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### GEOCHEMICAL CHARACTERIZATION OF ORGANIC MATTER IN VICTORIA HARBOUR SEDIMENTS, HONG KONG

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By

MICHAEL HSIEH Norman, Oklahoma 2006 UMI Number: 3207060

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### GEOCHEMICAL CHARACTERIZATION OF ORGANIC MATTER IN VICTORIA HARBOUR SEDIMENTS, HONG KONG

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### ABSTRACT

Victoria Harbour is located between Kowloon and Hong Kong Island in the southeastern prodelta region of the Pearl River system. Since the mid-1900s, the population in Hong Kong has grown rapidly, coastal areas surrounding Victoria Harbour have been reclaimed, and excess raw sewage has been disposed into the Harbour. Release of methane from harbour sediments during dredging activities instigated interest in studying the sources of methane trapped in Victoria Harbour sediments. Core MBH 54/2, from a heavily polluted area in Victoria Harbour's Kowloon Bay, was selected for this study. However, no methane was detected in sediments from this core. The project was redefined, using a detailed organic geochemical characterization approach to determine the sources of organic matter, evaluate changes in environmental conditions, and to ascertain whether remnants of bacterial lipids might be present to enhance our understanding of processes contributing to methane generation. Bulk properties (e.g., %C<sub>org</sub>, %N,  $\delta^{13}C_{org}$ , and  $\delta^{15}N$ ), lipid composition and profiles were applied to delineate changes in organic matter sources deposited in the Kowloon Bay area during the late Quaternary.

The organic carbon-to-nitrogen ratio demonstrated fluctuations in the sources of organic matter throughout the Holocene unit of MBH 54/2. High fluxes in the carbon-to-nitrogen ratio may reflect strong storms, where excess terrigenous plant material is transported into the area. Sediment intervals impacted by sewage waste had isotopic compositions (*i.e.*,  $\delta^{13}C_{org}$  and  $\delta^{15}N$ )

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consistent with those reported in the literature for sewage in coastal environments.

Sources of organic matter could be differentiated using free lipids, which consisted of sterols, n-alcohols, fatty acids, and n-alkanes. Environmental conditions in Kowloon Bay were inferred to be anoxic based on the relative abundance of stanols-to-sterols. Sewage contaminated sediments were confirmed by the presence of fecal sterols. Periods of improved environmental conditions were noted by the occurrence of sterols common to aquatic organisms. Bound lipids appear to retain lipid profiles descriptive of bacterial communities in the sediments. More in-depth comparisons to lipid profiles of bacterial strains might allow bacterial remnants in sediments to be identified, allowing us to better speculate on their role in the remineralization of organic matter in Recent sediments.

#### CHAPTER 1

### Introduction

### 1.1 Project Rationale and Objectives

Victoria Harbour is situated within the southeastern prodelta region of the Pearl River system (Fyfe *et al.*, 1999), between Kowloon and Hong Kong Island, in the Hong Kong Special Administrative Region (SAR) of China. A tidal channel runs west to east, through Victoria Harbour, with the mouth of the Pearl River to the west and the northern continental shelf of the South China Sea to the east (**Fig. 1.1**; Fyfe *et al.*, 1997; Yim *et al.*, 2002). The Pearl River system has played an important role in supplying sediments deposited in Victoria Harbour during the Quaternary (Chalmers, 1984; Davis, 1999; Fyfe *et al.*, 1999). Sediment transport in the harbour is controlled by tidal currents, with summer/autumn typhoons and winter/summer monsoons playing important roles in resuspending and redistributing sediments throughout the harbour (Huang and Yim, 1997; Yim *et al.*, 2002).

This region is of particular interest in that the Holocene unit of the innercontinental shelf in the Hong Kong SAR has been proposed to be a net carbon sink (Yim *et al.*, 2002). Continental margins, especially in regions in close proximity to deltas, are typically important reservoirs of sedimentary organic matter (Berner, 1989; Hedges, 1992; Pernetta and Milliman, 1995; Hedges *et al.*, 1997; Mudge and Norris, 1997). In studies of the global carbon cycle, the ocean has been identified as preserving approximately one-third of the total organic



**Fig. 1.1.** Map illustrating the location of Victoria Harbour, relative to the Pearl River, in the Hong Kong Special Administrative Region of China (map taken from Fyfe *et al.*, 2000).

carbon inventory (Hedges *et al.*, 1997). An estimated 80% of the organic carbon in the ocean is buried and preserved in deltaic and continental shelf environments (Berner, 1989; Hedges, 1992; Hedges and Oades, 1997; Mudge and Norris, 1997). Factors controlling the preservation of organic carbon in marine sediments have been debated, with primary arguments being between anoxia versus productivity. Anoxia focuses on the idea that organic carbon is less efficiently degraded under anaerobic conditions compared to aerobic conditions. Whereas those supporting productivity argue that conditions favoring the growth of organisms (*e.g.*, areas of coastal upwelling) are more important for the accumulation of organic matter (Demaison and Moore, 1980; Pedersen and Calvert, 1990; Calvert and Pedersen, 1993; Canfield, 1994). Additional parameters that have also been considered are the adsorption of organic matter to mineral surfaces and high sedimentation rates (Müller and Suess, 1979; Keil *et al.*, 1994; Rullkötter, 2000). It has been suggested that organic matter bound to mineral matrices is better protected from microbial attack and chemical alteration (Kawamura and Ishiwatari, 1984; Keil *et al.*, 1994). The thought behind better preservation of organic matter due to high sedimentation rate is that the organic matter will have a shorter residence time in the water column (where remineralization of organic matter to  $CO_2$  typically occurs) and will be rapidly buried in bottom sediments (Didyk *et al.*, 1978). At the same time, high sedimentation rates can also result in the dilution of organic matter due to the deposition of clastic material (Rullkötter, 2000).

