-environmental reconstructions and have since been applied in a variety of marine environments (Pancost et al., 2009; Willmott et al., 2010; Rampen et al., 2012, 2014b; Seki et al., 2012; Lopes dos Santos et al., 2013; Rodrigo-Gámiz et al., 2014, 2015, 2016; Nieto-Moreno et al., 2015; Plancq et al., 2015). The Long chain Diol Index (LDI) uses the fractional abundances of the C₂₈ 1,13-, C₃₀ 1,13- and C₃₀ 1,15-diols (Rampen et al., 2012) to determine past sea surface temperatures (SST). The Diol Index I and Diol Index II reconstruct upwelling intensities/ high nutrient conditions by use of the abundances of C₂₈ and C₃₀ 1,14-diols relative to C₂₈ and C₃₀ 1,13-diols

(Willmott et al., 2010) or to C₃₀ 1,15-diols (Rampen et al., 2008), respectively. More recently, the fractional abundance of the C₃₂ 1,15-diol (FC₃₂ 1,15-diol) has been proposed as a proxy for the reconstruction of riverine input into the marine environment (Lattaud et al., 2017b, 2017a; He et al., 2020). Additionally, the Diol Oxidation Index (DOXI) has been proposed to trace the occurrence of post-depositional oxidation (Ferreira et al., 2001). It is based on the ratio of LCDs and structurally similar mid-chain keto-ols (keto-ols) of identical chain length and mid-chain position of the functional group. Keto-ols are lipids occurring in a wide range of environments (Versteegh et al., 1997, 2000; Xu et al., 2007; Bogus et al., 2012) and because of their similar chemical structure and more oxidized state, have been hypothesized to represent oxidative products of LCDs (Ferreira et al., 2001; Sinninghe Damsté et al., 2003). Indeed, they have only been identified in trace amounts in one alga, i.e. *Nannochloropsis gaditana* (Méjanelle et al., 2003). However, differences in carbon chain lengths and position of functional moieties of LCDs and keto-ols have cast doubt on this hypothesis (Versteegh, 1997, 2019; Méjanelle et al., 2003).

Most studies using LCDs for paleoclimate reconstructions use a straightforward extraction procedure such as accelerated solvent extraction (ASE) or ultrasonic extraction. Consequently, they only take freely occurring LCDs into account. However, as shown by various studies (Ten Haven et al., 1987a; Ahmed et al., 2000; Grossi et al., 2001; Shimokawara et al., 2010), LCDs can occur not only as free lipids, but also bound in the extract and in the residue left after extraction. Typically, these LCDs are released by application of base and acid hydrolysis of the extract or residue (Volkman et al., 1992; Gelin et al., 1997b; Reiche et al., 2018). Therefore, most studies may have underestimated the amount of LCDs present in sediments, by not considering LCDs that are bound to organic matter by ester and glycosidic ether or amide bonds. Only few studies have examined LCDs bound in sediments. Ahmed et al. (2000) analyzed bound lipids from sediments of the Monterrey Formation, identifying C₃₀- and C₃₂ 1,15-diols and keto-ols as ester-bound lipids. Shimokawara et al. (2010) analyzed LCDs in a sediment core from Lake Baikal, focusing on how LCDs are bound to other lipids. Freely occurring LCDs accounted for 70-100 % of total LCDs throughout the core while LCDs that were ether linked to long chain compounds accounted for 0-30 %. Additionally, small amounts of ester-bound LCDs linked to more polar compounds, e.g. hydroxy fatty acids, were detected. The C₃₂ 1,15-diol was the most abundant freely occurring LCD, while the C₃₀ 1,13-diol was the most abundant ester- and ether-bound LCD. Ten Haven et al. (1987a) analyzed LCDs and keto-ols in sediments of an S7 sapropel, finding free-extractable and extractable-bound C₃₀ and C₃₂ 1,15-diols as well as C₃₀ 1,15-keto-ols.

Here, we studied the impact of mode of occurrence on LCDs and keto-ols in an S5 sapropel from the Eastern Mediterranean by analyzing not only freely occurring lipids, but also LCDs and keto-ols released by base and acid hydrolysis from the extract and extracted sediment residue. We use these results to evaluate the impact of the mode of occurrence of LCDs on the different proxies, i.e. the LDI, Diol Index I, FC₃₂ 1,15-diol and DOXI.

2. Material and methods

2.1. Sampling

The piston core studied here was collected during the NESSC 64PE406 cruise on board of R/V Pelagia at station 1 (33° 18.14898' N, 33° 23.71998' E, 1760 m water depth) in January 2016 (cf. Bale et al., 2019). The core, 9.2 m in length, was split into several sections, each measuring 1 m, which were subsequently analyzed for its bulk inorganic geochemistry using X-Ray Fluorescence (XRF) and inductively coupled plasma mass spectrometry (ICP-MS) as described in Bale et al. (2019). Section 6 was selected for this study, displaying a well preserved S5

sapropel lithostratigraphy in a core depth of 371 – 337 cm. Due to the focus on the S5 sapropel, 30 samples over a depth range of 49 cm were chosen with a resolution of 1 cm around the visually identified upper and lower boundaries of the sapropel (11 samples) and a resolution of 2 – 3 cm within the sapropel (19 samples). Samples were subsequently freeze dried.

2.2. Lipid analysis

Lipids were analyzed following a modified approach described by Gelin et al. (1997) and Goossens et al. (1986) (Fig. 1). Freeze-dried sediments were extracted by modified Bligh-Dyer extraction (Pitcher et al., 2011; Bale et al., 2015). After addition of a known volume of methanol (MeOH): dichloromethane (DCM): phosphate buffer (2:1:0.8; v,v,v), sediments were ultrasonically extracted for 10 min. The solvent mixture was separated from the residues by centrifugation, collected into a round bottom flask and the process was repeated three times. Subsequently, DCM and phosphate buffer were added to the collected solvents to induce phase separation, producing a new ratio of MeOH: DCM: phosphate buffer (1:1:0.9; v,v,v). After centrifugation, the DCM phase was collected into a round bottom flask and the remaining MeOH: phosphate buffer mixture was washed three times using DCM. The collected DCM phase and the residue were dried using rotary evaporation and drying under N₂. The extract produced by this work-up step was labelled as the free-extractable fraction.

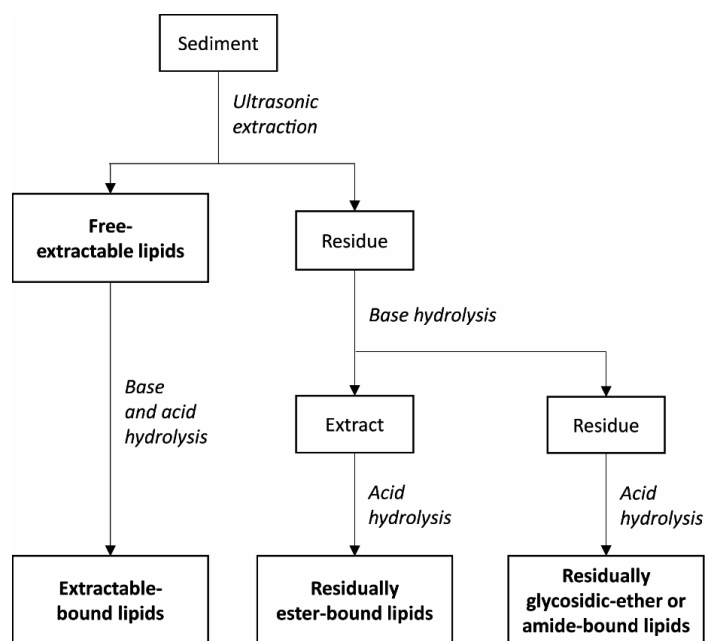


Fig. 1: Analytical scheme representing the work up for the analysis of free and bound LCDs in S5 sapropel sediments, as described in Section 2.2.

An aliquot of the Bligh-Dyer extract was subsequently base and acid hydrolyzed. In the first step of this procedure, the Bligh-Dyer extract was base hydrolyzed by refluxing with 1 M KOH in MeOH. After cooling, the extract was acidified to a pH of 3 using 2 N HCl in MeOH and furthermore extracted with DCM/ H₂O (1:1; v,v: 3x). The DCM phase was separated from the H₂O/ MeOH phase, subsequently combined and dried under N₂. In the second step, the base hydrolyzed extract was acid hydrolyzed using 1,5 M HCl in MeOH. After cooling, the extract was neutralized using 2 N KOH in MeOH and extracted with DCM/H₂O (1:1 v,v, 3x). The DCM phases were separated from the H₂O/ MeOH phase, the DCM phase was combined and dried under N₂. The extract resulting from the base and acid hydrolysis of the Bligh-Dyer extract was labelled as the extractable-bound and contained freely occurring as well as bound lipids present in the extract (Fig. 1).

The sediment residue left after the Bligh-Dyer extraction was also base and acid hydrolyzed. After base hydrolysis of the residue using 1 M KOH in MeOH, solvents were collected into a separatory funnel and acidified to a pH of 3 using 2 N HCl in MeOH. Residues were subsequently extracted using H₂O/ MeOH (1:1 v,v), MeOH (2x) and DCM (3x). The DCM layer was separated from the MeOH/ H₂O layer and collected, the MeOH/ H₂O layer was extracted with DCM (3x) and the combined DCM phases were dried under N₂. The extract resulting from the base hydrolysis of the residues was subsequently acid hydrolyzed following the procedure as outlined above. The extract resulting from the base hydrolysis of the residue and the subsequent acid hydrolysis of the resulting base hydrolyzed extract, was labelled as the residually ester-bound fraction, i.e. containing lipids bound via an ester bond to the organic matrix t (Fig. 1).

In a last step, the sediment residue resulting from the base hydrolysis of the Bligh-Dyer was acid hydrolyzed by refluxing with 1.5 M HCl in MeOH. After cooling, solvents were collected into a separatory funnel and neutralized using 2 N KOH in MeOH. Residues were subsequently extracted using H₂O/ MeOH (1:1 v,v), MeOH (2x) and DCM (3x) and solvents were collected into the separatory funnel. The DCM layer was separated from the MeOH/ H₂O layer and collected, the MeOH/ H₂O layer was extracted with DCM (3x) and the combined DCM phases were dried under N₂. The extract resulting from this last step was labelled as the residually glycosidic ether- or amide-bound fraction (Fig. 1).

All resulting extracts were eluted with DCM over a small Pasteur pipette containing Na₂SO₄ to remove remaining salts. A known amount of the C₂₂ 7,16- diol internal standard was added to aliquots which were subsequently methylated with diazomethane in diethylether. Very polar components were removed by eluting in ethyl acetate over a small pasteur pipette containing silica gel.

Prior to analysis by GC and GC/MS, extracts were silylated by adding BSTFA [N,O-bis(trimethyl-silyl)trifluoroacetamine] and pyridine and heated up to 60 °C for 20 min. Subsequently, samples were dissolved in ethyl acetate. Samples were analyzed for LCDs and keto-ols on an Agilent 7890B gas chromatograph coupled to an Agilent 5977A mass spectrometer, equipped with a fused silica capillary column (25 m x 0.32 mm) coated with CP Sil-5 (film thickness 0.12 µm) with a helium flow of 2mL/ min. Samples were injected at 70 °C oven temperature, increasing to 130 °C at 20 °C/min and then 320 °C at 4°/min which was held for 25 min. The mass spectrometer operated with an ionization energy of 70 eV. LCDs and keto-ols were quantified in selective ion monitoring (SIM) of *m/z* 187, 313, 339 and 341 for the different LCDs and *m/z* 300, 314 and 328 characteristic for the different keto-ols (Versteegh et al., 1997; Rodrigo-Gámiz et al., 2015, 2016).

