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Hydrogen isotopes in individual alkenones from the Chesapeake Bay estuary

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Abstract

Hydrogen isotope ratios of individual alkenones from haptophyte algae were measured in suspended particles and surface sediment from the Chesapeake Bay (CB) estuary, eastern USA, in order to determine their relationship to water δD values and salinity. δD values of four alkenones (MeC_{37:2}, MeC_{37:3}, EtC_{38:2}, EtC_{38:3}) from particles and sediments were between -165%and $-221\%_{00}$ and increased linearly ($R^2 = 0.7-0.9$) with water δD values from the head to the mouth of the Bay. Individual alkenones were depleted in deuterium by 156-188% relative to water. The MeC₃₇ alkenones were consistently enriched by $\sim 12\%$ relative to the EtC₃₈ alkenones, and the di-unsaturated alkenones of both varieties were consistently enriched by $\sim 20\%$ relative to the tri-unsaturated alkenones. All of the increase in alkenone δD values could be accounted for by the water δD increase. Consequently, no net change in alkenone-water D/H fractionation occurred as a result of the salinity increase from 10 to 29. This observation is at odds with results from culture studies with alkenone-producing marine coccolithophorids, and from two field studies, one with a dinoflagellate sterol in the CB, and one with a wide variety of lipids in saline ponds on Christmas Island, that indicate a decline in D/H fractionation with increasing salinity. Why D/H fractionation in alkenones in the CB showed no dependence on salinity, while D/H fractionation in CB dinsoterol decreased by 1% per unit increase in salinity remains to be determined. Two hypotheses we consider to be valid are that (i) the assemblage of alkenone-producing haptophytes changes along the Bay and each species has a different sensitivity to salinity, such that no apparent trend in $\alpha_{alkenone-water}$ occurs along the salinity gradient, and (ii) greater osmoregulation capacity in coastal haptophytes may result in a diminished sensitivity of alkenone–water D/H fractionation to salinity changes. © 2011 Elsevier Ltd. All rights reserved.

1. INTRODUCTION

The hydrogen isotopic composition of algal lipids closely co-varies with the hydrogen isotope composition of the water in which the organisms grew (Sessions et al., 1999; Englebrecht and Sachs, 2005; Zhang and Sachs, 2007; Schwab and Sachs, 2009; Sachs and Schwab, 2011). Algal lipid D/H ratios in sediments have thus been increasingly used to reconstruct the water cycle in past environments (e.g., Sauer et al., 2001; Huang et al., 2002; Pahnke et al., 2007; Van der Meer et al., 2007, 2008; Sachs et al., 2009). However, beside the biosynthetic pathways (Sessions et al., 1999) and species-specific differences (Schouten et al., 2006; Zhang and Sachs, 2007), environmental parameters including temperature (Wolhowe et al., 2009; Zhang et al., 2009), nitrogen limited growth rate (Zhang et al., 2009), growth phase (Wolhowe et al., 2009) and salinity (Schouten et al., 2006; Sachse and Sachs, 2008; Sachs and Schwab, 2011) have been shown to influence hydrogen isotope fractionation during lipid synthesis by algae, potentially complicating paleohydrologic interpretations.

For instance, culture (Schouten et al., 2006) and field studies (Sachse and Sachs, 2008; Sachs and Schwab, 2011) demonstrated that increased salinity resulted in decreased D/H fractionation between environmental water and the

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biosynthesized lipids. Paleosalinity and paleohydrologic reconstructions based on lipid δD values soon followed (Pahnke et al., 2007; Van der Meer et al., 2007, 2008; Sachs et al., 2009; Smittenberg et al., 2011). Internal cell water is the main donor of hydrogen atoms in all positions of the lipid chain (Duan et al., 2002; Baillif et al., 2009), and thus its hydrogen isotope composition is likely to be directly reflected in the biosynthesized lipids (Kreuzer-Martin et al., 2006). It has been thus proposed that the D-enrichment in algal lipids as salinity increases reflects progressive Denrichment of internal cell water due to its continual enrichment as biosynthetic reactions draw D-depleted hydrogen from the internal water pool in combination with reduced exchange with external water (Sachse and Sachs, 2008; Sachs and Schwab, 2011). A refinement put forth by Sachs and Schwab (2011) is that turgor regulation via increased osmolyte production siphons off isotopically depleted H, leaving internal water used for lipid synthesis enriched in D. Nevertheless, the mechanisms controlling the sensitivity of lipid δD values to salinity are still being evaluated and a general theory to explain the phenomenon is lacking.

Prymnesiophyte algae are the unique producers of alkenones (Volkman et al., 1980; Marlowe et al., 1984). In the modern ocean, they are produced primarily by the species *Emiliania huxleyi* and *Gephyrocapsa oceanica*, (Marlowe et al., 1984, 1990; Conte et al., 1995; Volkman et al., 1995) while in coastal and lacustrine environments, the main producers are likely the species *Isochrysis galbana* and *Chrysotila lamellosa* (e.g., Volkman et al., 1988; Li et al., 1996; Thiel et al., 1997; Schulz et al., 2000; Chu et al., 2005; Sun et al., 2007).

Heretofore, the measurement of alkenone δD values has been performed on a mixture of the C37:3 and C37:2 alkenones owing to the inability to achieve adequate separation between the two compounds during gas chromatography (Englebrecht and Sachs, 2005; Schouten et al., 2006; Pahnke et al., 2007; Van der Meer et al., 2007, 2008). Separation techniques using column argentation chromatography or recently developed high-performance liquid chromatographymass spectrometry (HPLC-MS) permit the purification of each individual alkenone for stable hydrogen isotope analyses based on the number of carbon atoms and double bonds (D'Andrea et al., 2007; Schwab and Sachs, 2009). Using the HPLC-MS technique, a systematic ca. 20% difference in the δD values of C_{x:3} and C_{x:2} alkenones (where x = 37 or 38) was measured in sediments and particles from a site in the CB estuary (Schwab and Sachs, 2009). An identical hydrogen isotopic offset between C37:3 and C37:2 alkenones $(\alpha_{C37:3-C37:2} = (\delta DC_{37:3}+1)/(\delta DC_{37:2}+1) = 0.97)$ was recently measured in cultured E. huxleyi and G. oceanica harvested at different growth stages (Wolhowe et al., 2009). The similar isotopic offset between $C_{37:3}$ and $C_{37:2}$ alkenones in both studies, despite the likely presence of alkenone producers in the Chesapeake Bay (CB) other than E. huxleyi and G. oceanica, led Wolhowe et al. (2009) to propose that a simple isotopic mass balance calculation could be used to determine δD values for individual alkenones measured from total C₃₇ alkenones (i.e., the combined $C_{37:2} + C_{37:3}$ alkenone peak eluting from a GC) without labor-intensive chemical separations.

To test the fidelity with which individual alkenones record water δD values and to evaluate the influence of salinity on D/H fractionation in alkenones, we measured the hydrogen isotope composition of these compounds along the salinity (10–29) and water δD (-37.6% to -10.6%) gradient in the CB estuary in Maryland and Virginia, Eastern USA. The same set of samples was used by Sachs and Schwab (2011) to demonstrate that about half of the increase in dinosterol δD values along the CB resulted from increasing water δD values, and the other half from decreasing D/H fractionation as salinity increased. Our results are then used to test the hypothesis of Wolhowe et al. (2009) that the hydrogen isotope offsets between individual alkenones are constant in an estuarine environment characterized by large salinity variations.