The present-day Victoria Harbour is a unique environment that has undergone many human-induced changes. Historical records indicate that the mid-1800s marked the beginning of reclamation activities and large-scale sewage discharge into Victoria Harbour. In the mid-1900s, the human population grew rapidly (**Table 1.1**) in Hong Kong, resulting in an increase of raw sewage and wastewater effluents being discharged into the harbour. This also led to the need for more land area resulting in large-scale dredging and coastal reclamation activities (Chalmers, 1984; Yim, 1984; Connell *et al.*, 1998; Lee and Liu, 1999; Tanner *et al.*, 2000; Yim *et al.*, 2002). During dredging of Victoria Harbour sediments, methane was observed escaping from the sediments to the surface

Year	Population	
1841 <sup>1</sup>	7,500	
1931 <sup>1</sup>	849,800	
1945 <sup>1</sup>	750,000	
1991 <sup>2,3,4</sup>	5.6 x 10 <sup>6</sup>	
1996 <sup>3,4</sup>	6.3 x 10 <sup>6</sup>	
1999 <sup>5</sup>	6.6 x 10 <sup>6</sup>	
2001 <sup>3,4</sup>	6.7 x 10 <sup>6</sup>	
2003 <sup>5</sup>	6.8 x 10 <sup>6</sup>	
2004 <sup>5</sup>	6.9 x 10 <sup>6</sup>	

**Table 1.1.** Estimated population in Hong Kong, 1841-2004.

<sup>1</sup>http://www.answers.com/topic/demographics-of-hong-kong

<sup>2</sup>http://www.china-tour.cn/cityguides/hongkong.htm <sup>3</sup>http://www.hk.cc.og.hk/eng/winter%202001/population%20Census.htm

<sup>4</sup>http://www.jil.go.jp/foreign/event\_r/event/documents/2004-sopemi\_e\_countryreport3.pdf

<sup>5</sup>http://www.info.gov.hk/censtatd/eng/hkstat/hkinf/population\_index.html

raising concerns in this area for its potential impacts as a greenhouse gas (Yim et al., 2002).

The increased sewage input has resulted in accelerated eutrophication of Victoria Harbour. Reclamation activities have not only reshaped the harbour (Figs. 1.2 and 1.3), but have also increased sedimentation rates in various parts of the harbour (Tanner et al., 2000; Yim et al., 2002). Based on the current amount of organic matter input, anoxic bottom waters, and high sedimentation rate, it would seem that marine sediments in Victoria Harbour should be wellsuited for the deposition and preservation of organic carbon (Didyk et al., 1978). Prior to human activities, the Pearl River, tidal currents, tropical storms and monsoons, and eustatic sea level changes were the likely factors controlling organic matter deposition in Victoria Harbour.



**Fig. 1.2.** Maps illustrating changes to the land area surrounding Victoria Harbour, 1903 to 1980, as a result of coastal reclamation (from Chalmers, 1984).



**Fig. 1.3.** Coastal reclamation history in Victoria Harbour (modified from Yim 2000). MBH 54/2 refers to the piston core used in this study. Sedimentation rates, based on <sup>210</sup>Pb data from MVC 74 (Tanner *et al.*, 2000), were used to estimate sediment ages in core MBH 54/2.

Lipid composition, elemental and isotopic measurements of organic matter at various depth intervals from a core in the inner-continental shelf region of Hong Kong have been used to reconstruct environmental changes and to speculate on the history of this region. The major objectives of this project were to: (1) characterize the various classes of lipids (*i.e.*, free-, ester-bound, and amide-bound lipids); (2) ascertain sources and changes in organic matter deposited and preserved during the late Quaternary in Victoria Harbour; (3) use elemental and bulk stable isotope compositions of carbon and nitrogen in sedimentary organic matter to infer changes in organic matter sources, to speculate on possible early diagenetic processes, and to identify periods of environmental change (*e.g.*, the interglacial-glacial boundary); (4) apply compound-specific carbon isotopes to differentiate sources of lipids in the core samples.

#### 1.2 Study Area – Victoria Harbour, Hong Kong SAR

Victoria Harbour is located in the Hong Kong Special Administrative Region (SAR) of China, between Kowloon and Hong Kong Island, and is one of the busiest shipping ports in the world. The total area within the Hong Kong territorial boundaries is about 3400 km<sup>2</sup>. Land coverage in this region, which is comprised of the New Territories, Kowloon, Hong Kong Island, Lantau Island, and other surrounding islands, totals an area of about 1100 km<sup>2</sup> (**Fig. 1.1**; Yim *et al.*, 2002). Victoria Harbour is in the southeastern prodelta region of the Pearl

River and has a tidal channel running west to east through the harbour into the northern continental shelf of the South China Sea (Fyfe *et al.*, 1997). Nearly all land surrounding Victoria Harbour has been reclaimed (**Fig. 1.3**).

### 1.2.1 Hong Kong – Late Pleistocene to Holocene

During the late Pleistocene, around the last glacial maximum (~25,000 years B.P.), the coastline along southern China was about 130 km south of Hong Kong (**Fig. 1.4**; Feng and Shi, 1997; Owen *et al.*, 1998). Shallow seismic profiles (Feng and Shi, 1997) unveiled buried ancient river channels demonstrating that the Pearl River palaeodelta once extended over a significant area on the continental shelf in the South China Sea (**Fig. 1.4**). Fyfe *et al.* (2000) and Owen (2005) have reported the occurrence of low sinuous braided river channels in the Hong Kong area, during the late Pleistocene (**Fig. 1.5**). Coarser grained sands were deposited in this area during this period of low sea level (Fyfe *et al.*, 2000).

After about 18,000 years B.P. sea level began rising, reaching at least -19.5m by around 8,080 years B.P. (Owen *et al.*, 1998; Owen, 2005). The rise in sea level resulted in a blanket of intertidal silty mud deposited over this area. By about 6,000 years B.P., the coastline along the southern shores of China extended as far north as Guangzhou, whereas the coastline surrounding Hong Kong was similar to what is seen in the present day (**Fig. 1.1**; Fyfe *et al.*, 1997; Owen *et al.*, 1998; Davis, 1999). Between 6,000 years B.P. and the present day,



**Fig. 1.4.** Ancient river channels and delta plain, extending out into the continental shelf of the South China Sea, during the late Pleistocene (from Feng and Shi, 1997).