Concentrations of LCDs and keto-ols were normalized on the total organic carbon (TOC) content, which has been determined by Bale et al. (2019). Residually ester-bound and residually

glycosidic ether- or amide-bound lipids were analyzed in a lower resolution (12 out of 30 samples) than free-extractable and free- and extractable-bound lipids.

The LDI was calculated and converted to SST as follows (Rampen et al., 2012):

$$LDI = \frac{[C_{30}1,15\text{-diol}]}{[C_{28}1,13\text{-diol}] + [C_{30}1,13\text{-diol}] + [C_{30}1,15\text{-diol}]} \quad [1]$$

$$SST = \frac{LDI - 0.095}{0.033} \quad [2]$$

The Index I was calculated as follows (Rampen et al., 2008):

$$\text{Diol Index I} = \frac{[C_{28}1,14\text{-diol}] + [C_{30}1,14\text{-diol}]}{[C_{28}1,14\text{-diol}] + [C_{30}1,14\text{-diol}] + [C_{30}1,15\text{-diol}]} \quad [3]$$

The FC₃₂ 1,15-diol was calculated as follows (Lattaud et al., 2017b):

$$FC_{32}1,15\text{-diol} = \frac{[C_{32}1,15\text{-diol}]}{[C_{28}1,13\text{-diol}] + [C_{30}1,13\text{-diol}] + [C_{30}1,15\text{-diol}] + [C_{32}1,15\text{-diol}]} \times 100 \quad [4]$$

The DOXI was calculated for LCDs and keto-ols showing identical carbon lengths and mid-chain functional groups (Ferreira et al., 2001):

$$DOXI = \frac{[\text{keto-ol}]}{[\text{keto-ol}] + [\text{diol}]} \quad [5]$$

3. Results

The boundaries of the S5 sapropel were based on the Ba/Al increase and decrease associated to the sapropel initiation and termination, respectively, following Bale et al. (2019). An influence of post-depositional oxidation, the oxidation of organic matter after deposition by a downward moving oxygen front (de Lange et al., 1989; Higgs et al., 1994; Thomson et al., 1995), could not be distinguished in the present S5 sapropel. Comparison of the TOC and Ba/Al profiles shows this absence of post-depositional diagenesis, as Ba/Al shows a similar profile to TOC while being resistant to post-depositional oxidation (Bale et al., 2019).

3.1. LCD and keto-ol concentration profiles

In order to assess the impact of mode of occurrence on the different LCDs and keto-ols, the sediment samples were worked up in a comprehensive approach following the methods described by Gelin et al. (1997) and Goossens et al. (1986) (Fig. 1). Ultrasonic extraction of the sediments resulted in free-extractable lipids, while a subsequent base and acid hydrolysis of the ultrasonic extract represented extractable-bound lipids. Residually ester-bound lipids were obtained after base hydrolysis of the ultrasonically extracted sediment, followed by acid hydrolysis of the base hydrolyzed extract. The residue left after base hydrolysis of the ultrasonically extracted sediment was finally acid hydrolyzed and yielded the residually glycosidic ether or amide-bound lipids. Since extractable-bound lipids represent both freely occurring lipids as well as lipids released by base and acid hydrolysis, extractable-bound LCDs and keto-ols were corrected for free-extractable lipids. However, correction of extractable-bound keto-ols resulted in a series of negative numbers and these were therefore excluded from our discussion. While work-up losses or a potential artifact formation by condensation reactions in alkaline or acidic conditions (Allard et al., 1997, 1998), have been discussed previously to account for minor losses during consecutive hydrolyses (Reiche et al., 2018), it is unclear, why extractable-bound keto-ols were as labile during work-up here.

Several LCDs were identified in the different extract fractions, i.e. the C₂₈ 1,12-, C₂₈ 1,13-, C₂₈ 1,14-, C_{28:1} 1,14-, C₃₀ 1,13-, C₃₀ 1,14-, C₃₀ 1,15- and C₃₂ 1,15-diols. The C₃₀ 1,15-diol was present in highest concentrations, followed by the C₃₂ 1,15-diol, while other LCDs were generally in low abundance. Of the two 1,14-diols identified, the C₃₀ 1,14-diol was generally the most abundant, while the C_{28:1} 1,14-diol was only present in trace amounts.

The free-extractable C₃₀ 1,15-diol (Fig. 2B) was present in low abundance below and above the S5 sapropel with concentrations of 0 to 4 µg/g TOC, while concentrations increased to maximum concentrations of 300 µg/g TOC at 345 cm core depth inside the sapropel. The extractable-bound C₃₀ 1,15-diol was present in lower concentrations of 6 to 64 µg/g TOC, except for a high value of 340 µg/g TOC at 345 cm core depth (Fig. 2B). The residually ester-bound (40 µg/g TOC at 360 cm core depth) and residually glycosidic ether- or amide-bound C₃₀ 1,15-diol (39 µg/g TOC at 358 cm core depth) showed lowest concentrations (Fig. 2B). The free-extractable C₃₂ 1,15-diol was present with concentrations of 0 to 2 µg/g TOC below and above the sapropel and inside the sapropel with maximum concentrations up to 30 µg/g TOC at 363 cm core depth (Fig. 2C), which represents an earlier concentration maximum in contrast to the C₃₀ 1,15-diol. Concentrations of the extractable-bound C₃₂ 1,15-diols were slightly lower with maximum values of 20 µg/g TOC at 345 cm core depth (Fig. 2C). The concentration for residually ester-bound and residually glycosidic ether or amide-bound C₃₂ 1,15-diol showed similar profiles but were lower in concentration than the free-extractable LCDs, i.e. maximizing at 15 µg/g TOC at 360 cm core depth and 12 µg/g TOC at 358 cm core depth respectively (Fig. 2C). The sum of 1,14-diols showed maximum concentrations up to 30 µg/g TOC at 338 cm core depth as free-extractable LCDs. These LCDs could not be detected in sediments below the sapropel and only in trace amounts above the sapropel. The maximum concentration was higher up in the sapropel than that of the 1,15-diols (Fig. 2A). Extractable-bound 1,14-diols were in higher maximum abundance (i.e. 50 µg/g TOC at 345 cm core depth) than free-extractable 1,14-diols (Fig. 2A). In comparison, residually ester-bound and residually glycosidic ether- or amide-bound 1,14-diols were in much lower abundances with maximum concentration at 6 µg/g TOC at 337 cm core depth and 2 µg/g TOC at 342 cm core depth, respectively (Fig. 2A).

The C₃₀ and C₃₂ 1,15-keto-ols are the two major keto-ols identified in the S5 sapropel. Both keto-ols were only detected in the sapropel and not in the sediments below or above it. Additionally, the keto-ols were only identified from 366 cm core depth onwards, therefore showing a slower onset in increasing abundances than the corresponding LCDs. The free-extractable C₃₀ 1,15-keto-ol was present in highest concentrations with maximum concentrations of 210 µg/g TOC at 353 cm core depth, whereas the free-extractable C₃₂ 1,15-keto-ol had maximum concentrations of 60 µg/g TOC at 353 cm core depth (Fig. 2D, E). Both keto-ols showed increasing concentrations towards a mid-sapropel maximum at 353 cm depth and decreasing concentrations thereafter. Concentrations for residually ester-bound C₃₀ 1,15- and C₃₂ 1,15-keto-ols were substantially lower at ca. 20 µg/g TOC at 360 cm core depth. Similarly, residually glycosidic ether- or amide bound C₃₀ 1,15- and C₃₂ 1,15-keto-ols reached similar concentrations at 358 cm core depth (Figs. 2D, E). As indicated above, the extractable-bound C₃₀ and C₃₂ 1,15-keto-ols were excluded from discussion.

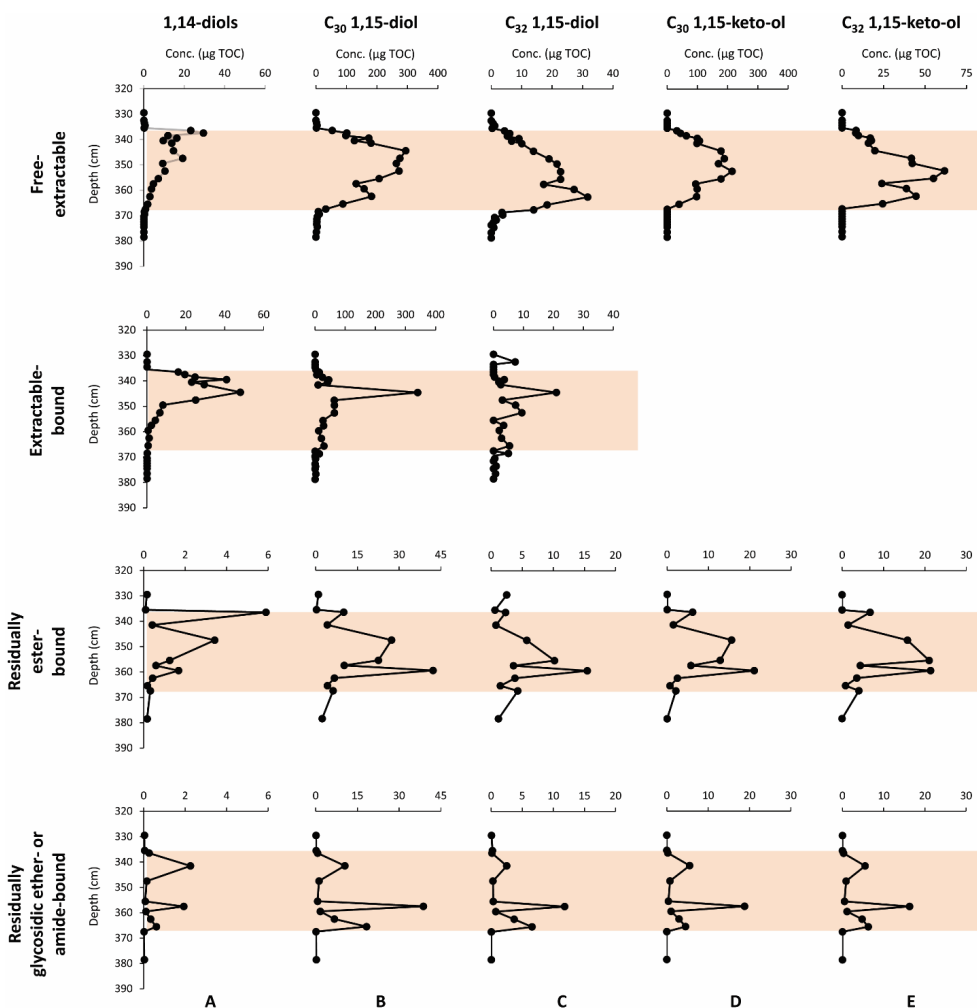


Fig. 2: Trends of 1,14- (A), C₃₀ 1,15- (B) and C₃₂ 1,15-diols (C), as well C₃₀ 1,15- (D) and C₃₂ 1,15-keto-ol (E) through the S5 sapropel as free-extractable, extractable-bound, residually ester-bound and residually glycosidic ether or amide-bound lipids. Shading indicates S5 sapropel.

3.2. LCD and keto-ol indices

A variety of LCD indices, i.e. the LDI, the Diol Index I, the FC₃₂ 1,15-diol and the DOXI, were calculated using these data. Since most LCDs and keto-ols used for their calculation were not detected outside of the sapropel, these indices could only be calculated for the sapropel sediments.