2. MATERIALS AND METHODS

2.1. Geological setting and water chemistry

Located in the Mid-Atlantic region of North America, the CB is a N–S elongate (300 km), shallow (average depth 8.2 m) estuary with an axial channel that is 12–30 m deep (Fig. 1). Freshwater from the Susquehanna River at the head of the Bay supplies 48% of the total riverine flux of water to the CB and creates typical partially mixed estuarine circulation with an outflow of freshwater in the upper layers and an inflow of seawater at depth. A salinity gradient along the length of the CB, and as a function of depth is thus formed (e.g., Pritchard, 1967; Austin, 2004).

Multi-decadal time series of monthly physical, chemical and biological parameters in the CB are available on the website of the CB Monitoring Program http://www.chesapeakebay.net/data_waterquality). Salinity, water δD values, and water temperature at multiple depths along the N-S transect of the Bay in May 22-25, 2006 are published in Sachs and Schwab (2011). Nutrient, Chl a concentration and primary production at the same sampling locations are available on the CB Monitoring Program website. Briefly, surface (0-2 m) salinity and δD values increased from 0 to -56.8% (station CB1.1) at the head of the CB, where the Susquehanna River enters, to 28.9 and -11.4%(station CB7.4N) at the mouth of the Bay (Fig. 1). The water δD values, previously published in Sachs and Schwab (2011), were linearly correlated with salinity (increasing with depth and along the length of the estuary) according to the relationship $\delta D_{\text{water}} = 1.63 \pm 0.04 \times \text{Salinity} - 52.3 \pm 0.7,$ $R^2 = 0.91$, with uncertainties represented as the standard errors of the slope and intercept of the regression. The observed relationship reflects fractionation due to evaporation toward more saline media and the mixing line between marine and freshwater (Craig, 1961; Craig and Gordon, 1965). Surface (0-2 m) water temperatures were between 14.9 and 20.1 °C. Lower temperatures occurred in the upper-bay (CB3.2) and lower-bay (CB7.4 N) and higher temperature in the mid-bay (CB6.2) (Fig. 1). A halocline and a thermocline were observed at all CB stations, with bottom waters as much as 15 saltier and 2 °C colder than surface waters. As a result, since freshwaters were depleted in deuterium by 50% relative to seawater, water δD values



Fig. 1. Maps of the CB showing surface (0-2 m) water (a) salinity, (b) δD_{water} values and (c) temperature at sample locations (red stars green circles). The station locations from which surface sediment and suspended particle were taken for this study are shown with red stars. Maps were produced with ArcGIS Spatial Analyst, with inverse distance weighted output cell size of 0.001° , 131 salinity points, 105 water δD values, and 122 temperature points. Salinity and temperature data are from the CB monitoring program (http://www.chesapeakebay.net/ data_waterquality). (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

increased with depth at all stations, with bottom waters as much as 15% heavier than surface waters. Total dissolved nitrogen and phosphate ranged from *ca*. 0.14 to 0.88 mg L⁻¹ and from 5.0 to 23.6 µg L⁻¹, respectively, with both nutrients most abundant at the head of the Bay and least abundant at the mouth. In May 2006, surface Chl *a* concentrations were between 1.07 and 31.4 µg L⁻¹, with lowest values at the mouth (station CB7.4 N) of the Bay and highest values in the mid-Bay region. Maximum chlorophyll concentrations occurred at 3–5 m water depth throughout the Bay. Higher primary production in the mid-Bay resulted in hypoxic (1–30% saturation of dissolved oxygen) or anoxic bottom waters.

2.2. Water, particle and sediment sampling in the Chesapeake Bay

Surface sediment and suspended particles were collected at 26 of the CB Monitoring Program stations along the central axis of the Bay from the R/V Kerhin during May 22– 25, 2006 (Fig. 1). Sample collection was timed to coincide with the haptophyte bloom that was expected to occur in late May, after peak runoff fluxes occurred (Mercer et al., 2005), and within two days of both the Maryland and Virginia Monitoring Program's monthly sampling of the Bay. The coordinates of each station and the detailed sampling procedure are published in Sachs and Schwab (2011). In brief, near surface sediments, collected with a grab sampler, were sub-sampled – typically at 0–2 and 0–10 cm – and subsequently frozen at -20 °C. Water and suspended particle samples were collected at multiple depths at each sampling site – typically 0–2, 2–4, 4–5, 14 and 30 m – by tubing attached to a conductivity–temperature–depth continuous profiling instrument and filtered through glass fiber filters immediately frozen at -20 °C. Water for hydrogen isotope analysis was collected with each filter and between stations. Salinity and temperature were continuously measured on board the vessel.

2.3. Coccolithophore identification

Samples were taken to determine coccolithophore taxonomic composition at each sample depth by vacuum filtering 1-2 L of water through cellulose nitrate and polycarbonates filters (0.45 µm pore size, 47 mm diameter). Each filter was rinsed with buffered water (63 mL of NH₄OH in 500 mL of distilled water) to remove salt, then dried at 40 °C. Filters were analyzed by polarized light microscopy at the University of Washington (Seattle, USA) and by scanning electron microscopy (SEM) at the University of Lausanne, Institute of Geology and paleontology (Switzerland).

2.4. Lipid extraction and pre-treatment

The detailed procedure for extracting and purifying alkenones is reported in Schwab and Sachs (2009). In brief, $n-C_{37}$ alkane, 2-nonadecanone (C_{19} -ketone) and cholesterol

(Sigma–Aldrich) were added as internal standards to freezedried sediment and filters prior to extraction with an Automated Solvent Extractor (ASE-200, Dionex Corp., Sunnyvale, CA, USA). The solvent was removed under a stream of nitrogen using a Turbovap system (Caliper, Hopkinton, MA, USA). The extracts were separated with column chromatography on pre-combusted Al_2O_3 into four fractions: F1 was eluted with hexane/DCM (9:1, v/v) and contained hydrocarbons, F2 was eluted with hexane/ DCM (1:1, v/v) and contained aromatic lipids and ketones (including alkenones), F3 was eluted with DCM/MeOH (1:1, v/v) and contained alcohols and F4 was eluted 100% MeOH and contained acids and polar lipids. Elemental sulphur was removed from F1 and F2 using activated Cu powder.

The alkenones and alkenoates contained in the F2 were identified by Gas Chromatography-Mass Spectrometry (GC–MS) using an Agilent (Santa Clara, CA, USA) 6890 N gas chromatograph equipped with an Agilent 5983 autosampler, a split-splitless injector operated in splitless mode, and a HP-5 ms column ($30 \text{ m} \times 0.32 \text{ mm}$ i.d. $\times 0.25 \mu \text{m}$ film thickness, Agilent) interfaced to an Agilent 5975 quadrupole mass selective detector (MSD). They were quantified with a Gas Chromatograph–Flame Ionization Detector (GC–FID). The Agilent 6890 GC was equipped with an Agilent 5983 autosampler and a programmable temperature vaporization inlet (PTV) operated in splitless mode. A 60 m Varian Chrompac CP-Sil 5 capillary column (0.32 mm $\times 0.25 \mu m$) was used with helium as the carrier gas (1.6 mL/min).