**Fig. 1.5.** Reconstructed environment around Hong Kong during the late Pleistocene (from Fyfe *et al.*, 2000).

sea level fell slightly resulting in the development of the current Pearl River delta system (**Fig. 1.1**; Fyfe *et al.*, 1999).

The rise and fall in sea level throughout the Quaternary, in Hong Kong, has resulted in the deposition of alternating units of marine and terrestrial deposits (Yim, 1984; Yim, 1994; Owen et al., 1998; Davis, 1999). The alternating units of marine sediments versus terrestrial sediments have been recognized by Yim (1994), based on selected features (*i.e.*, palaeontology, sedimentology, mineralogy, chemistry, and engineering properties). Using these parameters, Yim (1994) classified Quaternary sediments in Hong Kong as alternating units of marine and terrestrial sediments (denoted as "M" for marine, "T" for terrestrial, and numbered from 1 to 5). The youngest marine unit, "M1," has a maximum age of 8,100 years B.P. and is comprised of soft green, gray, and/or black colored silty clay, with abundant shell remnants (Yim, 1984 and 1994). Throughout much of Hong Kong, the "T1" unit is missing. The "T1" unit represents terrestrial sediments of the last glacial period (8,100 to 70,000 years B.P.), deposited during a time of low sea level (Yim, 1994; Davis, 1999). It has been suggested that sediments of the T1 unit are missing because either they were never deposited, or during the lowstand the water levels were still high enough not to expose the sediments (Davis, 1999). The top of the "M2" unit, the marine unit of the last interglacial period (90,000 to 140,000 years B.P.), has been identified by the presence of a desiccated crust. The desiccated crust refers to marine sediments that have been subaerially exposed during periods of low sea level. Sediments that have undergone desiccation, in pre-Holocene marine sediments,

will have a mottled appearance (*i.e.*, a mixture of white, yellow, orange, red, and brown colors) (Yim, 1994; Tovey and Yim, 2002).

# 1.2.2 Sedimentation Rates (Based on <sup>210</sup>Pb-Dating) and Approximate Age of Sediments

Sedimentation rates in the Kwun Tong Typhoon Shelter area have been measured by Tanner *et al.* (2000), using <sup>210</sup>Pb-dating, for core MVC 74 (**Fig 1.3**). <sup>210</sup>Pb-dating is a technique commonly used for estimating sedimentation rates based on the radioactive decay of <sup>210</sup>Pb, where the half-life ( $t_{\frac{1}{2}}$ ) is about 22.3y (Geyh and Schleicher, 1990). <sup>210</sup>Pb is a naturally occurring radionuclide which belongs to the <sup>238</sup>U decay series (see illustration below), and is produced in both the atmosphere and terrestrial environments.

 $^{238}U \xrightarrow{4.51 \times 10^9 y} \xrightarrow{226} \text{Ra} \xrightarrow{1602y} \xrightarrow{222} \text{Rn} \xrightarrow{3.8d} \xrightarrow{210} \text{Pb} \xrightarrow{22.3y} \xrightarrow{210} \text{Po} \xrightarrow{138.4d} \xrightarrow{206} \text{Pb}$ 

Radionuclides formed in the <sup>238</sup>U decay series (from Appleby, 2001).

Atmospheric <sup>210</sup>Pb originates from <sup>222</sup>Rn, a radioactive gas which diffuses through the subsurface into the atmosphere. <sup>222</sup>Rn has a short half-life ( $t_{\frac{1}{2}}$ =3.8d) and decays to <sup>210</sup>Pb, which then easily binds to particulate material and is returned to sediments by dry deposition or rain. The <sup>210</sup>Pb is believed to be immobile once redeposited, and undergoes further decay. Terrestrially derived <sup>210</sup>Pb refers to <sup>210</sup>Pb occurring in the sampled sediment intervals where <sup>222</sup>Rn undergoes *in situ* radioactive decay. <sup>210</sup>Pb-dating is calibrated using <sup>137</sup>Cs, an artificial radionuclide produced from the atmospheric testing of nuclear bombs. Peak deposition of <sup>137</sup>Cs in sediments occurred in 1964 (Noller, 2000; Appleby, 2001). Sedimentation rates, using <sup>210</sup>Pb, of recent sediments from lacustrine and marine environments have been estimated to be reliable between 5 to 150 years (Geyh and Schleicher, 1990; Noller, 2000).

The present-day sedimentation rate in Kwun Tong typhoon shelter (from the seafloor surface to 0.5m depth) was determined to be 3.5cm y<sup>-1</sup>. The mean sedimentation rate for 0.5m to 1.5m was estimated to be about 4.4cm y<sup>-1</sup> and corresponds to calendar years spanning 1957 to 1980. At depths of 1.5m to 2.1m, the sedimentation rate was estimated to be 1.9cm y<sup>-1</sup>, representing sediments deposited between 1928 and 1957. Sedimentation rates at depths greater than 2.1m could not be determined due to uncertainties with excess <sup>210</sup>Pb activity (Tanner *et al.*, 2000). The maximum Holocene age has been reported to be about 8,100y. The base of the Holocene unit is marked by a desiccated crust, which represents the boundary between the M1 and M2 units (Yim, 1994). If the base of the Holocene unit occurs at 3.7m, and the maximum Holocene age is 8,100y, then the average sedimentation rate between 2.1m and 3.7m would be about 0.2mm y<sup>-1</sup>.

### 1.2.3 Tropical Cyclones (Typhoons) in Hong Kong

Hong Kong is located on the northernmost region of the South China Sea and lies within the pathway commonly traversed by typhoons (Huang and Yim, 1997). Historical pathways of typhoons (also referred to as tropical cyclones) that have passed through Hong Kong (between 1957 and 1999), with wind speeds of at least 118km/hr, are summarized in **Fig. 1.6** (http://www.hko.gov.hk/informtc/ historical\_tc/no10track.htm). In general, the majority of typhoons tracked around the Hong Kong region approach Hong Kong from the southeast, and continues along a northwestern pathway (Huang and Yim, 1997).



**Fig. 1.6.** Map illustrating the pathways of typhoons, with wind speeds of at least 118 km/hr, that passed over Hong Kong between 1957 and 1999 (map taken from http://www.hko.gov.hk/ informtc/historical\_tc/no10track.htm).