The LDI for the different modes of occurrences yielded SST estimates ranging from 25.2 to 27.0 °C. These values are 2 to 5 °C higher than previously reported SST for the S5 sapropel calculated using different calibrations of the $U^{k_{37}}$ index (Emeis et al., 1998, 2003) and also up to 10 °C higher for calculated $U^{k_{37}}$ SST for the same S5 sapropel that was studied here (Witkowski et al. (submitted)).

The LDI calculated from free-extractable LCDs stayed relatively constant throughout the sapropel with SSTs of 26.8 to 27.0 °C (Fig. 3B). A slight decrease in the LDI can be discerned for the upper 1 cm of the sapropel. The absolute values and trends for the LDI calculated from extractable-bound, residually ester-bound and residually glycosidic ether- or amide-bound LCDs were similar to those of the LDI calculated from free-extractable LCDs (Fig. 3B). A slight shift to lower values of 25.2 to 26.6 °C was visible in the LDI calculated from residually ester-bound and residually glycosidic ether- or amide-bound LCDs, respectively.

The Diol Index I, used here rather than Diol Index II due to the predominance of the C₃₀ 1,15-diol and calculated from free-extractable LCDs, increased over the sapropel from 0.02 at 366 cm core depth to 0.30 at 337 cm core depth (Fig. 3A). The same trend and values were also visible for the Diol Index I calculated from extractable-bound, residually ester-bound and residually glycosidic ether- or amide-bound LCDs (Fig. 3A).

The FC₃₂ 1,15-diol calculated for free-extractable LCDs had an early maximum of 18% at 366 cm core depth gradually decreasing thereafter to 345 cm core depth reaching 5% (Fig. 3C). A slight increase to 8% was apparent at 337 cm core depth. A general decreasing trend over the sapropel was apparent for the FC₃₂ 1,15-diol of extractable-bound and residually ester-bound and residually glycosidic ether- or amide-bound LCDs (Fig. 3C). Values were slightly higher for the FC₃₂ 1,15-diol calculated from residually ester-bound and residually glycosidic ether- or amide-bound LCDs compared to the FC₃₂ 1,15-diol calculated from free-extractable LCDs.

The C₃₀ 1,15-DOXI calculated from free-extractable LCDs and keto-ols was approximately constant throughout the sapropel with values between 0.30 to 0.46 and maxima at 356 and 341 cm core depth (Fig. 3D). Similar trends were apparent in the C₃₂ 1,15-DOXI profile calculated from free-extractable LCDs and keto-ols, but values were slightly higher between 0.57 to 0.73. The C₃₀ 1,15-DOXI calculated from residually ester-bound and residually glycosidic ether- or amide-bound LCDs and keto-ols stayed approximately constant throughout the sapropel at values of 0.25 to 0.38 and 0.20 to 0.40, respectively (Fig. 3D). The C₃₂ 1,15-DOXI calculated from residually ester-bound and residually glycosidic ether- or amide-bound LCDs and keto-ols showed gradually increasing values throughout the sapropel with slight decreases in the upper sapropel. Values ranged between 0.48 to 0.74 for the C₃₂ 1,15-DOXI calculated from residually ester-bound and between 0.49 to 0.75 for residually glycosidic ether- or amide-bound LCDs and keto-ols (Fig. 3D). The DOXI calculated from extractable-bound LCDs and keto-ols could not be calculated, since the latter were excluded from the discussion.

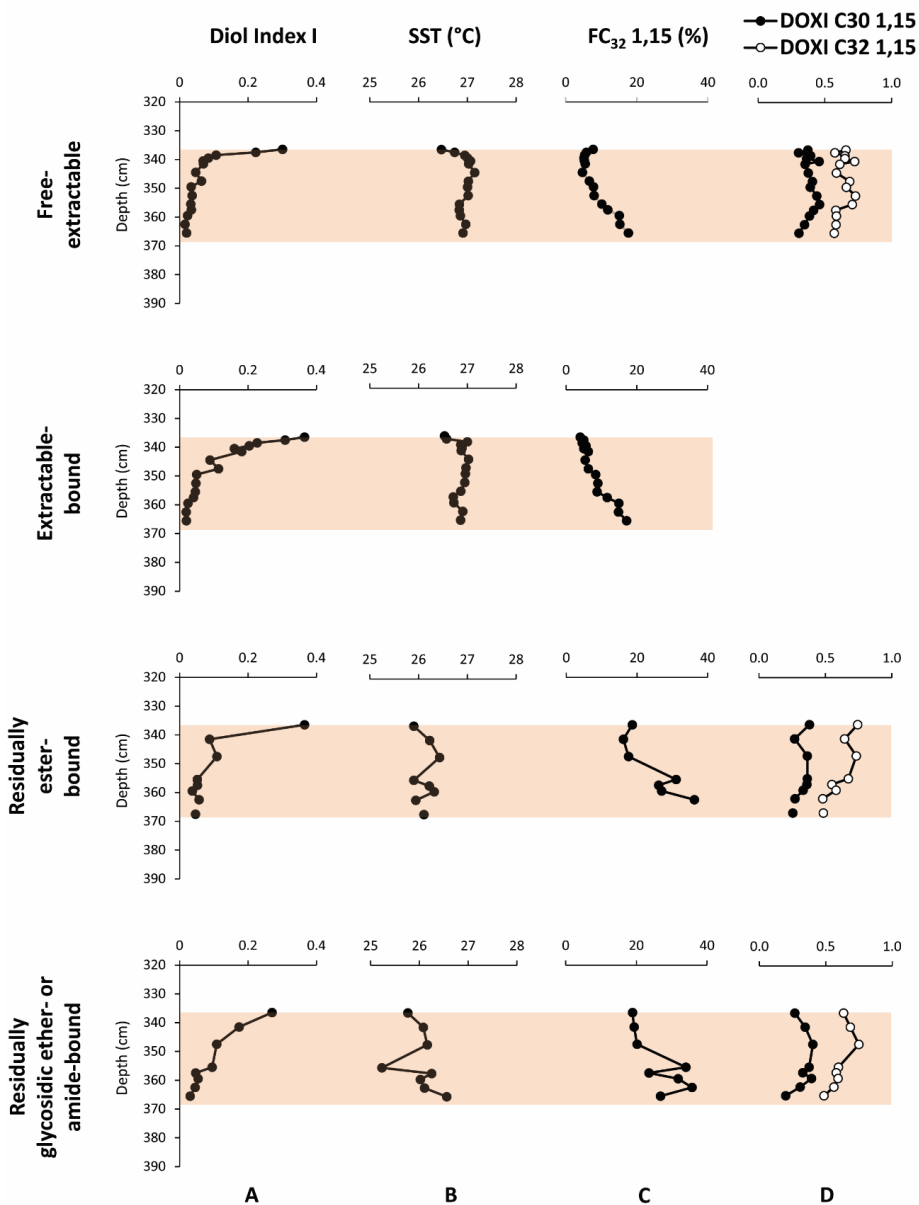


Fig. 3: Trends of Diol Index I (A), LDI SSTs (B), FC₃₂ 1,15-diol (C) and C₃₀ and C₃₂ 1,15-DOXI (D) through the S5 sapropel calculated from free-extractable, extractable-bound, residually ester-bound and residually glycosidic ether or amide-bound lipids. Shading indicates S5 sapropel.

4. Discussion

4.1. Impact of the mode of occurrence of LCDs on LCD profiles and indices

Most studies which analyzed LCDs in sediments, only report freely occurring LCDs extracted using common extraction techniques. Here, in addition, we studied bound LCDs released by hydrolysis of the extract as well as the sediment residue, both within and outside of the S5 sapropel. With respect to LCDs, it is apparent that extractable-bound 1,14-diols show higher concentrations than free-extractable 1,14-diols, suggesting some LCDs are bound to other molecules in the extract. The mode of occurrence of 1,14-diols has not been studied comprehensively to our knowledge. Rampen et al. (2007) applied a base hydrolysis to ultrasonic extracts of *Proboscia indica*, *Proboscia alata* and *Proboscia inermis*, thereby analyzing extractable-bound 1,14-diols, but did not study free-extractable LCDs released by direct extraction. Rampen et al. (2014) compared LCDs in ASE-extracted sediments with LCDs in ASE-extracted and subsequently base hydrolyzed sediments and found higher abundances of mono-unsaturated 1,14-diols after base hydrolysis, but did not discuss other LCD-isomers. Here, the C_{28:1} 1,14-diol was only present in trace amounts as a free-extractable lipid and could not be detected as a bound lipid.

In order to analyze the impact of the mode of occurrence on the different LCDs over depth of the S5 sapropel, we calculated the fractional abundances of 1,14-diols and 1,13- and 1,15-diols as free-extractable, extractable-bound, residually ester-bound and residually glycosidic ether- or amide-bound lipids below, inside and above the S5 sapropel (Fig. 4). Our results show that within the sapropel, the free-extractable and extractable-bound 1,14-diols (Fig. 4A) and 1,13- and 1,15-diols (Fig. 4B) represent 71 – 94 % and 74 – 92 %, respectively, making them the most dominant pool of LCDs within in the sapropel. 1,13- and 1,15-diols were predominantly present as free-extractable lipids (57 – 86 %) and in lower abundances as extractable-bound LCDs (6 – 30%), except in the sediments of 336 cm core depth (20 % free-extractable 1,13- and 1,15-diols and 60 % extractable-bound 1,13- and 1,15-diols). In contrast, 1,14-diols were present as both free-extractable and extractable-bound lipids, e.g. 30 – 62 % for free-extractable 1,14-diols and 9- 64 % for extractable-bound 1,14-diols. Residually bound LCDs may still represent a substantial portion of total LCDs within the sapropel; they vary from 6 – 29 % for 1,14-diols and 8 – 26 % for 1,13- and 1,15-diols.

In contrast, outside of the sapropel, freely occurring 1,14- and 1,13- and 1,15-diols are not detected and LCDs occur solely as bound lipids, predominantly bound as residually ester-bound lipids (83 – 89 % for 1,14-diols and 93 – 95 % for 1,13- and 1,15-diols) and in low abundances as residually glycosidic ether- or amide-bound lipids (11 – 17 % for 1,14-diols and 5 – 7 % for 1,13- and 1,15-diols) (Fig. 4). These residually bound LCDs may be predominant due to their recalcitrant and resistant nature as lipids are bound to the sedimentary matrix (Wakeham and Canuel, 2006) and therefore protected against microbial oxic degradation during deposition of the non-sapropel sediments in oxic bottom water. An impact of oxic degradation on LCDs and related proxies has been indeed been shown by a few studies, which mostly analyzed freely occurring LCDs (Ferreira et al., 2001; Hoefs et al., 2002; Sinninghe Damsté et al., 2002a; Bogus et al., 2012; Rodrigo-Gámiz et al., 2016) and have described decreasing concentrations of the individual LCDs upon oxygen exposure as well as decreasing LDI temperatures (Sinninghe Damsté et al., 2002a; Rodrigo-Gámiz et al., 2016). While Hoefs et al. (2002) have also shown decreasing concentrations of LCDs upon oxic degradation in Madeira Abyssal Plain turbidites, they also showed higher preservation efficiencies of residually ester-bound LCDs compared to free-extractable LCDs.