2.5. Alkenone purification by HPLC-MS

Detailed methods for alkenones purification by HPLC are given in Schwab and Sachs, 2009. Briefly, an Agilent (Santa Clara, CA, USA) high performance liquid chromatograph (HPLC) 1100 series, equipped with an autoinjector, a quaternary pump, an integrated fraction collector, and Chemstation chromatography manager software was used. Typically, one injection of 100 µL of the concentrated F2 (alkenone) fraction was enough for subsequent hydrogen isotope analyses of each individual alkenone. A Prevail Cyano column (250 \times 4.6 mm \times 5µ; Alltech, Deerfield CA) maintained at 30 °C was used for separation. Compound classes were separated using an isocratic mixture of hexane and 4% DCM in hexane. Eluting compounds were identified in real-time with an Agilent 1100 SL mass spectrometer equipped with a multimode source operated in positive atmospheric pressure chemical ionization (APCI+) mode on ca. 2% of the total flow that was split from the eluent using an adjustable flow splitter (ASI, El Sobrante, CA, USA). A Waters (Milford, MA, USA) 510 pump was used to add a flow of 0.3 mL/min 2,2,4-trimethylpentane to the 2% split in order to supply sufficient solvent to the APCI source to achieve optimal ionization efficiency.

2.6. Hydrogen isotope analyses

Alkenones were quantified by comparing their integrated peak areas with that of the n-C₃₇ standard using GC–FID operating at the same conditions previously described. The target compounds were dissolved in toluene to reach the adequate concentration (250–300 ng/ μ L of the individual alkenones) needed for sufficient signal intensity on the GC–IRMS system (Trace Ultra GC and DEL-TA V from Thermo Scientific, Waltham, MA, USA). The GC was equipped with a split-splitless injector operated in splitless mode at 300 °C, a TRIPLUS autosampler (Thermo Scientific, Waltham, MA, USA), and a DB-5ms capillary column (60 m × 0.32 mm × 0.25 μ m, Agilent). Helium was used as the carrier gas at a constant flow of 1 mL/min.

Instrument performance and the H_3^+ factor were determined on a daily basis using a tank of H₂ reference gas (Sessions et al., 2001) and a mixture of n-alkanes (n- C_{14} to *n*- C_{44}) of known isotopic composition. Hydrogen isotopic composition of *n*-C₁₄ to *n*-C₃₆ *n*-alkane was determined off-line using a thermochemical elemental analyzer (TC/EA) (Thermo-Fisher, Bremen, Germany) interfaced to the DELTA V PLUS irMS system via a Conflo III combustion interface (Thermo-Fisher, Bremen, Germany). The n-C₃₈, n-C₄₁ and n-C₄₄ standards used for δD corrections and added to the *n*-alkane mixture for H_3^+ factor calculation were purchased from A. Schimmelmann, (Indiana University, Bloomington, Indiana). The average offset between GC-irMS δD values and those determined off-line was 3.9_{00}° (*n* = 3630, 165 runs with 22 peaks). The H₃⁺ factor was lower than four and stable during the measurements reported here. The precision of alkenone δD measurements, as determined by the pooled standard deviation,

 $S_p = \sqrt{\frac{\sum_{i=1}^{k} ((n_i-1)s_i^2)}{\sum_{i=1}^{k} (n_i-1)}}$ of 363 analyses of 119 samples, seven

of which were measured in quadruple, 111 of which were measured in triplicate and one in duplicate, was 1.3%.

3. RESULTS

3.1. Alkenone producers in the Chesapeake Bay

No evidence of calcified haptophyte species, such as the common marine coccolithophorids E. huxleyi and G. oceanica, was found in 10 samples investigated by light microscopy or two samples investigated by SEM despite both those species having been reported in the Bay (Marshall, 1994; Marshall et al., 2005). Although the two samples investigated by SEM (station CB4.4 and station CB7.4) do not represent a comprehensive set of all sites in the Bay, they did contain substantial concentrations of alkenones. Light microscopy revealed the occurrence of uncalcified spherical cells in all samples tentatively identified as I. galbana or C. lamellosa. Both of those species have been identified in the Bay (Marshall, 1994; Marshall et al., 2005) and are known to produce alkenones (Versteegh et al., 2001; Sun et al., 2007). However, since E. huxlevi and G. oceanica may lose their coccoliths under certain environmental conditions and/or phases of growth (Green et al., 1996; Paasche, 2002), we cannot definitively exclude their presence in the CB. Doing so would require DNA analyses since taxonomic differentiations on naked cells are nearly impossible by SEM (Coolen et al., 2004).

3.2. Alkenone distributions along the bay

Alkenones were detected throughout the Bay in suspended particles from a variety of depths (Electronic annex Table EA-1) and in surface sediments (Electronic annex Table EA-3) collected between station CB3.1 and CB7.4N at salinities ranging from 3.7 to 30.9 (Fig. 1). Alkenone concentrations in water ranged from 0.01 to 0.36 μ g L⁻¹ in the Upper-Bay, from 0.06 to $2.1 \,\mu g \, L^{-1}$ in the Mid-Bay and from 0.04 to $1.3 \ \mu g \ L^{-1}$ in the Lower-Bay (Table EA-1). Converted to mass of suspended matter, alkenone concentration were between $0.4-23.6 \ \mu g \ g^{-1}$ in the Upper-Bay, 1.8–271.4 μ g g⁻¹ in the Mid-Bay and 3.2–86.3 μ g g⁻¹ in the Lower-Bay (Table EA-1). At individual stations highest alkenone concentrations usually occurred between 1 and 5 m, close to the depth of maximum of Chl a concentration. Alkenone concentrations in sediments averaged a factor of four lower than in filters, but tracked concentrations in the overlying water, with higher concentrations of 42.5 μ g g⁻¹ at station CB6.1, and lower concentrations of 0.8- $0.6 \ \mu g \ g^{-1}$ at station CB3.3E and CB7.4 in the Lowerand Upper-Bay, respectively (Table EA-3).

All particulate and sediment samples were characterized by similar alkenone distribution patterns to those shown in Schwab and Sachs (2009), with a predominance of methyl C_{37} alkenones over ethyl C_{38} alkenones, a higher relative concentration of tri- over di- and tetraunsaturated C_{37} and C_{38} alkenones, a small proportion of C_{39} alkenones, and trace amounts of tetraunsaturated C_{38} and C_{40} alkenones (Fig. 2). Only methyl and ethyl $C_{36;3}$ and $C_{36;2}$ alkenoates were detected, with ethyl $C_{36:2}$ alkenoate predominating. A distinct characteristic of all alkenone distributions was the lack of C_{38} methyl alkenones.

The relative abundance of the tetra-unsaturated C_{37} alkenone, expressed as $C_{37:4} = C_{37:4}/(C_{37:4} + C_{37:3} + C_{37:2})$ (Marlowe et al., 1984), was between 2.3–9.5% in particles (Table EA-2) and 2.0–10.5% in surface sediments (Table EA-3), and increased with salinity along the N–S transect of the Bay and with depth (Fig. 3A). For instance, between 0.5–3 m and 23 m at the station CB3.3C, salinity increased from 10.2 to 20.7 and $C_{37:4}$ increased 3.5 and 5.7 (Fig 3A).