### 1.2.4 Sewage Dumping in Victoria Harbour

The disposal of sewage waste into Hong Kong waters occurs on a rather large scale, primarily due to the immense population (~6.9 million people in 2004), and because existing sanitary landfill sites and sewage treatment plants are incapable of handling such large amounts of waste (Yim, 1984; World Wide Fund for Nature Hong Kong, 1993). In general, raw sewage is released into Hong Kong waters by seawall-type sewage outfalls (Fig. 1.7) and submarine-type sewage outfalls (Fig. 1.8), with minor or no treatment (Yim, 1984). About fifty percent of the raw sewage is released directly into Hong Kong waters. Of the remaining fifty percent of incoming sewage, about forty percent of larger size solid waste undergoes sedimentation (*i.e.*, "preliminary treatment"), and the remaining ten percent undergoes some type of further treatment (Wong and Tanner, 1997; World Wide Fund for Nature Hong Kong, 1993). Sometime after the mid-1970s, several of the seawall-type sewage outfalls were converted to submarine-type sewage outfalls and diverted further into the channel of Victoria Harbour. The goal was to dilute and better disperse sewage in Victoria Harbour (Yim, 1984). In a 1981 report, the two districts in Hong Kong generating and discharging the largest amount of sewage wastes were: Kwun Tong (~221,000 m<sup>3</sup>/day; via seawall-type sewage outfall) and Tsuen Wan/Kwai Chun (~243,000 m<sup>3</sup>/day; via submarine-type sewage outfall) (Yim, 1984). The total estimated raw sewage discharged throughout Victoria Harbour has been estimated to be at least 1.6 x 10<sup>6</sup> m<sup>3</sup>/day (Yim *et al.*, 2002). For comparison, the estimated daily sewage received by the waste water treatment facility in Norman, Oklahoma, is

about 38,611 m<sup>3</sup>/day, servicing a population of about 92,400 people (personal communications with Ralph Arnett, Darrell Schwartz, and Mark Daniels, from the Norman Waste Water Treatment Facility).



**Fig. 1.7.** Location of sewage outfalls (**\***) in Hong Kong, up to 1981 (modified from Yim, 1984 and 1993).


**Fig. 1.8.** Submarine-type sewage outfalls in Victoria Harbour, Hong Kong (from Yim, 1984).

# **1.3 Summary Remarks**

The area where the modern-day Victoria Harbour is located has undergone many changes from the late Pleistocene through the Holocene. The coastline along the southern regions of China was about 130km south of Hong Kong during the late Pleistocene (during the last glacial max). During this period, ancient river channels ran throughout the area, and a palaeodelta of the Pearl River extended over a significant region on the continental shelf in the South China Sea. Sea level began rising after ~18,000 years B.P., reaching at least -19.5m by ~8,080 years B.P. The coastline reached as far north as Guangzhou by about 6,000 years B.P., then receded slightly with the fall in sea level. With the rise in sea level, a blanket of Holocene intertidal silty mud was deposited throughout the area surrounding Hong Kong. Alternating layers of marine and terrestrial sediments, resulting from changes in sea level, can be observed in the Hong Kong marine sediments.

In more recent times, rapid population growth within Hong Kong has resulted in the need for more land, and the generation and disposal of significant amounts of raw sewage waste. Reclamation activities have altered the harbour profile and increased sedimentation rates in many areas around Hong Kong. Transitions of seawall-type sewage outfalls to submarine-type sewage outfalls further into the channel of Victoria Harbour have also had an impact on the concentration and dispersion of sewage waste in Hong Kong waters.

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# **CHAPTER 2**

# Samples and Methodology

# 2.1 Core Samples

Four intervals of a piston core section (MBH 54/2; 0.5m-4.2m) were obtained from the Kwun Tong Typhoon Shelter in Victoria Harbour's Kowloon Bay (**Fig. 1.3**). Core MBH 54/2 (**Fig. 2.1**) was supplied from the collection of Dr. W. W. -S. Yim, from the Department of Earth Sciences, at the University of Hong Kong. MBH 54/2 was collected in 1996 by rotary boring in stainless steel casings, wax sealed, and sent to the University of Oklahoma where it was stored in a freezer at -21°C.



**Fig. 2.1.** Core MBH 54/2-Kwun Tong Typhoon Shelter, Kowloon Bay. Estimated sedimentation rates were reported by Tanner *et al.* (2000) for MVC 74.

#### 2.1.1 Sediment Core Description – MBH 54/2

The majority of sediments in core MBH 54/2 (**Fig. 2.1**) are made up of dark greenish-gray to black, soft silty clays, with the presence of shell fragments. The base of the core section is composed of more compact, lighter colored, coarse grained sediments. Sediments in the uppermost unit (0.5m to 2.4m) were dark greenish-gray to black silty clays, underlain by light brownish-gray silty clays (2.5m to 3.1m), dark gray silty clays (3.1m to 3.5m) which transitioned into lighter gray sandy clays with red streaks and coarse quartz grains (3.9m to 4.2m). The dark greenish-gray to black sediment color observed throughout much of the core section is due to the presence of sulfides (*i.e.*, pyrite), from sulfate reduction. The light gray sandy clays with red streaks have been observed in desiccated pre-Holocene sediments formed during a low sea-level stand (Yim, 1994).

#### 2.2 Overview of Experimental Method

Cores were sectioned into 5 cm intervals and stored at -21°C. Samples from various depths were freeze-dried using a Labconco Freeze Dryer 5. The following sections will provide detailed procedures used to isolate and analyze free, ester- and amide-bound lipids. Surrogate standards (cholestane and cholanic acid) were added to a small number of sediment samples prior to extraction. An external deuterated standard ( $C_{24}D_{50}$ ) was added to the lipid extracts prior to analysis by gas chromatography (GC) and gas chromatography-

mass spectrometry (GCMS) to assess the extraction efficiency. Results summarizing surrogate standard recovery were misplaced. However, the percent recovered ranged between 90% and 110%. The flowchart in **Fig. 2.2** summarizes the experimental procedures.