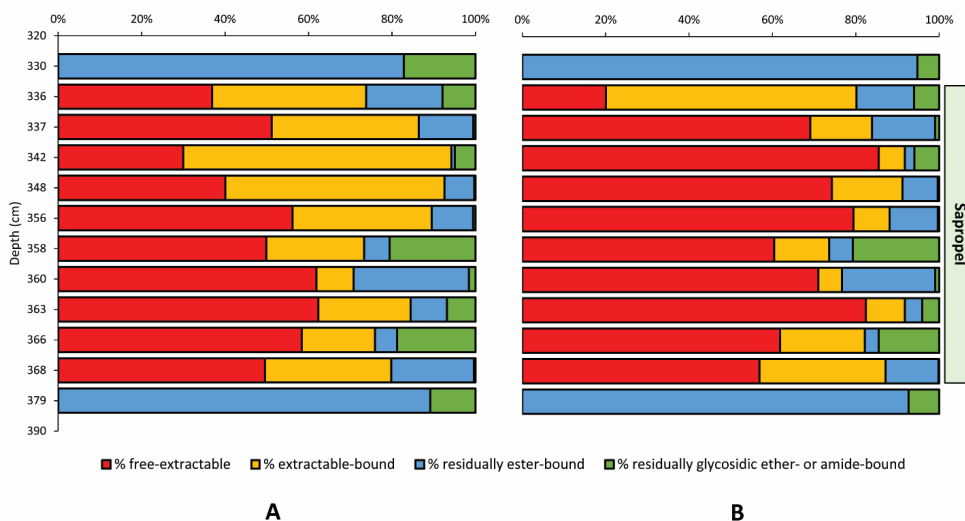


Fig. 4: Fractional abundances of A: 1,14-diols and B: 1,13- and 1,15-diols as free-extractable, extractable-bound, residually ester-bound and residually glycosidic ether or amide-bound LCDs over depth of sapropel.

Despite these substantial differences in the mode of occurrence, little difference is apparent in the various LCD indices, i.e. Diol Index I, LDI and FC₃₂ 1,15-diol, calculated from either freely occurring or bound LCDs show similar trends and values (Fig. 3), indicating that the mode of occurrence does not have a large impact on the application of LCD indices in paleoenvironmental reconstructions. These results are somewhat different to the results of an aerobic incubation of *Nannochloropsis oculata* biomass, where the LDI varied strongly when calculated from free- extractable, bound LCDs and residually bound LCDs (Reiche et al., 2018). Possibly, the sedimentary LCDs are sourced from multiple species of algae with different LCD distributions as well as different amounts of free-extractable, bound LCDs and residually bound LCDs.

4.2. Paleo-environmentally significant trends in LCD concentration profiles

The presence of 1,14-diols, the C₃₂ 1,15- and the C₃₀ 1,15-diols has also been shown in previous studies of sapropels (Ten Haven et al., 1987a, 1987b, 1987c; Ferreira et al., 2001; Menzel et al., 2003; Plancq et al., 2015). All three LCDs showed different trends throughout the sapropel studied here, e.g. different maximum concentrations as free-extractable lipids at different depths. The concentration of the C₃₂ 1,15-diol maximizes at the base of the sapropel, i.e. their production/delivery maximized during the early phase of sapropel deposition. In contrast the C₃₀ 1,15-diol has a mid-sapropel maximum and the 1,14-diols showing late sapropel maxima. The C₃₀ 1,15-diol follows the trend of TOC and the general productivity trends as indicated by Ba/Al (cf. Bale et al., 2019), suggesting that its producer bloomed in concert with the majority of phytoplankton. In contrast, the concentration maximum of 1,14-diols in the upper sapropel (Fig. 2A) indicates that its producers, i.e. *Proboscia* diatoms (Sinninghe Damsté et al., 2003; Rampen et al., 2007), did not follow general productivity trends and proliferated at a later stage. It has been suggested that rhizosolenid diatoms are important contributors to the phytoplankton

community at the time of the sapropel deposition (Kemp et al., 1999). Potentially, *Proboscia* diatoms were outcompeted by other diatoms and could only proliferate, when certain nutrient conditions were more favorable. Lower silica concentrations have been discussed to favor the growth of *Proboscia* diatoms, i.e. *Proboscia alata*, as due to their weakly silicified frustule, less silica may be required for growth than for heavily silicified diatom species (Sakka et al., 1999; Sinninghe Damsté et al., 2003). Indeed, higher concentrations of lightly silicified diatoms were shown at the end of sapropel deposition of an S5 sapropel in a core collected south of Crete (Sangiorgi et al., 2006). Additionally, increased abundances of *Proboscia* following periods of elevated diatom abundances have been shown for a sedimentary record of the Peru upwelling region (de Bar et al., 2018). Thus, the late maximum in the 1,14-diols may be due to more favorable nutrient conditions for *Proboscia* diatoms.

The early sapropel concentration maximum of the C₃₂ 1,15-diol can be linked to the conditions leading to the formation of the sapropel itself. Sapropels in the Eastern Mediterranean are generally thought to result from a combination of surface water freshening, reducing deep water ventilation, and an increased export production, increasing oxygen consumption at depth. The reduced ventilation is probably intimately linked to an intensification of the summer monsoon and therefore an increased input of freshwater into the Mediterranean basin by drainage from large river catchments like the river Nile during precession minima (Rossignol-Strick, 1983, 1985; Rohling et al., 2015). Indeed, several studies using e.g. δD of alkenones (van der Meer et al., 2007), $\delta^{18}O$ of foraminifera carbonate tests (Rossignol-Strick et al., 1982, 1985; Rohling et al., 2002) and biomarkers for freshwater cyanobacteria (Bale et al., 2019) have shown large freshwater inputs into the Mediterranean basin at the onset of sapropels in general, and sapropel S5 specifically. The C₃₂ 1,15-diol can be derived from riverine sources in coastal regions as it is a dominant LCD in rivers (de Bar et al., 2016; Lattaud et al., 2017a, 2017b). The relatively high levels of the C₃₂ 1,15-diol (19%) at the onset of the sapropel are therefore a strong indication for the input of Nile freshwater and the following decrease suggest a reduced input of riverine freshwater, as evidenced by other proxies in other S5 sapropels (Rohling et al., 2002; van der Meer et al., 2007)

4.3. Keto-ols as potential oxidation products of LCDs

Two major keto-ols were identified in this sapropel, the C₃₀ and C₃₂ 1,15-keto-ol, with the C₃₀ 1,15-keto-ol being the more abundant one. Both were not detected outside of the sapropel. The occurrences of keto-ols agree with previously published results on keto-ols in different sapropels from the Mediterranean (Smith et al., 1983; Ten Haven et al., 1987a, 1987b, 1987c; Versteegh et al., 1997; Ferreira et al., 2001) and similar trends of keto-ols over sapropel depth have also been shown in S1 sapropel sediments (Ferreira et al., 2001).

Since keto-ols are structurally related to LCDs, their formation by oxidation of LCDs (Ferreira et al., 2001; Sinninghe Damsté et al., 2003) has been proposed. Analysis of an S1 sapropel, heavily impacted by post-depositional oxidation as indicated by profiles of TOC and Ba/Ti (Ferreira et al., 2001), showed an enrichment of keto-ols relative to LCDs, expressed in the DOXI, at the base of the oxidized part of the sapropel, indicating a potential formation of keto-ols as oxidative intermediate products from LCDs. Here, in comparison, profiles of TOC and Ba/Al do not indicate a strong impact of post-depositional oxidation, i.e. at the top of the sapropel TOC levels drop in concert with Ba/Al levels (cf. Bale et al. (2019)). Indeed, the DOXI did not show any significant trends in contrast to those of Ferreira et al. (2001) while values were on average higher here compared to the C₃₀ and C₃₂ 1,15-DOXI in the unoxidized interval of S1 sapropel sediments, reported by Ferreira et al. (2001). Differences in DOXI values between both sapropels may be related to the extent of photic zone euxinia in the water column.

While the S5 sapropel has been discussed to be deposited from a fully anoxic water column (Rohling et al., 2006; Sinninghe and Hopmans, 2008), this was likely not the case during the S1 sapropel (Ferreira et al., 2001).

While the DOXIs for both chain lengths show similar trends when calculated from free-extractable LCDs and keto-ols, a slight diverging trend is visible for the C₃₂ 1,15-DOXI calculated from residually ester-bound and residually glycosidic ether- or amide-bound LCDs and keto-ols indicating a higher preservation of the C₃₂ 1,15-keto-ol versus C₃₂ 1,15-diol. If a formation of keto-ols by oxidation of LCDs is assumed, the high riverine input of the C₃₂ 1,15-diol likely led to an increased formation of the respective keto-ol due to a longer transport and therefore increased impact of oxic degradation. Since the subsequently decreasing concentration of the C₃₂ 1,15-diol indicate an increased degradation of this LCD and likely its derived keto-ol, residually ester-bound and residually glycosidic ether- or amide-bound keto-ols might be better preserved due to their more recalcitrant bonds to the sedimentary matrix. Finally, no 1,14-keto-ols were detected in this study. This agrees with previous studies and has been suggested to be due to the rapid sinking through the water column due to the larger size of the diatoms, in comparison to keto-ols potentially produced by the Eustigmatophytes, thereby reducing the time to oxygen exposure (Sinninghe Damsté et al., 2003).

5. Conclusions

In order to elucidate the impact of the mode of occurrence on long chain LCDs, keto-ols and related indices, we analyzed freely occurring as well as extractable-bound, residually ester-bound and residually glycosidic ether or amide-bound LCDs and keto-ols in an S5 sapropel from the Eastern Mediterranean. The C₃₀ 1,15-diol as the most abundant LCD identified, showing a mid-sapropel maximum in concentration while the C₃₂ 1,15-diol showed a maximum at the base of the sapropel. The latter is in line with an increased riverine freshwater input during sapropel onset into the Mediterranean basin. The 1,14-diols showed a late maximum in the sapropel, indicating a late establishment of the *Proboscia* diatom community. In the sapropel, freely occurring and extractable-bound LCDs were predominant, whereas above and below the sapropel LCDs were present as residually ester-bound and residually glycosidic ether or amide-bound LCDs. The LDI, LCD Index I and FC₃₂ 1,15-diol calculated from freely occurring and bound LCDs showed similar values and trends irrespective of the bond of LCDs used to calculate the indices, indicating that matrix sequestration does not have an impact on LCD indices in paleoenvironmental reconstructions. Keto-ols were only present in the sapropel and calculated DOXIs did not show significant trends throughout this S5 sapropel, indicating a lack of post-depositional oxidation.

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Chapter 6

The effect of post-depositional oxidation on long-chain diol proxies in the f-turbidite of the Madeira Abyssal Plain

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Abstract

Long-chain diols (LCDs) are increasingly used in paleo-environmental reconstructions. However, few studies have considered the impact of diagenesis on the application of LCD proxies. Here, the impact of post-depositional oxidation on the abundance and distribution of LCDs and mid chain keto-ols (keto-ols) as well as LCD proxies are studied in the f-turbidite of the Madeira Abyssal Plain. Comparison of the upper part, which is affected by post-depositional oxidation, with the lower part that remained anoxic showed that the C₂₈ 1,13-diol, C₃₀ 1,15-diol and the C₃₀ 1,15-keto-ol were better preserved than other LCDs and keto-ols. The Diol Index I (an indicator of upwelling) and the FC₃₂ 1,15-diol (an indicator for river influence on the marine environment) showed decreasing values with increased oxygen exposure, while the LDI (a proxy for sea surface temperature reconstruction) showed slightly increasing values. This impact of post-depositional oxidation likely resulted from preferential preservation of the C₃₀ 1,15- and C₃₂ 1,15-diol over 1,13- and 1,14-diols. A significant increase of the C₃₀ 1,15-keto-ols relative to the 1,15-diols was apparent in the upper part of the turbidite impacted by post-depositional oxidation, suggesting that keto-ols may be intermediate products in the oxidation of LCDs. Our results show that care has to be taken in interpreting LCD proxies from sediments deposited under changing oxygen conditions.