The relative abundance of C_{37} alkenones relative to C_{38} $C_{37}/C_{38} = (C_{37:3} + C_{37:2})/$ alkenones. expressed as $(C_{38\cdot3} + C_{38\cdot2})$ (Marlowe et al., 1984) was between 1.2–4.5 in particles and 1.1-2.1 in sediments (Tables EA-2 and EA-3). Samples between stations CB3.1 and CB7.4 indicated a decrease of C37/C38 (from 3.1 to 1.2) as salinity increased from 3.7 to 28.9 along the N-S transect of the Bay and with depth (Fig. 3B). A different trend was observed at station CB7.4N, where two particle samples collected at 5 and 11 m displayed much higher C_{37}/C_{38} ratios of 4.2 and 4.5, respectively, in the salinity range 29.7 at 5 m to 30.9 at 11 m. No correlation between either $%C_{37:4}$ or C_{37}/C_{38} with water temperature or nutrient concentration was observed.

In particles and sediments, alkenone indices such as $U_{37}^{K} = C_{37:2} - C_{37:4}/C_{37:2} + C_{37:3} + C_{37:4}$ (Brassell et al., 1986) and $U_{37}^{K'} = C_{37:2}/C_{37:2} + C_{37:3}$ (Prahl and Wakeham, 1987) displayed large variations throughout the Bay. Contrary to an earlier report by Mercer et al. (2005) U_{37}^{K} and $U_{37}^{K'}$ in particles and surface sediment showed no correlation with temperature (e.g. Fig. 4). In suspended particles, U_{37}^{K}



Fig. 2. Partial gas chromatogram showing representative alkenone distribution in surface sediments of the CB; with Me and Et being Methyl and Ethyl alkenone, respectively and Me-and Et-oate being Methyl and Ethyl alkenoate, respectively.



Fig. 3. (A) Plot showing variations of $%C_{37:4}$ ratios versus salinity in water particle (black lozenge dots) and sediment (yellow triangle dots) samples. (B) Plot showing variations of C_{37}/C_{38} ratios versus salinity in water particle (black lozenge dots) and sediment (yellow triangle dots) samples. The blue star dots show the variation of these former ratios in water particle samples collected at 0.5–3, 7–4 and 20–24 m in station CB3.3C. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)



Fig. 4. Plot showing variations of $U_{37}^{K'}$ versus measured surface temperature in water particle (black dots) and sediment (yellow dots) samples. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

and $U_{37}^{K'}$ ranged from 0.06 to 0.22 and from 0.12 to 0.26, respectively (Table EA-2). In sediment, U_{37}^{K} and $U_{37}^{K'}$ ranged from 0.12 to 0.28 and from 0.18 to 0.33, respectively (Table EA-3). Water surface temperatures derived from the relationship $U_{37}^{K'} = 0.034T + 0.039$ (Prahl et al., 1988) were between 2.3–8.4 °C in particles (Table EA-1) and

4.2-8.7 °C in sediment (Table EA-3), all 7.7-17.1 °C (average 12.4 °C) colder than the measured water temperature. Improved estimates of the water temperature were obtained using the alkenone temperature calibrations $U_{37}^{K} = 0.0377T - 0.5992$ (Sun et al., 2007) and $U_{37}^{K'} = 0.016T - 0.0607$ (Versteegh et al., 2001) calibrated with C. lamellosa and I. galbana, respectively. Applying the C. lamellosa calibration, $U_{37}^{K'}$ -derived water temperatures were between 17.6-21.7 °C in particles and 15.9-23.3 °C in sediments, which together averaged 2.7 ± 1.9 °C warmer than measured temperatures. Applying the *I. galbana* temperature calibration, $U_{37}^{K'}$ -derived water temperatures were between 11.0-24.1 °C in particles and 15.1-24.7 °C in sediments, on average just 0.3 ± 2.9 °C colder than measured temperatures (Tables EA-1 and EA-3).

3.3. Alkenone δD values

Alkenone δD values were measured in sediment and particle samples with sufficient alkenone abundance, which in this study was approximately greater than 1 µg of the total lipid extract. Such abundances occurred between stations CB3.3C and CB7.4N where surface salinities were between 10–29 and water δD values were between -37.6% and -10.6%. The δD values of individual alkenones (MeC_{37:2}; MeC_{37:3}; EtC_{38:2}; EtC_{38:3}) in suspended particles collected at different depths are shown in Table EA-4A along with the associated water δD values from surface sediments (0– 10 cm) are listed in Table EA-4B along with the salinity and water δD values from water depths of 2–5 m, where highest alkenone concentrations occurred (Table-EA-1).

 δD values of alkenones from particles and sediments were between $-207\%_{00}$ to $-182\%_{00}$ for MeC_{37:3}, $-185\%_{00}$ to $-165\%_{00}$ for MeC_{37:2}, $-221\%_{00}$ to $-196\%_{00}$ for EtC_{38:3}, and $-197\%_{00}$ to $-175\%_{00}$ for EtC_{38:2}. No significant difference in the δD value of individual alkenones was observed between surface sediments and particles in the overlying water column. Individual alkenones were depleted in deuterium relative to water by an amount that was lipid-specific but relatively constant throughout the Bay. That amount was $-177 \pm 2.9\%_{00}$ (n = 35) for MeC_{37:3}, $-156 \pm 3.2\%_{00}$ (n = 25) for MeC_{37:2}, $-188 \pm 3.2\%_{00}$ (n = 28) for EtC_{38:3}, and $-168 \pm 3.0\%_{0}$ (n = 30) for EtC_{38:2}. These isotopic fractionations are expressed as $\varepsilon_{alkenone-water}$ values ($\varepsilon_{alkenone-water} = (\alpha_{alkenone-water} - 1) = [(\delta D_{alkenone} + 1)/(\delta D_{water} + 1)$ - 1] in Tables EA-4A and EA-4B.

Particulate and sedimentary alkenone δD values were linearly correlated with water δD values with R^2 values of 0.8–0.9 in particles and 0.7–0.9 in sediments (Table 1, Fig. 5A). As salinity was linearly correlated with water δD values (Sachs and Schwab, 2011), alkenone δD values were linearly correlated with salinity with R^2 values of 0.6–0.8 in particles and 0.8 in sediments (Table 1, Fig. 5B). Linear regression analyses of $\delta D_{\rm alkenone} = \alpha_{\rm slope} \delta D_{\rm water} + \varepsilon_{\rm intercept}$ relationships indicated similar slopes ($\alpha_{\rm slope}$) of 0.703 ± 0.07 (n = 8) for each type of alkenone in both particles and sediments. But, intercepts ($\varepsilon_{\rm intercept}$) differed for each type of alkenone: $-178\%_{00}$ for

Equations of the relationships between individual alkenones and water δD values, individual alkenone δD values and salinity, and $\alpha_{alkenone-water}$ and salinity in sediment and water particle samples. Relationships between coeluting alkenones and water δD from others studies are shown for comparison.