**Fig. 2.2.** Flowchart summarizing methodology for separating and isolating lipid groups from marine sediments.

# 2.3 Free Lipid Extraction and Fractionation

Cellulose extraction thimbles, glasswool, and boiling chips were pre-

extracted with dichloromethane:methanol (1:1 v/v, at least 6hrs). Copper

trimmings were activated in dilute hydrochloric acid (10% HCI), followed by

ultrasonication in deionized water (3x), methanol (3x), and dichloromethane (3x).

Frozen sediments were transferred to Lyph-Lock flasks (100ml) with 24/40 joints and connected to valve ports on the drying chamber of a Labconco Freeze-Dry system (duration on the freeze-dry system was typically 24hrs). Dried sediments were ground to a fine powder using a ceramic mortar and pestle, weighed and transferred to pre-extracted cellulose thimbles for extraction of free lipids.

Free lipids were extracted from freeze-dried sediments using a mixture of dichloromethane:methanol (2:1 v/v, 48hrs). Activated copper was used to remove elemental sulfur extracted with the free lipid fraction. Excess solvent was removed using a Yamato RE-51 rotary evaporator under vacuum, and transferred to pre-weighed vials where solvents were completely evaporated under a stream of nitrogen gas. Sample weights were recorded.

# 2.3.1. Separation of Total Free Lipid Extracts into Non-Saponifiable and Saponifiable Fractions

A fraction of the total free lipid extract was saponified by refluxing with 6N potassium hydroxide (15ml; 6hrs) in 10% aqueous methanol. Non-saponifiable lipids, sometimes referred to as "neutral lipids," were extracted with dichloromethane (5x30ml) in a separator funnel (vigorously shaken, 1min). The aqueous phase was acidified to pH~2 by the addition of 4N HCl to release saponifiable lipids (also referred to as "acidic lipids"). Saponifiable lipids were recovered from the aqueous phase by liquid-liquid extraction using dichloromethane. Excess solvent was removed from lipid fractions by rotary

evaporation under vacuum, transferred to pre-weighed 4ml vials, dried under a stream of nitrogen gas, and weighed.

# 2.3.2. Fractionation of Non-Saponifiable Lipids to Saturate, Aromatic, and Polar Fractions

Samples with sufficient total free lipid extracts were separated into saturate, aromatic, and polar fractions by column chromatography. Free non-saponifiable lipids were eluted through a 100ml glass column, packed with pre-conditioned alumina (14g; 80-200 mesh). Solvents of increasing polarity were used to elute saturate (n-pentane, 50ml), aromatic (n-pentane:dichloromethane 7:3 v/v, 50ml) and polar compounds (dichloromethane:methanol, 97:3 v/v, 50ml). Lipid fractions were concentrated by rotary evaporation and transferred to pre-weighed 4ml vials. Both saturate and aromatic fractions were screened by GC and GCMS.

The polar fraction was separated into three aliquots. The first aliquot was screened by GC and GCMS. The second aliquot was methylated using BF<sub>3</sub>-methanol (see section 2.5.1 for methylation procedure) and screened by GC and GCMS, followed by silylation (see section 2.5.2 for silylation procedure) of a small fraction of the methylated free non-saponifiable polar lipids. Free polar lipid compounds that were methylated and silylated were analyzed by GC and GCMS.

#### 2.4 Extraction of Ester- and Amide-Bound Lipids

Ester- and amide-bound lipids were released and extracted from residual sediments using alkaline and acid hydrolysis, respectively. The technique used to extract the bound lipids is based on a slight modification of the procedure described by Lattuati *et al.* (2002). A Soxhlet extraction step was added at the end of each procedure to recover ester- or amide-bound lipids not extracted during the filtration step.

"Residual sediment-1" was refluxed in a solution of 1N potassium hydroxide (KOH) in methanol (30ml, 2hrs), to cleave ester-linkages of lipids present in the sediments. The reaction mixtures were filtered through vacuum flasks, using pre-combusted glass fiber filters (Whatman 934-AH). Residual sediment-1 was rinsed with methanol (100ml) and the combined filtrates were transferred to a large separator funnel (1000ml). Combined filtrates in the separator funnel were acidified using 10% aqueous HCI. Residual sediment-1 was then rinsed with dichloromethane (50ml), which was transferred to the separator funnel containing the combined filtrates. The solution mixture in the separator funnel was shaken vigorously (1min), producing a monophasic solution. The solution mixture was then diluted with deionized water (50ml) and additional dichloromethane (50ml). The solution mixture was shaken vigorously (1min), resulting in a biphasic solution, where the ester-bound lipids were extracted into the dichloromethane layer. The remaining residual sediment and glass fiber filter were transferred to a pre-extracted cellulose extraction thimble, and Soxhlet extracted with dichloromethane:methanol (2:1 v/v, 24hrs) to recover

any remaining released ester-bound lipids. All released ester-bound lipids were combined, concentrated, and weighed in 4ml vials, representing the total esterbound lipid fraction.

"Residual sediment-2" (**Fig. 2.2**), was hydrolyzed by reflux in 4N HCl (40ml, 6hrs) to release amide-bound lipids. Hydrolyzed sediments were filtered through pre-combusted glass fiber filters (Whatman 934-AH) and rinsed with methanol (100ml). Combined filtrates were transferred to a large separator funnel (1000ml). The residual sediment was washed with dichloromethane (50ml), which was then transferred to the separator funnel and shaken vigorously (1min). Deionized water (50ml) and additional dichloromethane (50ml) were added to the separator funnel and shaken vigorously (1min), to recover the organic phase containing the amide-bound lipids. Any freed amide-bound lipids remaining in the sediment were recovered by Soxhlet extraction (dichloromethane:methanol; 2:1 v/v, 24hrs) and combined with the previously isolated amide-bound lipids. The total amide-bound lipid fraction was concentrated and weighed in 4ml vials.