6

1. Introduction

In order to reconstruct past environmental changes, such as sea surface temperatures (SST), several organic proxies are used in paleoclimate studies. For example, SST reconstructions using the U^K₃₇ index are based on the degree of unsaturation of long-chain alkenones, produced by haptophyte algae (Brassell et al., 1986; Prahl and Wakeham, 1987), while SST reconstructions using the TEX₈₆ proxy are based on the number of cyclopentane moieties in glycerol dialkyl glycerol tetraether lipids (GDGTs) (Schouten et al., 2002), which in the marine environment are produced by Thaumarchaeota (Schouten et al., 2000; Sinninghe Damsté et al., 2002b; Besseling et al., 2019; Zeng et al., 2019).

More recently, proxies based on long-chain alkyl diols (LCDs) have been developed for palaeoceanographic reconstructions. LCDs occur in a large variety of marine and freshwater environments (Versteegh et al., 1997; Sinninghe Damsté et al., 2003; Zhang et al., 2011; Romero-Viana et al., 2012; Rampen et al., 2014a, 2014b; de Bar et al., 2016, 2020; Lattaud et al., 2017b) and are generally composed of C₂₈ to C₃₂ *n*-alkyl carbon chains with alcohol moieties at the first carbon atom and at a mid-chain position (i.e. at C₁₂ to C₁₅). Saturated and mono-unsaturated C₂₈ and C₃₀ 1,14 diols have been shown to be predominantly produced by *Proboscia* diatoms (Sinninghe Damsté et al., 2003; Rampen et al., 2007) and saturated C₂₈ to C₃₂ 1,14-diols have been identified in *Apedinella radians* (Rampen et al., 2011). LCDs with the mid-chain moiety at an odd numbered carbon atom, i.e. 1,13- and 1,15-diols, have been identified in cultures of marine and freshwater Eustigmatophyte alga and range in chain length from C₂₄ to C₃₆ (Volkman et al., 1992, 1999; Gelin et al., 1997b; Versteegh et al., 1997; Méjanelle et al., 2003; Shimokawara et al., 2010; Rampen et al., 2014a; Balzano et al., 2017). However, LCD distributions of 1,13- and 1,15 diols in marine environments differ from LCD distributions in marine Eustigmatophyte cultures, suggesting unknown marine sources (Volkman et al., 1992; Versteegh et al., 1997; Rampen et al., 2012, 2014a; Balzano et al., 2018; Reiche et al., 2018).

Several indices based on LCD distributions have been proposed to reconstruct paleo-environmental conditions. The Long-chain Diol Index (LDI), based on the fractional abundances of C₂₈ 1,13-, C₃₀ 1,13- and C₃₀ 1,15-diols, has been proposed as an SST proxy (Rampen et al., 2012), while the Diol Indices using the abundances of C₂₈ and C₃₀ 1,14-diols

relative to C₂₈ and C₃₀ 1,13-diols (Willmott et al., 2010) or to C₃₀ 1,15-diols (Rampen et al., 2008), have been used to reconstruct upwelling/nutrients and productivity (de Bar et al., 2018; Zhu et al., 2018; 2020). More recently, the fractional abundance of the C₃₂ 1,15-diol (FC₃₂ 1,15-diol) has been used to study riverine inputs into marine environments (de Bar et al., 2016; Lattaud et al., 2017a, 2017b; Häggi et al., 2019). These indices have since their development been applied in paleo-environmental reconstructions of different marine environments (Pancost et al., 2009; Rampen et al., 2012; Seki et al., 2012; Lopes dos Santos et al., 2013; Rodrigo-Gámiz et al., 2014; Plancq et al., 2015; Jonas et al., 2017; Dauner et al., 2019; de Bar et al., 2019; Lattaud et al., 2019a; Zhu et al., 2020). Additionally, the Diol Oxidation Index (DOXI), based on the ratio of LCDs and structurally related mid-chain keto-ols (keto-ols) of identical chain length and position of functional groups, has been proposed as a tool to assess the impact of post-depositional oxidation (Ferreira et al., 2001), although this has been questioned (Versteegh and Lipp, 2019).

Although LCD indices are now regularly used in paleo-environmental reconstructions, only few studies consider the impact of diagenesis on the distribution and abundance of LCDs and the application of LCD indices. In general, oxic degradation has been shown to have a profound effect on the distribution of biomarkers (Sun and Wakeham, 1994; Hoefs et al., 1998a; Prah et al., 2003; Rontani et al., 2009, 2013), including paleoclimate proxies such as the U^K₃₇ proxy (Gong and Hollander, 1999; Hoefs et al., 2002; Huguet et al., 2009; Kim et al., 2009a). Several studies have analyzed the impact of oxic degradation on LCDs and LCD indices (Ferreira et al., 2001; Hoefs et al., 2002; Sinninghe Damsté et al., 2002a; Rampen et al., 2007; Bogus et al., 2012; Rodrigo-Gámiz et al., 2016; Reiche et al., 2018). Sinninghe Damsté et al. (2002) analyzed LCDs in sediment cores from the Arabian Sea finding relatively high preservation efficiencies of LCDs compared to steroids and alkenones. Hoefs et al. (2002) studied the impact of oxygen exposure on the distribution of LCDs in Madeira Abyssal Plain turbidites, which consist of anoxic shelf sediments deposited in oxic waters thereby creating an oxidation front. By comparing the oxidized part with the unoxidized part of the turbidite, a higher preservation of residually ester-bound LCDs compared to freely extractable LCDs was shown. Rodrigo-Gámiz et al. (2016) showed a decrease in LCD abundances with increasing oxygen exposure in Arabian Sea surface sediments under contrasting redox conditions and a higher preservation of 1,14- and 1,13-diols compared to 1,15-diols, which resulted in a decrease in LDI with increasing oxygen exposure time. The latter result is in contrast to a laboratory incubation study by Reiche et al. (2018), investigating the impact of oxygen exposure on LCD-containing biomass, which showed a negligible effect of oxic degradation on the LDI.

Here, we analyze the impact of post-depositional oxic degradation on LCD and keto-ol proxies by analyzing sediments from a turbidite from the Madeira Abyssal Plain (MAP), which was previously investigated for GDGTs and associated proxies (Lengger et al., 2013). The results are compared with those of previous studies investigating the impact of post-depositional oxidation on LCDs and LCD indices.

2. Material and methods

2.1. Study sites and sampling

A piston core was retrieved from the Madeira Abyssal Plain (MAP) in the eastern North Atlantic at 31°26.8'N 024°48.8'W during cruise JCR209 onboard R/V James Clark Ross in September 2007 and the f-turbidite was sub-sampled directly after retrieval of the core (see Lengger et al., 2013 for details). Its location is in the area of well-studied turbidite deposits which in previous studies have been analyzed to ascertain oxic degradation effects on various biomarkers (de Lange et al., 1987, 1998; Cowie and Hedges, 1994, 1995; Prah et al., 1997,

Hoefs et al., 1998a, 2002; Sinninghe Damsté et al., 1998; Huguet et al., 2008, 2009; Lengger et al., 2013).

2.2. Lipid analysis and quantification

Aliquots of the sediment layers of core MAP-1 stored frozen, were extracted specifically for this study. Freeze dried sediment samples were extracted by accelerated solvent extraction (ASE 200, DIONEX) with a mixture of dichloromethane (DCM) and methanol (MeOH) (9:1; v/v). A known amount of the internal standard (C₂₂ 7,16 – diol, MolPort SIA, Latvia) was added to the extracts, which were subsequently separated into apolar, ketone and polar fractions by elution over an Al₂O₃ column using hexane: DCM (9:1, v/v), hexane: DCM (1:1, v/v) and DCM: MeOH (1:1, v/v). Prior to analysis by gas chromatography/mass spectrometry (GC/MS), polar fractions were silylated by adding BSTFA [N,O-bis(trimethyl-silyl)trifluoroacetamide] and pyridine and heated up to 60 °C for 20 min. Subsequently, samples were dissolved in ethyl acetate.

Samples were analyzed for LCDs and keto-ols on an Agilent 7890B gas chromatograph coupled to an Agilent 5977A mass spectrometer, equipped with a fused silica capillary column (25 m x 0.32 mm) coated with CP Sil-5 (film thickness 0.12 µm) with a helium flow of 2 mL/min. Samples were injected at 70 °C oven temperature, increasing to 130 °C at 20 °C/min and then programmed at 4°/min 320 °C which was held for 25 min with a helium flow of 2 mL/min. The mass spectrometer operated with an ionization energy of 70 eV. Lipids were quantified in selective ion monitoring (SIM) of *m/z* 187, 299, 313, 327 and 341 characteristic for the different LCDs and the diol standard and *m/z* 300, 314, 328 for the different keto-ols (Versteegh et al., 1997; Rodrigo-Gámiz et al., 2015; 2016).

LCD and keto-ol concentrations in the sediments of the MAP-1 f-turbidite were normalized on dry weight of the sediment (µg/g dry sediment).

The LDI was calculated according to Rampen et al. (2012):

$$\text{LDI} = \frac{[\text{C}_{30}1,15\text{-diol}]}{[\text{C}_{28}1,13\text{-diol}] + [\text{C}_{30}1,13\text{-diol}] + [\text{C}_{30}1,15\text{-diol}]} \quad [1]$$

using the calibration of de Bar et al. (2020):

$$\text{SST} = \frac{\text{LDI} - 0.1082}{0.0325} \quad [2]$$

The Diol Index I was calculated according to Rampen et al. (2008):

$$\text{Diol Index I} = \frac{[\text{C}_{28}1,14\text{-diol}] + [\text{C}_{30}1,14\text{-diol}]}{[\text{C}_{28}1,14\text{-diol}] + [\text{C}_{30}1,14\text{-diol}] + [\text{C}_{30}1,15\text{-diol}]} \quad [3]$$

The fractional abundance of the C₃₂ 1,15-diol was calculated according to Lattaud et al. (2017a):

$$\text{FC}_{32}1,15\text{-diol} = \frac{[\text{C}_{32}1,15\text{-diol}]}{[\text{C}_{28}1,13\text{-diol}] + [\text{C}_{30}1,13\text{-diol}] + [\text{C}_{30}1,15\text{-diol}] + [\text{C}_{32}1,15\text{-diol}]} \times 100 \quad [4]$$

The DOXI was calculated as proposed by Ferreira et al. (2001):

$$\text{DOXI} = \frac{[\text{keto-ol}]}{[\text{keto-ol}] + [\text{diol}]} \quad [5]$$

3. Result and discussion

3.1. Impact of post-depositional oxic degradation on LCDs

We used sedimentary intervals of the MAP f-turbidite which have previously been studied by Lengger et al. (2013) to study the impact of post-depositional oxidation on GDGT- based proxies, i.e. TEX₈₆ and the BIT index. Since MAP turbidites represent organic-matter rich homogenous turbidites (Buckley and Cranston, 1988) that are subsequently exposed to downward diffusing oxygen from the oxygen rich deep abyssal waters (de Lange, 1992), they are ideal to study the impact of oxidation on lipid distribution and abundance (Cowie et al., 1995, 1998; Prahl et al., 1997; Hoefs et al., 2002). The stratigraphic boundary between the oxidized and unoxidized f-turbidite was set at 1105 cmbsf, following Lengger et al. (2013) and was based on statistical clustering of XRF data, organic carbon content and grain size distribution.