Compound	Type of samples	Equation line δD alkenone versus δD water and salinity	R^2	n	$\alpha_{alkenone-water}-slope$	$\alpha_{alkenone-water} -$ intercept	Equation line $\alpha_{alkenone-water}$ versus salinity	R^2	n
MeC _{37:3}	Sediment	$\delta D_{\rm alk} = 0.793 \delta D_{\rm water} - 177.2^*$	0.79	16	0.793	0.823	$\alpha_{alkenone-water} = 0.0000 salinity + 0.824$	0.1	16
MeC _{37:3}	Filter	$\delta D_{\rm alk} = 0.710 \delta D_{\rm water} - 179.5^*$	0.85	19	0.710	0.821	$\alpha_{\text{alkenone-water}} = -0.0002 \text{salinity} + 0.827$	0.2	19
MeC _{37:2}	Sediment	$\delta D_{\rm alk} = 0.585 \delta D_{\rm water} - 164.4^*$	0.72	11	0.585	0.836	$\alpha_{alkenone-water} = -0.0004$ salinity $+ 0.849$	0.5	11
MeC _{37:2}	Filter	$\delta D_{\rm alk} = 0.701 \delta D_{\rm water} - 158.0^*$	0.87	16	0.701	0.842	$\alpha_{alkenone-water} = -0.0002 salinity + 0.848^*$	0.1	16
EtC _{38:3}	Sediment	$\delta D_{\rm alk} = 0.776 \delta D_{\rm water} - 190.6^*$	0.85	13	0.776	0.809	$\alpha_{alkenone-water} = -0.0001$ salinity $+ 0.812$	0.1	13
EtC _{38:3}	Filter	$\delta D_{\rm alk} = 0.618 \delta D_{\rm water} - 191.1^*$	0.77	15	0.618	0.809	$\alpha_{alkenone-water} = -0.0002 \text{salinity} + 0.817^*$	0.2	15
EtC _{38:2}	Sediment	$\delta D_{\rm alk} = 0.680 \delta D_{\rm water} - 172.5^*$	0.78	14	0.680	0.828	$\alpha_{alkenone-water} = -0.0003$ salinity $+ 0.835$	0.2	14
EtC _{38:2}	Filter	$\delta D_{\rm alk} = 0.761 \delta D_{\rm water} - 168.3^*$	0.83	16	0.761	0.832	$\alpha_{alkenone-water} = -0.0001 salinity + 0.834^*$	0.1	16
MeC _{37:3}	Sediment	$\delta D_{\rm alk} = 1.307 {\rm sal} - 218.3^*$	0.81	16	1.307	0.782			
MeC _{37:3}	Filter	$\delta D_{\rm alk} = 0.915 \text{sal} - 212.3^*$	0.63	19	0.915	0.788			
MeC _{37:2}	Sediment	$\delta D_{\rm alk} = 0.900 \text{sal} - 193.9^*$	0.85	11	0.900	0.806			
MeC _{37:2}	Filter	$\delta D_{\rm alk} = 0.958 \text{sal} - 192.0^*$	0.65	16	0.958	0.808			
EtC _{38:3}	Sediment	$\delta D_{\rm alk} = 1.121 {\rm sal} - 228.0^*$	0.78	13	1.121	0.772			
EtC _{38:3}	Filter	$\delta D_{\rm alk} = 0.911 {\rm sal} - 221.1^*$	0.78	15	0.911	0.779			
EtC _{38:2}	Sediment	$\delta D_{\rm alk} = 1.102 {\rm sal} - 207.3^*$	0.85	14	1.102	0.793			
EtC _{38:2}	Filter	$\delta D_{\rm alk} = 1.172 \text{sal} - 206.3^*$	0.83	16	1.172	0.794			
$MeC_{37:3} + MeC_{37:2}$		$\delta D_{\rm alk} = 0.732 \delta D_{\rm water} - 225.2$	0.99	5	0.732	0.775	From Englebrecht and Sachs (2005)		
$EtC_{38:3} + EtC_{38:2}$		$\delta D_{\text{alk}} = 0.745 \delta D_{\text{water}} - 232.8$	0.99	5	0.745	0.767	From Englebrecht and Sachs (2005)		
$MeC_{37:3} + MeC_{37:2}$		$\delta D_{\text{alk}} = 2.6 \delta D_{\text{water}} - 215$	0.85	11	2.600	0.785	From Schouten et al. (2006) (E.huleyi)		
$MeC_{37:3} + MeC_{37:2}$		$\delta D_{\rm alk} = 2.9 \delta D_{\rm water} - 242$	0.76	6	2.900	0.758	From Schouten et al. (2006) (G. oceanica)		

 $\alpha_{\text{alkenone-water}} - \text{intercept} = 1 - (-\epsilon/1000).$ * =equation in a confidence level of at least 95%.

MeC_{37:3}, -161_{00} for MeC_{37:2}, -191_{00} for EtC_{38:3}, and -170_{00} for EtC_{38:2}. Calculation of the fractionation factor using the intercept ($\alpha_{intercept} = 1 - (-\varepsilon_{intercept}/1)$) gave



Fig. 5. CB individual alkenones δD values in water particle (black dots) and sediment (yellow dots) samples plotted against water δD values (A) and salinity (B). Equation lines and R^2 of these relationships are listed in Table 1. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Table 2

D/H fractionation ratios between the different alkenones and water ($\alpha C_{3x:y-water}$) and between the different alkenones ($\alpha_{C3x:y-C3x:y}$) in water particles and sediments.

Ratios	Sample type	D/H fractionation	Number of samples
αC _{37·3-water}	Filter	0.823 ± 0.002	19
$\alpha C_{37:3-water}$	Sediment	0.823 ± 0.003	16
$\alpha C_{37\cdot 2-water}$	Filter	0.846 ± 0.002	16
$\alpha C_{37:2-water}$	Sediment	0.842 ± 0.003	11
$\alpha C_{38\cdot 3-water}$	Filter	0.814 ± 0.002	15
$\alpha C_{38:3-water}$	Sediment	0.810 ± 0.003	13
$\alpha C_{38:2-water}$	Filter	0.833 ± 0.002	16
$\alpha C_{38\cdot 2-water}$	Sediment	0.831 ± 0.003	14
α _{C37:3-C37:2}	Filter	0.973 ± 0.02	14
α _{C37·3-C37·2}	Sediment	0.974 ± 0.04	9
α _{C37·2} -C38·2	Filter	1.015 ± 0.03	12
α _{C37·2} -C38·2	Sediment	1.014 ± 0.08	10
α _{C38·3} _C38·2	Filter	0.977 ± 0.04	14
α _{C38·3-C38·2}	Sediment	0.969 ± 0.14	13
α _{C37·3-C38·3}	Filter	0.992 ± 0.08	15
α _{C37:3-C38:3}	Sediment	0.981 ± 0.09	10

 $\alpha C_{3x:y-water} = (1 + \delta D_{alk})/(1 + \delta D_{water})$ and $\alpha_{C3x:y-C3x:y} = (C_{3x:y} + 1)/(C_{3x:y+1})$ with x = 7 or 8 and y = 2 or 3.