#### 2.5 Derivatization of Functionalized Lipids

Functionalized lipids (*e.g.*, alcohols, sterols, fatty acids, and hydroxy fatty acids) typically require derivatization procedures in order to produce less polar, more volatile compounds which can be separated on gas chromatographic columns. Methylation and silylation procedures, respectively, are commonly used in the derivatization of compounds with carboxyl or hydroxyl groups. Lipids

containing both carboxyl and hydroxyl groups are typically methylated, then silylated, to produce methyl ester-trimethylsilyl (TMS) ether compounds.

#### 2.5.1 Methylation of Saponifiable, Ester- and Amide-Bound Lipids

Aliquots of saponifiable, ester- and amide-bound lipids were methylated using BF<sub>3</sub>-methanol (14% borontrifluoride, 86% methanol). In general, for 5mg of lipids, 500µl of BF<sub>3</sub>-methanol was used for methylation reactions. Lipid fractions (*i.e.* saponifiable, ester- and amide-bound lipids) were transferred and weighed in 4ml vials. BF<sub>3</sub>-Methanol was added to the vials, capped, and heated at 60°C (15min). The reaction mixture was allowed to cool, transferred to a separatory funnel (125ml), and diluted with deionized water (20ml). Lipids, as methyl esters, were recovered by liquid-liquid extraction with dichloromethane (3x40ml). Samples were concentrated using a rotary evaporator, transferred to preweighed vials, and any remaining solvent removed under a stream of nitrogen gas. Sample weights were recorded and small aliquots of samples were set aside for GC, GCMS, and gas chromatography-isotope ratio mass spectrometry (GCIRMS) analyses. All methylated samples were stored in a refrigerator.

#### 2.5.2 Silylation of Non-Saponifiable, Ester- and Amide-Bound Lipids

Non-saponifiable lipids, ester- and amide-bound lipid-methyl esters were silylated to produce trimethylsilyl ethers. Lipid samples (up to 1mg) were heated

to 50°C (30min) in N,O,-bis(trimethylsilyl)trifluoro-acetamide with 1% trimethylchlorosilane (BSTFA with 1%TMCS; 100µl) and pyridine (100µl), which was used as a catalyst for the reaction. After silylation, samples were dried under a flow of nitrogen gas, diluted in dichloromethane, and analyzed by GC and GCMS. Remaining samples were stored in a refrigerator.

#### 2.6 Gas Chromatography

Lipid fractions were initially screened on a Hewlett-Packard 5890 gas chromatograph equipped with an on-column injector and a flame ionization detector (FID) set at 310°C. Samples were chromatographed on an Agilent/J&W HP-5MS fused silica capillary column (30m x 0.25mm i.d. x 0.5µm film thickness), which has a non-polar stationary phase composed of (5%-phenyl)methylpolysiloxane. The oven temperature was programmed from 40°C to 310°C, at a rate of 4°C/min, and held isothermally at 310°C (32min). Data were acquired with a PE Nelson-900 series acquisition interface and transferred to a Windows-based computer at the end of the run. Chromatograms were processed and plotted using PE Nelson Chromatography Software.

#### 2.7 Gas Chromatography – Mass Spectrometry

Lipid fractions were analyzed by gas chromatography-mass spectrometry with a Varian 3400 gas chromatograph interfaced via transfer line to a Finnigan

MAT triple stage guadrupole mass spectrometer (TSQ-70). Gas chromatography was performed on an Agilent/J&W Scientific DB-5MS fused silica capillary column (60m x 0.32mm i.d. x 0.25µm film thickness), which utilizes a (5%phenyl)-methylpolysiloxane equivalent non-polar stationary phase. The split/splitless injector on the gas chromatograph was temperature programmed from 40°C to 310°C at a rate of 180°C/min, then held isothermal at 310°C (99min). The GC oven program was initially held at 40°C (1.5min), then heated to 310°C at a rate of 4°C/min, where it was held at 310°C (31min). The transfer line was isothermal (310°C) for the entire length of the run (100min). Analyses were completed in full-scan mode, where compounds were ionized by electron impact ionization (EI @ 70eV). The electron multiplier was set to detect and measure ions over the mass range of m/z 50 to m/z 550 each second. GCMS results were acquired, processed, and interpreted on a DEC Alpha Workstation, using the ICIS/ICL (*i.e.* "Interactive Chemical Information System"/"Interactive Control Language") data acquisition software.

#### 2.8 Gas Chromatography – Isotope Ratio Mass Spectrometry (GCIRMS)

Compound specific carbon isotopes were measured on two GCIRMS systems – a Varian 3410 GC interfaced via combustion reactor to a Finnigan MAT-252 isotope ratio mass spectrometer; and a HP6890A GC interfaced via ThermoQuest Finnigan GC Combustion III furnace to a ThermoQuest Delta<sup>plus</sup>XL isotope ratio mass spectrometer. GCIRMS allows  $\delta^{13}$ C values to be measured for

individual components in complex compound mixtures. In this project, fatty acids in the ester- and amide-bound lipid fractions were analyzed by GCIRMS as methyl esters. Samples were introduced into the GC, where the oven was temperature programmed from 40°C to 310°C at a rate of 4°C/min (total run time = 100min). The Varian 3410 was equipped with a DB-1 fused silica capillary column (60m x 0.32mm I.D. x 0.25µm film thickness), which has a non-polar stationary phase (100% dimethylpolysiloxane); and the HP6890A was equipped with a HP-5MS fused silica capillary column (30m x 0.25mm i.d. x 0.5µm film thickness), which has a non-polar stationary phase composed of (5%-phenyl)methylpolysiloxane. Components separated on the GC columns passed through a ceramic combustion reactor (980°C), and completely combusted to  $CO_2$  and  $H_2O$ . The water was removed with a water separator prior to introducing the  $CO_2$ into the isotope ratio mass spectrometer, where the relative proportions of <sup>13</sup>CO<sub>2</sub> to <sup>12</sup>CO<sub>2</sub> were determined relative to the Pee Dee Belemnite (PDB) standard. The data acquisition system converted isotopic ratios of  ${}^{13}C/{}^{12}C$  to delta ( $\delta$ ) notation, using equation 2.8.1 (Hoefs, 1997).