3.1.1. LCD and keto-ol distributions

A suite of LCDs and keto-ols were identified in the MAP turbidite, in varying concentrations (Table 1). Of all LCDs, the C₃₀ 1,15-diol (Fig. 1B) was present in highest abundance with concentrations of 39 ± 15 µg/g in the oxidized zone and tenfold higher concentrations (i.e. 390 ± 60 µg/g) in the unoxidized zone. Other abundant LCDs were the C₂₈ 1,14-, the C₃₀ 1,14- (Fig. 1A) and the C₃₂ 1,15-diols (Fig. 1C) with concentrations of 10 ± 4 µg/g, 9 ± 3 µg/g and 8 ± 3 µg/g in the oxidized zone and 128 ± 11 µg/g, 137 ± 30 µg/g and 106 ± 24 µg/g in the unoxidized zone, respectively. The C₂₈ 1,13-, C₂₈ 1,12- and C₃₀ 1,13-diols were present in lower concentrations, while the C_{28:1} 1,14-diol was only present in trace amounts. Four different keto-ols were identified, of which the C₃₀ 1,15- (Fig. 1D) and C₃₂ 1,15-keto-ols (Fig. 1E) were most abundant with concentrations of 32 ± 20 µg/g and 17 ± 12 µg/g in the oxidized zone and 171 ± 147 µg/g and 210 ± 187 µg/g in the unoxidized zone, respectively. The C₃₀ 1,13- and C₃₀ 1,14-keto-ols were present in lower concentrations at 13 ± 8 µg/g and 5 ± 3 µg/g in the oxidized zone and 104 ± 92 µg/g and 32 ± 26 µg/g in the unoxidized zone, respectively (Table 1). No C₂₈ keto-ols were detected. Keto-ols showed slightly different trends in the unoxidized zone, when compared to LCDs, with lower values at the top and the deepest part of the unoxidized zone.

With these data, the relative preservation efficiency (RPE) between lipids in the oxidized and unoxidized zone was calculated (cf. Lengger et al., 2013):

$$RPE (\%) = 100 \times \frac{\text{average lipid concentration in oxidized zone}}{\text{average lipid concentration in unoxidized zone}} \quad [6]$$

For LCDs, RPEs varied between 6.4 to 10.8 %, with lowest RPEs for the C₂₈ 1,12-diol, C₃₀ 1,13- and 1,14-diols with 7.1 %, 7.1 % and 6.4 %, respectively, and highest RPEs with 10.8 % and 10.2 % for the C₂₈ 1,13-diol and C₃₀ 1,15-diols, respectively (Table 1, Fig. 2). A much wider range of RPEs was apparent for keto-ols, with a RPE of 8.0 % for the C₃₂ 1,15-keto-ol and 18.9 % for the C₃₀ 1,15-keto-ol. The C₃₀ 1,13- and C₃₀ 1,14-keto-ols showed RPEs of 12.3 % and 15.5 %, respectively.

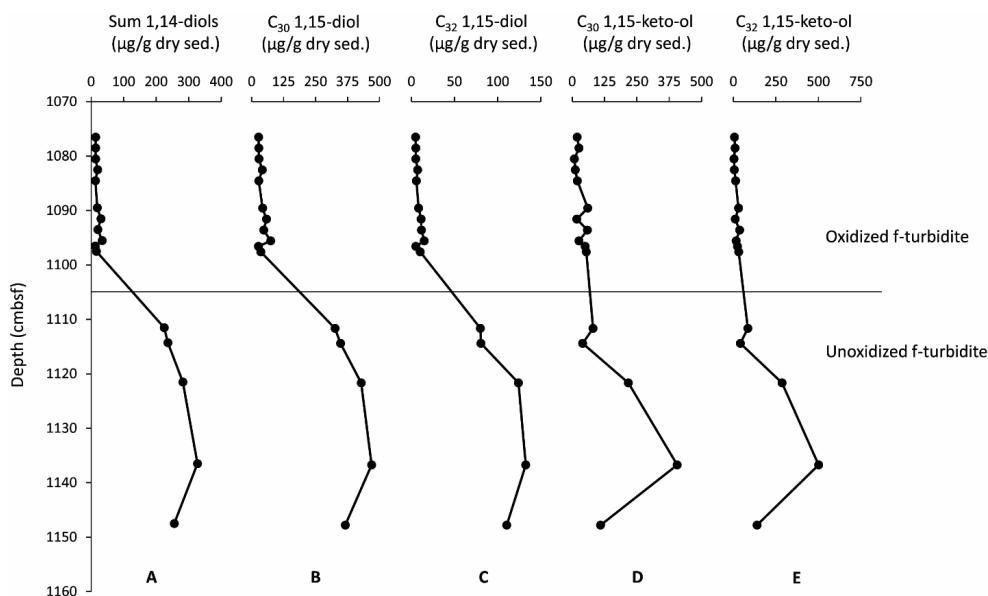


Fig. 1: Concentrations of LCDs and keto-ols with depth in the f-turbidite of the MAP-1 core. A: Sum of 1,14-diols, B: C_{30} 1,15-diol, C: C_{32} 1,15-diol, D: C_{30} 1,15 keto-ol and E: C_{32} 1,15 keto-ol. Horizontal line marks stratigraphic boundary of the oxidized f-turbidite and the unoxidized f-turbidite (*cf.* Lengger et al., 2013).

Our results indicate that the C_{28} 1,13-diol, C_{30} 1,15-diol and C_{30} 1,15-keto-ol are, among LCDs and keto-ols, potentially least affected by oxic degradation, while the C_{28} 1,12-diol, C_{30} 1,13- and C_{30} 1,14-diols and the C_{32} 1,15-keto-ol are more affected. Since 1,14-diols and 1,13-/1,15-diols are produced by different biological source organisms (Volkman et al., 1992; Gelin et al., 1997b; Versteegh et al., 1997; Sinninghe Damsté et al., 2003; Rampen et al., 2007), a difference in degradation behavior under oxic conditions between both groups might be expected. However, differences between 1,13-diols and 1,15-diols, which presumably have the same biological source as they are often found together in cultured algae (Volkman et al., 1992; Gelin et al., 1997b; Rampen et al., 2012; Reiche et al., 2018), and show a strong temperature correlation in marine sediments (Rampen et al., 2012), are harder to explain. Our results are also in contrast to Rodrigo-Gámiz et al. (2016), who analyzed Arabian Sea sediments and showed a better preservation of 1,13-diols (RPE of 11 %) over 1,14-diols (RPE of 6 %) and 1,15-diols (RPE of 4 %), irrespective of chain length. This difference in degradation behavior among different LCDs has been explained by potentially higher abundances of certain LCDs occurring as bound lipids in contrast to free lipids (Rodrigo-Gámiz et al., 2016), i.e. implying that LCDs bound to organic matter by e.g. ester-bonds were released during oxic degradation. However, more recent results by Reiche et al. (2018) showed that oxic degradation did not alter the composition of both free and bound LCDs. The major differences between our results and the study of Rodrigo-Gámiz et al. (2016) are the study area and different oxygen exposure times. Whereas Buckley and Cranston (1986) estimated an oxygen exposure time of 10 kyr for the MAP f-turbidite, Lengger et al. (2014) reported an oxygen exposure time of just up to 200 years for Arabian Sea surface sediments. However, it is unclear how the difference in oxygen exposure time would lead to different RPEs for different LCDs.

Hoefs et al. (2002) studied LCDs in older turbidites of the Madeira Abyssal Plain of Pliocene age and showed much lower RPEs for both freely extractable LCDs (C_{28} , C_{30} and C_{32} 1,15-diols) and keto-ols (C_{30} and C_{32} 1,15-keto-ols) than found here, i.e. for LCDs RPEs of < 0.5 to 2.5 % (vs. 7.1 % to 10.2 % reported here for the f-turbidite) and keto-ols 1.7 to 3.1 % (vs. 8.0 to 18.9 % for the f-turbidite). However, residual ester-bound LCDs and keto-ols (not studied here) showed significantly higher RPEs (Hoefs et al., 2002). The reason for the difference between the RPEs of extractable LCDs and keto-ols in the f-turbidite and the Pliocene turbidites is not clear and may be related to the older age and/or different oxygen exposure times.

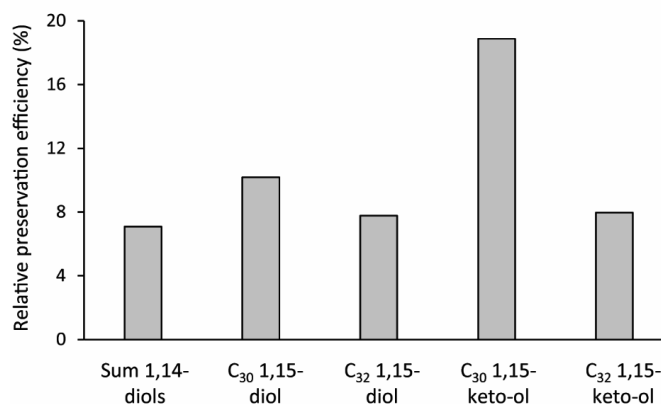


Fig. 2: Relative preservation efficiency (%) of LCDs and keto-ols in f-turbidite of MAP-1 core.

3.1.2. LCD and keto-ol indices

Commonly used LCD and keto-ol indices were calculated for the f-turbidite, i.e. the Diol Index I (Eq. 2) (Rampen et al., 2008), the LDI (Eq. 1) (Rampen et al., 2012), the fractional abundance of the C_{32} 1,15 LCD (Eq. 4) (Lattaud et al., 2017b) and various DOXIs (Eq. 5) (Ferreira et al., 2001). The Diol Index I, used here due to the predominance of the C_{30} 1,15-diol, showed a lower average value of 0.32 ± 0.01 in the oxidized zone than in the unoxidized zone (0.41 ± 0.01 , Fig. 3A). The LDI showed similar, slightly higher values (0.84 ± 0.04) in the oxidized zone than in the unoxidized zone (0.81 ± 0.03 , Fig. 3B). This would translate into reconstructed SSTs of 22.4 ± 1.3 °C in the oxidized zone and 21.7 ± 1.1 °C in the unoxidized zone of the turbidite using the SST calibration of de Bar et al. (2020). The FC_{32} 1,15-diol (Fig. 3C) decreased somewhat with increased oxygen exposure time with average values of 14.7 ± 1.6 % in the oxidized zone and 18.0 ± 1.6 % in the unoxidized zone. These indices, therefore, show an effect of post-depositional oxidation, which likely results from preferential preservation of the C_{30} 1,15-diol over 1,13- and 1,14-diols (Fig. 1).

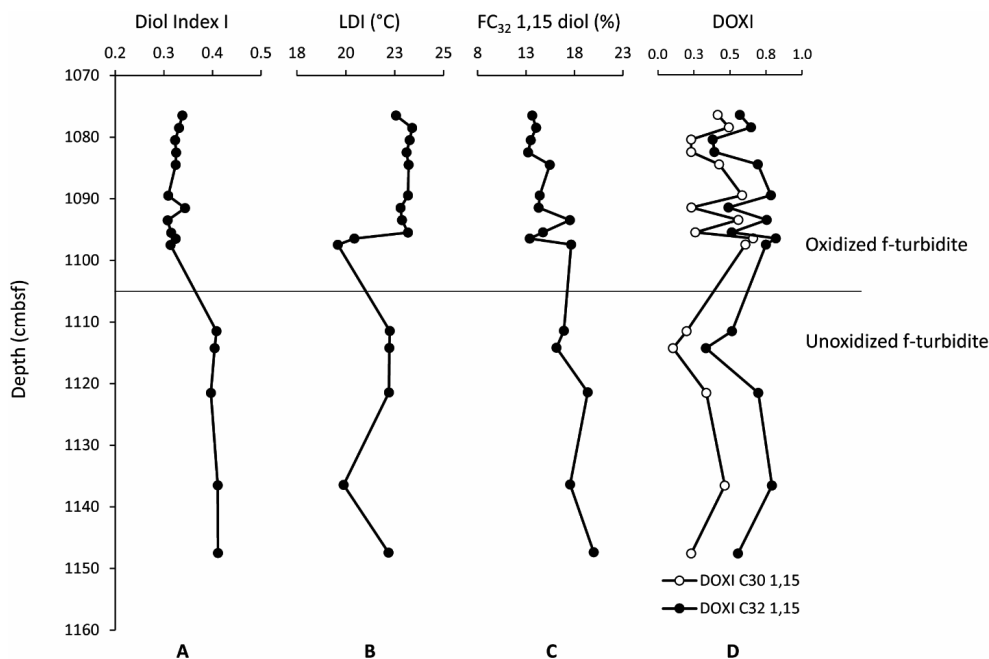


Fig. 3: Comparison of LCD and keto-ol indices over depth in f-turbidite of MAP-1 core. A: Diol Index I, B: LDI in °C, C: FC₃₂ 1,15-diol in %, D: DOXI C₃₀ 1,15 (depicted with open circles) and DOXI C₃₂ 1,15 (depicted with black circles). Horizontal line marks stratigraphic boundary of the oxidized f-turbidite and the unoxidized f-turbidite (*cf.* Lengger et al. (2013)).