Fig. 6. CB $\alpha_{alkenones-water}$ in water particle (black dots) and sediment (yellow dots) samples plotted against salinity (A) and temperature (B). Equation lines and R^2 of these relationships are listed in Table 1. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

values close to the fractionation factors derived from the slope (α_{slope}), with $\alpha_{intercept}$ on average 0.1 higher than α_{slope} (Table 1). The *D/H* fractionation between the different alkenones and water ($\alpha_{alkenone-water} = (1 + \delta D_{alk})/(1 + \delta D_{water})$) was nearly constant along the salinity gradient of the Bay (Table 2, Fig. 6A). Hydrogen isotopic offsets between diand tri-unsaturated C₃₇ and C₃₈ alkenones ($\alpha_{C37:2} - C_{37:3}$ and $\alpha_{C38:2} - C_{38:3}$, where $\alpha_{C37:x} - C_{38:x} = (C_{37:x} + 1)/(C_{37:x} + 1)$ with x = 2 or 3) and between C₃₇ and C₃₈ alkenones ($\alpha_{C37:2} - C_{38:2} = (C_{37:3} + 1)/(C_{37:2} + 1)$ with x = 7 or 8) were similar in all particulate and sediment samples collected in the CB (Table 2 and Electronic annex Tables EA-5A and EA-5B).

4. DISCUSSION

4.1. Alkenones and alkenone-producing haptophytes in the Chesapeake Bay

All particulate and sediment samples were characterized by low abundances of C_{38} methyl alkenones and the $C_{38:4}$ alkenone, implying minor contributions of alkenones from oceanic haptophyte species such as *E. huxleyi* and *G. oceanica* (Marlowe et al., 1984; Rontani et al., 2004) relative to those favoring fresh or brackish water (Marlowe et al., 1984; Volkman et al., 1995; Li et al., 1996; Thiel et al., 1997; Schulz et al., 2000; Sun et al., 2007). Similar distributions of alkenones have been reported in marginal and lacustrine environments characterized by uncalcified haptophytes such as C. lamellosa, I. galbana and Chromulina sp. (Cranwell, 1985; Volkman et al., 1988; Versteegh et al., 2001; Sun et al., 2007). The light microscopic and SEM-based identification of uncalcified haptophyte cells in our samples and reports of C. lamellosa and I. galbana in the Bay (Marshall, 1994; Marshall et al., 2005) support the notion that these species contributed substantially to the production of alkenones in our samples. The haptophyte bloom was reported to occur in late spring and to account for the bulk of the alkenone production during the year in the CB (Mercer et al., 2005). Therefore, alkenone indices in surface sediments likely represent long-term (i.e., 1-10 years, based on a linear sedimentation rate of $0.1-1 \text{ cm yr}^{-1}$ (Colman and Bratton, 2003; Cronin et al., 2010) averaged surface conditions occurring in late spring when runoff of D-depleted freshwater into the CB is high.

4.1.1. C_{37}/C_{38} and $%C_{37:4}$

The ratio of C_{37}/C_{38} alkenones in the CB was 1.2–4.5 in particles and 1.1-2.4 in sediments, averaging 2.2 (Tables EA-2 and EA-3). Open ocean ratios are typically between 0.5 and 1.5 (Conte et al., 1998; Chu et al., 2005), similar to those found in the most common marine alkenone producers, E. huxlevi (0.86-2.16, mean of 1.16 (Conte et al., 1994)) and G. oceanica (0.59-0.81, mean of 0.7 (Volkman et al., 1995)). Coastal haptophytes tend to have higher C₃₇/C₃₈ ratios, such as C. lamellosa (1.4-9.5, mean of 5.6 (Prahl et al., 1988)) and I. galbana (5.4-15.1, mean of 10.9 (Marlowe et al., 1984)), and values in excess of two have been reported in the Baltic Sea (Schulz et al., 2000) and some lacustrine environments (Cranwell, 1985: Thiel et al., 1997; Chu et al., 2005 and references therein) and attributed to alkenone production by species similar to C. lamellosa and I. galbana. We thus tentatively attribute average C_{37}/C_{38} ratios of 2.2 in the CB to substantial alkenone production by coastal haptophytes such as C. lamellosa and I. galbana, noting, however, that Theroux et al. (2010) suggested that C_{37}/C_{38} ratios better reflect algal growth status than species identity.

Relatively high concentrations of $C_{37:4}$ compared to $(C_{37:3} + C_{37:2})$ have been reported in low salinity marginal, coastal and estuarine environments (Rosell-Melé, 1998; Volkman et al., 1988; Schulz et al., 2000; Blanz et al., 2005; Mercer et al., 2005), while very high % $C_{37:4}$ values of up to 95% have been found in lacustrine sediments (Cranwell, 1985; Chu et al., 2005 and references therein). It was thus proposed that % $C_{37:4}$ values could be used as a paleosalinity or water mass proxy (Rosell-Melé, 1998; Schulz et al., 2000; Blanz et al., 2005). Furthermore, analysis of fossil DNA from sediments in hypersaline Ace Lake, Antarctica, showed that higher % $C_{37:4}$ values are correlated with higher relative abundance of freshwater versus oceanic haptophyte phylotypes (Coolen et al., 2004).

According to these studies, high values of both C_{37}/C_{38} and $%C_{37:4}$ are commonly associated with a predominance of freshwater and coastal haptophytes (Chu et al., 2005 and references therein). In the CB, a negative correlation between C_{37}/C_{38} and salinity (Fig. 3B) supports these observations. But a positive correlation between $%C_{37:4}$ and salinity (Fig. 3A) is inconsistent with these results. These opposite trends imply that physiological and/or species changes in alkenone producers along the salinity gradient in the CB differ from those in other settings. Furthermore, Toney et al. (2010) found a complete absence of $C_{37:4}$ in some lacustrine settings that was attributed to the presence of unidentified freshwater alkenone producers. Higher contributions of similar species in the upper reaches (i.e., low salinity region) of the CB may be an alternative explanation for the uncommon trend of $%C_{37:4}$ as a function of salinity.

4.1.2. $U_{37}^{K'}$ and U_{37}^{K} temperature indices

A vast amount of literature demonstrates the high fidelity with which the alkenone unsaturation ratio $U_{37}^{K'}$ co-varies with sea surface temperature (SST) in the ocean, and the suitability of a single global temperature calibration equation $(U_{37}^{K'} = 0.034T + 0.039)$ (see review by Herbert (2001)). When E. huxleyi and G. oceanica are not the predominant alkenone producers very different temperature calibrations have been reported (e.g., Rosell-Melé, 1998; Schulz et al., 2000; Chu et al., 2005; Mercer et al., 2005), making alkenone-based SST reconstructions uncertain when the identity of alkenone producers is unknown. Reconstructed SSTs in the CB are closer to measured values when temperature calibrations for I. galbana (Versteegh et al., 2001) and C. lamellose (Sun et al., 2007) are used than when marine (Prahl et al., 1988) calibrations are used. This provides further evidence that these coastal species may predominate the assemblage of alkenone-producing haptophytes in the CB. The absence of correlation between U_{37}^{K} or $U_{37}^{K'}$ and SSTs (e.g. Fig. 4) may be due to variations in the growth phase or the assemblage of alkenone producers along the CB (see discussion below), both of which are known to influence U_{37}^{K} and $U_{37}^{K'}$ (e.g., Conte et al., 1994; Mercer et al., 2005; Toney et al., 2010), or alternatively to the small range of temperature observed during sampling.