$$\delta^{13} C = \left(\frac{{}^{13} C / {}^{12} C_{sample}}{{}^{13} C / {}^{12} C_{standard}} - 1\right) \times 1000 \quad (eq. 2.8.1)$$

 $\delta^{13}$ C values for fatty acids were corrected for the addition of a methanol carbon, from the methylating reagent BF<sub>3</sub>-methanol. Bulk stable isotopes of a C<sub>24:0</sub> fatty acid standard and fatty acid methyl ester product formed by the methylating reagent were measured by Rick Maynard from the Organic

Geochemistry Laboratory at the University of Oklahoma. The isotopic composition of the methanol carbon  $\delta^{13}C_{MEOH}$  was determined using **equation 2.8.2**, where x is the fractional carbon contribution of fatty acid to fatty acid methyl ester (*e.g.* tetracosanoic acid would have a fractional carbon contribution where x=24/25) (Abrajano *et al.*,1994). The isotopic composition of fatty acids ( $\delta^{13}C_{FA}$ ) was calculated (**equation 2.8.2**) using the known  $\delta^{13}C_{MEOH}$  value and measured  $\delta^{13}C_{FAME}$  compositions.

$$\delta^{13}C_{FAME} = [x]\delta^{13}C_{FA} + (1-x)\delta^{13}C_{MEOH}$$
 (eq. 2.8.2)

# 2.9. Elemental Analysis, and Bulk Organic Carbon ( $\delta^{13}C_{org}$ ) and Total Nitrogen ( $\delta^{15}N$ ) Stable Isotope Measurements

Freeze-dried sediment samples were sent to the Sedimentary Coastal and Oceanic Organic Biogeochemistry Laboratory at Texas A & M University for elemental analysis, and bulk stable isotope measurements of carbon and nitrogen. A Costech EA (Model # ECS4010), interfaced to a Finnigan MAT-252 dual inlet isotope ratio mass spectrometer via a Thermo-Finnigan Conflo III interface, was used to measure total organic carbon ( $C_{org}$ ), total nitrogen (N),  $\delta^{13}C_{org}$  and  $\delta^{15}N$  of decarbonated sediment samples. The following temperature and flow parameters were used in the analyses: combustion furnace = 1020°C; reduction furnace = 650°C; column temperature = 40°C; and flow-rate = 98ml/min. <sup>13</sup>C/<sup>12</sup>C and <sup>15</sup>N/<sup>14</sup>N isotopic ratios were compared to standards (PDB and

atmospheric nitrogen, respectively), and converted to delta ( $\delta$ ) notation.  $\delta^{13}$ C was calculated using **equation 2.8.1**, and  $\delta^{15}$ N was calculated using **equation 2.9.1**.

$$\delta^{15} N = \left(\frac{{}^{15} N / {}^{14} N_{\text{sample}}}{{}^{15} N / {}^{14} N_{\text{standard}}} - 1\right) \times 1000 \quad \text{(eq. 2.9.1)}$$

Samples were weighed in silver boats and decarbonated in glass desiccators, with a small amount of 12N HCl at the base of the desiccator. After 72hrs, acid vapors were removed by placing samples in a vacuum oven (<30°C, 24hrs). Samples free of acid vapors were then placed into tin boats, crushed, closed and ready for analysis. Since nitrogen is typically present in low amounts relative to carbon, it is the limiting component. Thus, sample sizes were adjusted to obtain adequate measurements for  $\delta^{15}$ N (*i.e.*, about 100µg to 200µg N is needed for an adequate signal on the isotope ratio mass spectrometer). Carbon, however, could be diluted with helium gas to reduce the signal (*i.e.*, to about 75µg to 190µg C) for a reliable  $\delta^{13}C_{org}$  measurement.

## 2.10. References

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#### **CHAPTER 3**

# Elemental and Stable Isotope Analyses of Organic Carbon and Total Nitrogen in Sediments from Kowloon Bay, Hong Kong

#### 3.1 Introduction

Bulk properties of organic matter in marine sediments provide insights into the history and changes in environmental conditions in the sedimentary record. The piston core investigated in this study (*i.e.*, core MBH 54/2) provides a record of organic matter deposited during the late Quaternary, revealing changes in organic source material, periods affected by anthropogenic activities, and natural changes to this region. The bulk properties utilized include elemental and stable isotope compositions of organic carbon and total nitrogen in sedimentary organic matter. Sedimentary organic matter is comprised of complex mixtures which can include lipids, proteins, cellulose, lignin, and/or other components originating from various organisms or anthropogenic wastes. The majority of organic matter introduced into the marine environment undergoes remineralization (*i.e.*, organic matter is oxidized, resulting in the production of  $CO_2$ ,  $H_2O_1$ , and nutrients) in the water column, where less than 10% of the original organic matter is incorporated and preserved in the sediments (Meyers and Lallier-Vergès, 1999; Meyers, 2003). While remineralization occurs during sedimentation, bulk properties of deposited sedimentary organic matter still retain important information for delineating the sources of organic matter and possible processes that may have occurred (e.g.,

denitrification or enhanced productivity; Teranes and Bernasconi, 2000; Bratton *et al.*, 2003; Meyers, 2003).

#### 3.2 Review of Literature

#### 3.2.1 Organic Carbon and Total Nitrogen in Marine Sediments

The total organic carbon (TOC) content in marine sediments measures the amount of organic matter that survived remineralization processes in the water column and was preserved in the sediments (Meyers and Lallier-Vergès, 1999). The amount of organic matter in sediments has been estimated to be twice the amount of measured TOC (*i.e.*, 50% of sedimentary organic matter is composed of carbon; Meyers, 2003). Changes in TOC content reflect periods of higher or lower influx of organic matter, which includes the total mixture of terrigenous plant material, algae, bacterial biomass, sewage effluents, and/or other sources of organic carbon.