This impact of post-depositional oxidation on the LDI differs from results reported for Arabian Sea surface sediments (Rodrigo-Gámiz et al., 2016). The latter study showed a decrease for the LDI, showing differences of 2.0 – 3.5 °C, due to a better preservation of the 1,13-diols. Furthermore, no impact of oxygen exposure was noted in that study for the Diol Index I as 1,15- and 1,14-diols were equally well preserved. As discussed above, the main differences are the different study regions, and possibly therefore different marine sources of diols, as well as the quite different oxygen exposure times of the turbidites versus the Arabian Sea sediments but it remains unclear how this would impact the different RPEs of LCDs. In any case, it suggests that oxic degradation can have variable effects on the different LCD indices.

The DOXI index has been applied to study paleo-oxicity and post-depositional oxidation (Ferreira et al., 2001; Bogus et al., 2012; Rodrigo-Gámiz et al., 2016), which is based on the idea that keto-ols are likely intermediate products formed by oxidation of LCDs as, for example, has been demonstrated by the study of post-depositional oxidation of sapropels (Ferreira et al., 2001). Indeed, a biological source of keto-ols is still unknown (Volkman et al., 1992; Versteegh et al., 1997; Méjanelle et al., 2003; Sinninghe Damsté et al., 2003; Versteegh and Lipp, 2019). The C₃₀ 1,15-DOXI (Fig. 3D) showed higher average values in the oxidized turbidite, i.e. 0.43 ± 0.17 in the oxidized zone vs. 0.27 ± 0.14 in the unoxidized zone. The C₃₂ 1,15-DOXI (Fig. 3D) also showed higher average values (0.62 ± 0.16 in the oxidized zone vs. 0.58 ± 0.18 in the unoxidized zone) but the difference is not significant. Therefore, especially the C₃₀ 1,15-DOXI showed the impact of post-depositional oxidation. While the DOXI values in the unoxidized

zone likely reflect oxidation of LCDs during water column settling, deposition in the surface shelf sediments and transport to the abyssal plain during turbidite deposition, the higher DOXI values in the oxidized zone may be the result of additional formation of keto-ols from LCDs by post-depositional oxidation of the turbidite. The higher overall values for the C₃₂ 1,15-DOXI, in comparison to C₃₀ 1,15-DOXI, may be due to the different origin of the C₃₂ 1,15-diols, i.e. they can be derived from an input of distant riverine sources (Lattaud et al., 2017b, 2018) and thus be oxidized to keto-ols already during transport from the river mouth to the shelf sediment. The C₃₀ 1,14-DOXI also showed higher values of 0.35 ± 0.15 in the oxidized turbidite than in the unoxidized zone (0.17 ± 0.08). If the keto-ols are formed by post-depositional oxidation of LCDs, then the somewhat lower overall values for the 1,14-DOXI, compared to the C₃₀ and C₃₂ 1,15-DOXI may be the result of a different matrix preservation of 1,14-diols over 1,13- and 1,15-diols. 1,14-diols are hypothesized to being initially less exposed to oxygen during settling through the water column due to the initially protective silica skeleton and generally heavier cells of *Proboscia* diatoms and therefore less production of corresponding keto-ols, as previously discussed by Rampen et al. (2007) and Sinninghe Damsté et al. (2003). Small changes in DOXI trends are visible in the unoxidized zone, similar to the trends of LCDs, keto-ols and the other LCD indices, and suggest some inhomogeneity in the unoxidized turbidite.

Our data therefore are in contrast with another study which showed only a weak relationship of DOXI with oxygen exposure time in the Arabian Sea (Rodrigo-Gámiz et al., 2016). In other studies, differences between LCD and keto-ol carbon chain lengths, i.e. C₂₆ to C₃₄ for LCDs and C₂₈ to C₃₆ for keto-ols, and mismatches between LCDs and keto-ols, i.e. an unsaturated C_{32:1} keto-ols without matching parent diol or C_{30:1} 1,14-diols without matching keto-ols, as evidenced by HPLC-APCI-MS analyses (Versteegh and Lipp, 2019) have also been reported.

3.1.3. Comparison of LCDs with GDGTs and GDGT proxies

RPE values for GDGTs were measured for the same f-turbidite, with RPE values of 3.5 % and 4.4 % for core lipid derived and intact polar lipid derived branched GDGTs, respectively, and RPEs of 0.25 to 0.31 % and 0.49 to 2.2 % for core lipid and intact polar lipid isoprenoid GDGTs, respectively (Lengger et al., 2013). The higher RPEs for branched GDGTs have been suggested to be the result of enhanced preservation of terrigenous derived GDGTs in contrast to marine derived GDGTs in combination with some minor contributions by in-situ production (Huguet et al., 2008; Lengger et al., 2013). In comparison, LCDs and keto-ols (RPE from 6.4 % to 18.9 %) seem to be better preserved than branched and isoprenoid GDGTs in the f-turbidite. This mirrors results on post-depositional oxidation studies of Arabian Sea sediments, which also showed a better preservation of LCDs and keto-ols over GDGTs (Sinninghe Damsté et al., 2002a).

Our analysis of the f-turbidite of the Madeira Abyssal Plain showed an impact of oxic degradation on the different LCD indices. An impact of post-depositional oxidation of the turbidite was also apparent for the BIT index (Huguet et al., 2008; Lengger et al., 2013), a proxy for the river- and soil contribution to marine environments based on branched GDGTs and crenarchaeol (Hopmans et al., 2004; Walsh et al., 2008). The preferential preservation of terrestrial-derived GDGTs over marine isoprenoid GDGTs resulted in an increase of BIT values upon post-depositional oxidation. In contrast, the FC₃₂ 1,15-diol, a tracer for input of riverine organic matter (de Bar et al., 2016; Lattaud et al., 2017b), showed a decrease. This difference may be explained by the different origins of the branched GDGTs versus the C₃₂ 1,15-diol, i.e. when derived from the continent it is mainly originating from rivers and lakes (Rampen et al., 2014a; Lattaud et al., 2017b, 2018, 2019b), while the former is, when derived from land, also sourced from soil run-off (Hopmans et al., 2004). Irrespective of the cause, our results suggest

that changes in oxygen exposure time may cause a difference between these two terrestrial proxies, i.e. the BIT index might increase, while the FC₃₂ 1,15-diol would decrease upon post-depositional oxidation, although the latter decrease is relatively minor.

With respect to the temperature proxies, the effect of oxic degradation on the TEX₈₆ was negligible in the f-turbidite (Lengger et al., 2013), which is partly in agreement with Huguet et al. (2009), who showed only small increases of TEX₈₆ in the Madeira Abyssal Plain f-turbidite with increased oxygen exposure. The small increase in the study of Huguet et al. (2009) was explained by preferential preservation of soil-derived isoprenoid GDGTs suggesting that oxic degradation may lead to small changes in the TEX₈₆. In the same study, a slight preferential degradation of the C_{37:3} alkenone upon oxygen exposure was reported, resulting in a small increase in the U^K₃₇ index, previously also shown by Hoefs et al. (1998b). These changes of the U^K₃₇ index translated to changes in reconstructed SST < 0.5 °C, while for the LDI we show changes of ca. <1 °C with post-depositional oxidation, which are just slightly larger. In contrast, Rodrigo-Gámiz et al. (2016) indicated a decrease in the LDI, equivalent to 2.0–3.5 °C, in sediments from the Arabian Sea. Together, these results suggest that oxic exposure may result in unpredictable changes in LDI, although mostly within the calibration error of the proxy (de Bar et al., 2020). Overall, a change in oxygen exposure time in sediments will therefore cause (mostly small) changes in different organic temperature proxies, i.e. LDI, TEX₈₆ and U^K₃₇, but not always in a uniform direction. This may partly explain observed mismatches between LDI, TEX₈₆ and U^K₃₇ SST records (Lopes dos Santos et al., 2013; Jonas et al., 2017; de Bar et al., 2018).

4. Conclusions

The impact of post-depositional oxidation on the abundances and distributions of LCDs and LCD proxies were determined by analysis of a turbidite from the Madeira Abyssal Plain. The C₂₈ 1,13-diol, C₃₀ 1,15-diol and the C₃₀ 1,15-keto-ol showed the highest preservation. The Diol Index I, the LDI and the fractional abundance of the C₃₂ 1,15-diol were all impacted by post-depositional oxidation to some degree, likely due to a preferential preservation of the C₃₀ 1,15-diol over 1,13- and 1,14-diol, suggesting that care has to be taken in applying these proxies in sediment cores deposited under changing oxygen exposure time. The DOXI indices showed increases in varying degrees with post-depositional oxidation, depending on the LCD chain length and mid-chain functional position, suggesting immediate formation of keto-ols.

Acknowledgments

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Table 1: Average LCD and keto-ol concentrations (variation indicated by standard deviation, $n = 11$ for oxidized turbidite, $n = 5$ for unoxidized turbidite), their derived relative preservation efficiency (RPE) and values for LCD indices in the oxidized and unoxidized zones of the f-turbidite.