4.2. CB alkenone δD values

The CB estuary provides an ideal setting to evaluate (i) the fidelity with which alkenone lipids record the hydrogen isotopic composition of water and (ii) the influence of salinity on D/H fractionation in those lipids, because ample alkenone concentrations exist for hydrogen isotope analyses over a ~200 km estuary in water spanning a δD range of 27‰ (-38‰ to -11‰) and a salinity range of 19 (10.0–28.8). Below we first discuss the relationship between water δD values and δD values in four different alkenones from CB particles and surface sediments, demonstrating that these lipids faithfully track the isotopic composition of water. A discussion of the invariant D/H fractionation between water and alkenones along the salinity gradient of 19 follows.

4.2.1. Relationship between alkenone and water δD values

Our sampling in late May was timed to occur during the maximum of the annual alkenone production that occurs in late spring when runoff of D-depleted freshwater into the CB is at a maximum (Mercer et al., 2005). Therefore, the similitude of alkenone δD values in particulate and sediment samples suggest that alkenone δD values in sediment record long-term (i.e., 1–10 years, based on a linear sedimentation rate of 0.1–1 cm yr⁻¹ (Colman and Bratton, 2003; Cronin et al., 2010) average surface D-depleted water by late spring freshwater runoff.

Different intercept (ε) but similar slope (α) values for each of the four alkenones in the equation δD_{alke-} $_{none} = \alpha \delta D_{water} + \varepsilon$ indicate that, while different for each of the four alkenones, the magnitude of lipid-water D/Hfractionation for each remains constant along the length of the CB (Table 1). Furthermore, previous culture experiments of the oceanic haptophytes E. huxlevi and G. ocea*nica* showed higher alkenone-water D/H fractionation of 233 ± 6 for C_{37:2 + 37:3} in Englebrecht and Sachs (2005), 220 ± 16 for C_{37:2 + 37:3} in Schouten et al. (2006), 198 ± 20 for $C_{37:3}$ and 180 ± 16 for $C_{37:2}$ in Wolhowe et al. (2009) than reported in this study in which we find $\epsilon_{C37:3-water}$ of 180 ± 4 and $\epsilon_{C37:2-water}$ of 158 ± 3 in water particles (Table 1). As coastal haptophytes are likely predominant in our samples, lower alkenone-water D/H fractionation (by up to 60% relative to Englebrecht and Sachs (2005)) suggests that coastal haptophytes may fractionate less than oceanic haptophytes.

4.2.2. Lack of co-variation between salinity and D/H fractionation in CB alkenones

The D/H fractionation between water and all four alkenones was invariant over the salinity range of 10-29 (Fig. 6A and Table 2). This result is at odds with three recent studies indicating a 1-3% decrease in D/H fractionation in lipids per unit increase in salinity (Schouten et al., 2006; Sachse and Sachs, 2008; Sachs and Schwab, 2011). In batch-cultured E. huxleyi and G. oceanica Schouten et al. (2006) observed a positive linear correlation between salinity and $\alpha_{alkenone-water}$, the slope of which implied a decrease in D/H fractionation in $C_{37:3} + C_{37:2}$ alkenones of 3.3% and 3.0%, respectively, per unit increase of salinity over the salinity range 25-35. In the same CB sample set on which we report here we observed a 0.99% decrease in D/H fractionation in dinosterol (4 α ,23,24trimethyl-5α-cholest-22E-en-3β-ol) per unit increase of salinity over the salinity range 10-29 (Sachs and Schwab, 2011). And in hypersaline ponds on Christmas Island in the central equatorial Pacific (Republic of Kiribati), a 0.8-1.1% decrease in D/H fractionation per unit increase in salinity was observed in phytene, n-C₁₇ alkane and a mixture of total extractable lipids over the salinity range 35-150 (Sachse and Sachs, 2008). While the 3-4-fold higher sensitivity of D/H fractionation to salinity in Schouten et al. (2006)'s alkenones as compared to these other studies was likely due in (large) part to the associated changes in growth rate, there is strong evidence that D/H fractionation decreases in algal lipids in response to higher salinity.

We consider three hypotheses to explain the lack of covariation between $\alpha_{alkenone-water}$ and salinity in the CB. (i) Growth rate and temperature changes cause D/H fractionation in alkenones that counteract the fractionation arising from salinity changes. (ii) The assemblage of alkenoneproducing haptophytes changes along the Bay and each species has a different sensitivity to salinity, such that no apparent trend in $\alpha_{alkenone-water}$ occurs along the salinity gradient. (iii) Coastal haptophytes that likely predominate the assemblage of alkenone producers in the CB may have an enhanced osmoregulation capacity that circumvents the mechanisms that cause a change in alkenone-water D/H fractionation in response to salinity changes in other species of phytoplankton and cyanobacteria.

- (i) Growth rate and temperature changes cause D/Hfractionation in alkenones that counteract the fractionation arising from salinity changes. Culture and field studies have demonstrated that growth rate and temperature can influence D/H fractionation in lipids (Schouten et al., 2006; Wolhowe et al., 2009; Zhang et al., 2009), but the data are few and the results contradictory.Zhang et al. (2009) showed that acetogenic lipids (C14; C16:1; C16; C18 fatty acids) from two species of freshwater green algae (E. unicocca and V. aureus) grown in batch culture at 15 °C were enriched in deuterium by 20-40% relative to those grown at 25 °C. Similarly, Wolhowe et al. (2009) observed that individual alkenones from batch-cultured E. huxleyi and G. oceanica at 17 °C were enriched in deuterium by 40-50% relative to those grown at 25 °C. However, Schouten et al. (2006) observed no significant effect of temperature on alkenone-water D/H fractionation in batch-cultured E. huxlevi and G. oceanica, but growth rate changed along with temperature in these experiments. In the CB, the lack of any trend between alkenone-water D/H fractionation and temperature also suggests that temperature was not a first order influence on individual alkenone δD values (Fig. 6B). Growth rate has also been shown to influence D/Hfractionation. Schouten et al. (2006) observed a substantial increase in D/H fractionation as growth rate increased in batch-cultured E. huxlevi and G. oceanica in response to different temperatures and salinities. In continuous culture experiments (i.e., chemostats), Zhang et al. (2009) found that D/H fractionation also increased substantially as growth rate increased in a sterol from the diatom T. pseudonana, but little or not at all in fatty acids from the same cultures. Growth rate trends from other phytoplankton groups in the CB indicate highest values in the mid-Bay and lowest values near the mouth of the Bay (Xu and Hood, 2006), opposite to those required to counteract the expected decrease of alkenone-water D/H fractionation as salinity increases along the Bay.
- (ii) The assemblage of alkenone-producing haptophytes changes along the Bay and each species has a different sensitivity to salinity, such that no apparent trend in α_{alkenone-water} occurs along the salinity gradient. Different species of alkenone producers and other phytoplankton are known to produce the same lipids with different δD values when grown in water with the same salinity (cf., Schouten et al., 2006) or in water with the same δD value (e.g., Zhang and Sachs,

2007). Schouten et al. (2006) demonstrated that D/H fractionation in alkenones from *E. huxleyi* and *G. oceanica* grown in water with the same salinity differed by about 30%. Though we know of no study reporting D/H fractionation in alkenones from coastal species such as *C. lamellosa* and *I. galbana*, it seems reasonable to assume that they too fractionate hydrogen differently, and consequently that a change in the assemblage of alkenone producing haptophytes along the Bay influenced $\alpha_{alkenone-water}$.