Parameter	Oxidized Zone	Unoxidized zone	RPE (%)
<i>Concentrations ($\mu\text{g/g dry sed.}$)</i>			
C ₂₈ 1,12-diol	3 ± 1	37 ± 3	7.1
C ₂₈ 1,13-diol	3 ± 1	28 ± 9	10.8
C ₂₈ 1,14-diol	10 ± 4	128 ± 11	7.8
C ₃₀ 1,13-diol	4 ± 2	64 ± 27	7.1
C ₃₀ 1,14-diol	9 ± 3	137 ± 30	6.4
C ₃₀ 1,15-diol	39 ± 15	387 ± 60	10.2
C ₃₂ 1,15-diol	8 ± 3	106 ± 24	7.8
C ₃₀ 1,13-keto-ol	13 ± 8	104 ± 92	12.3
C ₃₀ 1,14-keto-ol	5 ± 3	32 ± 26	15.5
C ₃₀ 1,15-keto-ol	32 ± 20	171 ± 147	18.9
C ₃₂ 1,15-keto-ol	17 ± 12	210 ± 187	8.0
<i>LCD indices</i>			
Diol Index I	0.32 ± 0.01	0.41 ± 0.01	
LDI	0.84 ± 0.04	0.81 ± 0.03	
LDI SST (°C)	22.4 ± 1.3	21.7 ± 1.1	
FC ₃₂ 1,15-diol (%)	14.7 ± 1.6	18.0 ± 1.6	
C ₃₀ 1,13 DOXI	0.69 ± 0.14	0.56 ± 0.14	
C ₃₀ 1,14 DOXI	0.35 ± 0.15	0.17 ± 0.08	
C ₃₀ 1,15 DOXI	0.43 ± 0.17	0.27 ± 0.14	
C ₃₂ 1,15 DOXI	0.62 ± 0.16	0.58 ± 0.18	

Chapter 7

Synthesis

Since their development as proxies for paleo-reconstructions, the long-chain diols (LCD) proxies, i.e. the LDI and the Diol Index as well as newly established proxies such as the FC₃₂ 1,15-diol index, are increasingly used. The LDI has since the start of work described in this thesis been applied in a variety of regions to reconstruct SST, along with the UK'₃₇ and TEX₈₆ proxies (Nieto-Moreno et al., 2015; Plancq et al., 2015; Rodrigo-Gámiz et al., 2015; de Bar et al., 2016, 2018; Jonas et al., 2017; Warnock et al., 2018; Zhu et al., 2018, 2019; Dauner et al., 2019; Lattaud et al., 2019a; Crampton-Flood et al., 2020; García-Alix et al., 2020; He et al., 2020). Furthermore, the LDI calibration has been refined (de Bar et al., 2020) and regional calibrations have been applied (Lattaud et al., 2019a; Yang et al., 2020). The FC₃₂ 1,15-diol index has been recently developed to ascertain the riverine input into marine environments and applied in a variety of regions (de Bar et al., 2016; Lattaud et al., 2017a, 2017b; 2018; Häggi et al., 2019; He et al., 2020). Additionally, a number of studies have analyzed upwelling, nutrient and productivity conditions using the Diol Index and the newly developed Nutrient Diol Index (de Bar et al., 2018; Gal et al., 2018, 2019; Zhu et al., 2020). However, a lack of studies analyzing the impact of diagenesis on the application of LCD proxies is apparent. While a development and refinement of new and existing proxies is necessary and application in paleoenvironmental studies leads to new insights when developing climate proxies, the level of confidence in the proxy can only be raised when targeted studies are employed to constrain possible biases and diagenetic effects. In this thesis, the impact of diagenesis and sequestration on LCDs and associated proxies was studied in an effort to ascertain possible biases in the application of LCD proxies in paleoclimate reconstructions. In particular, the impact of oxygen exposure and the mode of occurrence was analyzed in algal biomass and marine sediments.

Research described in chapter 2, 3 and 4 revealed a number of issues that represent some of the underlying problems in using LCDs in paleo-reconstructive studies. The large-scale aerobic degradation of *N. oculata* biomass showed that LCDs are in fact not degraded in the time span of the incubation (271 days), nor is the structure of the algaenan affected (chapter 2 and 3). This suggests that both are well preserved, at least in suspended particulate matter and settling particles in the water column. Recently, Balzano et al. (2018) used 18S rRNA amplicon sequencing in suspended particulate matter from the western tropical North Atlantic Ocean to identify potential sources of 1,13- and 1,15-diols. However, only weak relationships between the abundance of sequences and the concentrations of the C₃₀ 1,15-diols were shown. This was attributed to the low degradability of the C₃₀ 1,15-diol compared to DNA, in agreement with the observation of low degradability of LCDs on short times scales described in thesis (chapter 2). Additionally, this suggests that it will be difficult to determine the unknown producers of marine 1,13- and 1,15-diols by comparing LCD abundances with DNA abundance or even microscopic analysis as the degradability of the LCDs is much lower than that of DNA or intact cells.

LCDs are commonly considered to be the building blocks of the biopolymer algaenan (de Leeuw et al., 1981; Tegelaar et al., 1989b; Gelin et al., 1997b, 1999; Volkman et al., 1998; Scholz et al., 2014; Zhang and Volkman, 2017). Algaenan is thought to play a role in the cell protection due to its non-hydrolysable and highly resistant character (Tegelaar et al., 1989b; de Leeuw and Largeau, 1993; Largeau and de Leeuw, 1995; Blokker et al., 2000; van Bergen et al., 2004). Indeed, a significant structure change in the algaenan of *N. oculata* was not observed, although it could not be ruled out that the abundance of algaenan diminished (chapter 3). Furthermore, it is likely that outer parts of the algaenan were affected as indicated by the release of LCDs (chapter 2). While the pyrolysis GC-MS analysis represented a qualitative approach, a quantitative approach is required to ascertain if the actual abundance of algaenan decreased during oxidic degradation. This could for example be achieved by relative quantitation using authentic standards (Hoefs et al., 1998b). Additionally, the application of additional pyrolysis GC-MS techniques might be advantageous, i.e. by tetramethylammonium hydroxide (TMAH)

pyrolysis (De Leeuw and Baas, 1993; Clifford et al., 1995; Nierop, 1998). TMAH pyrolysis might enable a more targeted breakdown of the algaenan and due to the additional methylation of more polar lipids, might lead to a more in-depth structural analysis of the algaenan. Furthermore, application of chemically degradative methods, i.e. by HI or RuO₄ degradation (Blokker et al., 1998b, 2000, 2006), could shed further light on the potentially aerobically degraded algaenan structure and composition.

Commonly, LCDs are thought to be extractable-bound in algal biomass and to only occur in low concentrations as freely extractable lipids (Volkman et al., 1992). However, as shown here, results indicate otherwise (chapter 4) as high concentrations of freely occurring LCDs were detected. Interestingly, analysis of different strains of *Nannochloropsis* spp. highlighted that application of a saponification and subsequent acid hydrolysis, which should in theory release additional lipids, in contrast led to a decrease in lipid concentration (chapter 2, 4, 5). While losses during the work-up might have led to a decrease, e.g. from shaking against water and subsequent dissolution or micelle formation, the formation of melanoidin-like polymers induced by harsh consecutive hydrolyses (Allard et al., 1997, 1998) could potentially lead to the binding of the LCDs to other compounds. The use of trifluoroacetic acid in hydrolyses (Allard et al., 1998) has been applied previously in the context of algaenan isolation as an alternative and may represent another methodological approach to be used in the analysis of bound LCDs both in biomass and sediment studies. Nevertheless, the lack of soluble macromolecules, equivalent to the algaenan (chapter 4), is somewhat surprising in that it would be expected that a less densely cross-polymerized algaenan would be soluble and degradable by chemical agents. Possibly, different approaches during the cultivation of the algal biomass, preparation of the biomass, i.e. freeze drying and storage until the work-up, and the eventual chemical work-up might lead to further insights if there are soluble parts of the algaenan.

Intriguingly, the results of the first chapters describing the analysis of LCDs in mostly aerobically degraded biomass of *Nannochloropsis* spp. (chapter 2, 3, 4) were often in contrast to the results from the natural environment (chapter 5 and 6). For example, in contrast to the results of the aerobic incubation experiment, LCDs were not only predominant as freely occurring lipids, but also occurred abundantly extractable-bound LCDs inside of the sapropel (chapter 5). In fact, above and below the sapropel, at low TOC values, LCDs occurred as residually ester-bound and residually glycosidic ether- or amide bound lipids only, although present in low concentrations. Interestingly, the finding of a higher abundance of extractable-bound 1,14-diols, compared to freely occurring 1,14-diols, in the S5 sapropel sediments is surprising as *Proboscia* diatoms represent a silica-based organism, which likely does not contain a biopolymer similar to the algaenan, in contrast to Eustigmatophyte alga. However, studies analyzing *Proboscia* biomass similarly to the approach outlined in chapters 2, 4 and 5, do not exist and could highlight how 1,14-diols are bound to the biomass.

While an impact of oxygen exposure could not be shown for the LDI in the aerobic incubation experiment (chapter 2), oxic degradation had a significant impact on LCD proxies in turbidites of the Madeira Abyssal Plain, which had been impacted by post-depositional oxidation (chapter 6). These results were furthermore mirrored by Rodrigo-Gámiz et al. (2016), who analyzed the impact of oxic degradation in Arabian Sea sediments and similarly showed an effect of oxygen exposure on LCD proxies. While both environmental studies indicated a potential preferential preservation of certain LCD isomers, the impact of oxic degradation affected the LCDs and LCD proxies differently in both studies and showed different preservation for the different isomers. Therefore, the change in values of LCD proxies upon oxygen exposure was not unidirectional nor consistent with the short-term aerobic degradation experiment (chapter 2). These differences may point to different modes of breakdown of LCDs in biomass and sediment. While the release of LCDs in the oxic degradation experiment (chapter 2) and therefore, the impact of oxygen exposure, was possibly governed by the external parts of the

algaenan, additional mechanisms of selective degradation seem to take place in the sediment. However, up to now, the knowledge of degradation mechanisms upon oxygen exposure is restricted to degradative time spans of under a year (chapter 2, 3), about 200 years (Rodrigo-Gámiz et al., 2016), and about 10 kyr (chapter 6). Therefore, further studies are needed to analyze the impact of oxygen exposure over time spans longer than 200 years and up to 10 kyr, i.e. by analysis of suitable sediments or further experimental studies. Furthermore, these studies comprise different regions with possibly different LCD producer compositions leading to a heterogeneity in the starting LCD pool prior to degradation. Possibly, the analysis of fossilized algaenan in sediments, could shed light on possible changes of algaenan structure upon longer oxygen exposure times. Pyrolysis GC-MS studies have shown a selective preservation of the aliphatic signatures, revealed by a series of *n*-alkene/ *n*-alkane pyrolysis products, likely pertaining to algaenans with increased oxygen exposure time (Hoefs et al., 1998b). A similar, targeted approach, analyzing and contrasting the LCD signatures of the algaenan of Eustigmatophytes, i.e. mid-chain ketones and *n*-alkan-1-ones, in sediments impacted by post-depositional oxidation could highlight possible changes over longer oxygen exposure times, but also further our knowledge of the algaenan of not only recent biomass but also fossilized biomass.

Overall, the application of LCDs and LCD proxies in paleoenvironmental proxies shows great promise and they will likely be increasingly used in future research, especially complimentary to already existing paleoclimate proxies. However, this thesis also highlighted that a number of issues need to be taken into consideration. An underlying problem of LCD proxy application remains the unidentified producer of 1,13- and 1,15-diols, since the mechanisms for their distribution and composition in biomass and the natural environment can only be properly understood when knowing the source organism. This also applies to gaining a better understanding of the biosynthetic pathways that are producing LCDs in the producer organism and comprehension of the nature of the biopolymer algaenan. However, this thesis especially highlighted the need for additional studies analyzing the impact of diagenesis. While open questions remain concerning the impact of oxygen exposure on LCD proxies, additional diagenetic processes have not yet been explored and need to be addressed, such as the impact of lateral transport on LCDs.

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About the author

Sophie Reiche was born on 4th March 1988 in Greifswald, Germany. She graduated from Europaschule Landesgymnasium Latina August Hermann Francke in Halle/Saale in 2007 and subsequently moved to the University of Greifswald to study Environmental Sciences. After changing her major to Geology in 2009, she graduated in 2012 with a Bachelor of Science in Geology with a specialization in Mineralogy and Quaternary Geology. Subsequently, she moved to Sweden to study at the University of Lund. She completed her Master of Science in Geology with a specialization in Quaternary Geology in September 2014 when she defended her Master thesis titled “Ascertaining the lithological boundaries of the Yoldia Sea of the Baltic Sea – a geochemical approach”. Two weeks later she started her PhD position at the Royal Netherlands Institute for Sea Research on Texel, the Netherlands, under the guidance of Prof. Stefan Schouten and Prof. Jaap Sinninghe Damsté, which has led to this doctoral thesis.