(iii) Coastal haptophytes that likely predominate the assemblage of alkenone producers in the CB may have an enhanced osmoregulation capacity that circumvents the mechanisms that cause a change in alkenone-water D/H fractionation in response to salinity changes in other species of phytoplankton and cyanobacteria. The observation that D/H fractionation in dinosterol of the CB dinoflagellates diminished as salinity increased (Sachs and Schwab, 2011), but remained constant in alkenones of the CB haptophytes, suggests that the effect of salinity on D/H fractionation in lipids is species dependant. Increasing salinity is particularly challenging for phytoplankton since lower water potential occurring during hyperosmotic stress reduces external water availability and increases harmful ion concentrations in the cell (Epstein, 1980). Sachse and Sachs (2008) and Sachs and Schwab (2011) proposed that algal lipid D-enrichment with increasing salinity results from the progressive D-enrichment of internal cell water as H is preferentially drawn from the internal water pool for biosynthetic reactions, and the flux of external water into the cell slows. Furthermore, since enzyme activities are affected by elevated cytoplasmic ion concentrations (Niu et al., 1995; Kirst, 1989) and ultimately control lipid δD values (Baillif et al., 2009; Chikaraishi et al., 2009; Schwab and Sachs, 2009), it is likely that elevated cytoplasmic ion concentrations occurring during hyperosmotic stress may hamper vital enzymes involved in lipid synthesis and their resultant δD values. As both cellular internal water volume and cytoplasmic ion concentration affect water potential during hyperosmotic stress (Haines, 1994; Jahnke and White, 2003), lipid δD values are expected to be affected by the capacity of algae to reduce water loss and internal concentrations of incompatible ions (i.e., osmoregulate). Conversely, at low external salinities the flux of external water into the cell increases and deleterious ion concentrations in the cytoplasm decrease. For instance, in the marine alga, Dunaliella tertiolecta, a high flux of external water into the cell under extreme hypoosmotic stress caused a larger cell volume and a higher concentration of α -tocopherol, indicative of heightened stress within the chloroplast membrane (Jahnke and White, 2003). Under this condition, the D/Hratio of the internal cell water is likely to be similar to the D/H ratio of the external water, resulting in lipid δD values that are more likely to reflect those of the environmental water. In CB, such hypo- and

hyper-osmotic stress was probably not so intense, but the mechanisms that drove water flux into and out of the cell were likely similar.

4.3. Paleoenvironmental implications

The difference in sensitivity of D/H fractionation to salinity in CB haptophytes versus CB dinoflagellates indicates that salinity reconstructions based on paleo-lipid δD values may be species dependant. Some species may synthesize lipids with D/H ratios that are sensitive to salinity variations, others may synthesize lipids with δD values that are independent of salinity. The latter are likely to be good targets for water isotopic reconstructions.

Furthermore, the consistency of individual D/H fractionation factors between alkenones and water ($\alpha_{alkenone-water}$) in suspended particles and in sediment of the CB, and the similarity between slope- and intercept-derived fractionation factors (Table 2), suggest that water δD values are the primary control on changing alkenone δD values along the Bay and thus, CB water δD values may be determined using individual alkenone δD values with the equations listed in Table 1.

However, constant hydrogen isotope offsets between alkenones with different chain lengths but the same degree of unsaturation ($\alpha_{C37:2-C38:2}$ and $\alpha_{C37:3-C38:3} \approx 1.01$) and between those with the same chain length but different deof unsaturation grees and $(\alpha_{C37:3} - C37:2)$ $\alpha_{C38:3 - C38:2} \approx 0.97$) suggest that single, well-defined values represent hydrogen isotope fractionation associated with enzymatic dehydrogenation and carboxylation reactions during alkenone biosynthesis in the CB (Rontani et al., 2006; Chikaraishi et al., 2009; Schwab and Sachs, 2009). This observation strengthens the argument of Wolhowe et al. (2009) that these values may be used in isotopic mass balance calculations to determine δD values for individual alkenones without the labor intensive LC-MS separation developed by Schwab and Sachs (2009) and used in this study. However, further evaluation of the universality of the isotopic offset between different alkenones ought to be conducted in the lab and field before extending it to other coastal environments and corresponding species, since different salinity effects on $\alpha_{alkenone-water}$ in oceanic and coastal species suggest that physiological or morphological differences between alkenone producers may greatly influence alkenone δD values.

5. CONCLUSION

Alkenone distributions characterized by a high abundance of $C_{37:4}$ and high C_{37}/C_{38} ratios, in addition to the occurrence of uncalcified cells, support substantial contributions of coastal haptophyte species, such as *C. lamellosa* and *I. galbana*, in all studied surface sediment and suspended particle samples in the CB estuary.

The δD values of individual alkenones (MeC_{37:2}; MeC_{37:3}; EtC_{38:2}; EtC_{38:3}) closely co-varied with water δD values. Constant hydrogen isotope offsets ($\alpha_{C37:2 - C38:2}$ and $\alpha_{C37:3 - C38:3} = 1.01$ and $\alpha_{C37:2 - C37:3}$ and $\alpha_{C38:2 - C38} =$

0.97) were observed between the different alkenones in all studied samples along the Bay. This value was used in an isotopic mass balance calculation to determine δD values for individual alkenones in the CB when a mixture of two alkenones is analyzed (e.g., C_{37:2} + C_{37:3})

Slope- and intercept- derived fractionation factors of the correlations ($\delta D_{alkenone} = \alpha_{alkenones-water} \delta D_{water} + \varepsilon$) describing the relationship between water and individual alkenone δD values did not differ significantly. No change in D/H fractionation ($\alpha_{alkenone-water}$) was observed along the salinity gradient of the Bay, suggesting that individual alkenone D/H ratios in the CB were primarily determined by source water δD values and that they may be used as proxy to reconstruct CB water δD values.

The difference in sensitivity of D/H fractionation to salinity in haptophytes relative to dinoflagellates in the CB indicates that the mechanisms of hydrogen isotope fractionation in different phytoplankton are influenced to a varying extent by salinity. By extension, paleosalinity reconstructions from lipid δD values are expected to be species dependant. Two valid hypotheses we consider to explain the lack of a salinity influence on alkenone-water D/H fractionation in the CB are: (i) the assemblage of alkenone-producing haptophytes changes along the Bay and each species has a different sensitivity to salinity, such that no apparent trend in $\alpha_{alkenone-water}$ occurs along the salinity gradient, and (ii) greater osmoregulation capacity in coastal haptophytes may result in little or no sensitivity of the D/Hfractionation process during alkenone synthesis to salinity changes. Laboratory culture studies with oceanic, coastal and lacustrine alkenone-producing prymnesiophytes ought to provide a test of these hypotheses.

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APPENDIX A. SUPPLEMENTARY DATA

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.gca.2011.09.031.

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