Nitrogen is an important nutrient and plays an important role in productivity in the marine environment (Minagawa and Wada, 1986; Teranes and Bernasconi, 2000; Talbot, 2001). If nitrogen availability is too low, then primary production is limited; if nitrogen is abundant this can result in intense algal blooms and eutrophication of the marine system (Sigleo and Macko, 2002). Nitrogen is typically characterized in sediments as total nitrogen, or as the ratio of organic carbon-to-nitrogen (C/N), and provides important information to aid in delineating sources and past variations of nitrogen in the environment (Meyers, 1994; Talbot,

2001). Atmospheric nitrogen is plentiful, comprising about 78% of the earth's atmosphere, and is the primary source of nitrogen for terrigenous plants (Peters *et al.*, 1978; Sweeney *et al.*, 1978; Hoefs, 1997; Meyers and Lallier-Vergès, 1999). Terrigenous plants are able to utilize atmospheric nitrogen after bacteria, located around the plant roots, convert the nitrogen to ammonia via nitrogen fixation (**eq. 3.1**; Peters *et al.*, 1978; Sweeney *et al.*, 1978; Sweeney *et al.*, 1978; Whelan and Farrington, 1992; Bickert, 2000). Terrigenous plants can also uptake nitrogen, in the form of nitrates, around the plant roots (Moore, 2004).

Nitrogen Fixation:  $N_{2-atmos} + 3H_2O \rightarrow 2NH_3 + \frac{3}{2}O_2$  (eq. 3.1)

Marine organisms assimilate dissolved inorganic nitrogen in the form of nitrate  $(NO_3^-)$ , ammonium  $(NH_4^+)$ , or nitrite  $(NO_2^-)$ , although nitrate is the most common form of assimilated nitrogen (Peters *et al.*, 1978; Fogel and Cifuentes, 1993; Bickert, 2000; Talbot, 2001). Nitrate is reduced to nitrogen gas  $(N_2)$  by anaerobic bacteria via denitrification processes (**eq. 3.2**), returning nitrogen to the atmosphere (Hoefs, 1997; Bickert, 2000; Teranes and Bernasconi, 2000; Talbot, 2001). Denitrification is an important mechanism for balancing the natural processes of nitrogen fixation (Sweeney *et al.*, 1978). Ammonia (NH<sub>3</sub>) and ammonium  $(NH_4^+)$  in aquatic environments are produced by bacterial decomposition of organic matter under anaerobic conditions (Teranes and Bernasconi, 2000; Talbot, 2001). The ammonia formed from the mineralization of organic nitrogen is an important component utilized by aerobic bacteria during

nitrification processes (Sweeney *et al.*, 1978). Conversion of ammonia to nitrate can be expressed by **equation 3.3** (Bickert, 2000).

Denitrification: 
$$5CH_2O + 4NO_3^- + 4H^+ \rightarrow 5CO_2 + 7H_2O + 2N_2$$
 (eq. 3.2)

Nitrification: 
$$\frac{NH_3 + \frac{3}{2}O_2 \rightarrow HNO_2 + H_2O}{HNO_2 + \frac{1}{2}O_2 \rightarrow HNO_3}$$
 (eq. 3.3)

# 3.2.2 Carbon-to-Nitrogen Ratio (C/N)

TOC and total nitrogen are commonly utilized together, where the ratio of organic carbon-to-total nitrogen functions as a tool for distinguishing the source of organic matter. The primary nitrogen components in marine organisms (*e.g.*, phytoplankton and zooplankton) are proteins. Vascular terrigenous plants, however, have low protein content and are enriched in cellulose and lignin. The low cellulose, high protein composition of phytoplankton results in C/N ratios between 4 and 10, whereas bacterioplankton have C/N ratios ranging between 2.6 and 4.3. High cellulose and lignin composition of vascular terrigenous plants, along with low protein content, results in C/N ratios greater than 15 (Sampei and Matsumoto, 2001; Meyers, 2003; Wu *et al.*, 2003). The downcore profile of the C/N ratio in a sediment core section can illustrate shifts through time as the principal contributor of organic matter changes between terrigenous and algal material. Preferential loss of nitrogen relative to carbon can occur as a result of the decomposition of algal biomass settling through the water column.

Consequently, organic matter deposited in the sediments may appear to have a higher C/N ratio (Sampei and Matsumoto, 2001).

# 3.2.3 Bulk Stable Isotope Analyses

Bulk stable isotope composition of organic carbon and total nitrogen complement the C/N ratio. Carbon and nitrogen each have two stable isotopes,  $^{12}$ C (98.89%) and  $^{13}$ C (1.11%), and  $^{14}$ N (99.64%) and  $^{15}$ N (0.36%) (Hoefs, 1997). Stable isotopes of carbon and nitrogen are expressed using the delta ( $\delta$ ) value, where the ratio of heavy-to-light isotope values (*i.e.*,  $^{13}$ C/ $^{12}$ C and  $^{15}$ N/ $^{14}$ N) are calibrated relative to international standards (**eq. 2.8.1** and **eq. 2.9.1**; Hoefs, 1997).

 $\delta^{13}C_{org}$  values have been used to determine the sources of sedimentary organic matter (*i.e.*, marine or terrigenous) and to identify photosynthetic pathways utilized by terrigenous plants (*i.e.*, C3 plants or C4 plants). C3 plants incorporate carbon into organic matter using the Calvin pathway, where a molecule of CO<sub>2</sub> reacts with the enzyme ribulose 1,5-bis-phosphate carboxylase to produce two molecules of 3-phosphoglycerate (**Fig. 3.1**; Fogel and Cifuentes, 1993; Hoefs, 1997).



**Fig. 3.1.** Incorporation of carbon into organic matter utilizing the C3 pathway (from Fogel and Cifuentes, 1993).

C4 Plants follow the Hatch-Slack pathway, where the incorporation of carbon into organic matter occurs when CO<sub>2</sub> reacts with phosphoenolpyruvate carboxylase to form oxaloacetate (**Fig. 3.2**; Fogel and Cifuentes, 1993; Hoefs, 1997).  $\delta^{13}$ C values of C3 terrigenous plants range between -33°/<sub>oo</sub> and -22°/<sub>oo</sub>; whereas C4 terrigenous plants are more <sup>13</sup>C-enriched and range between -22°/<sub>oo</sub> and -8°/<sub>oo</sub> (Meyers, 1994; Hoefs, 1997; Huang *et al.*, 1999; Meyers and Teranes, 2001). Marine organic matter (*e.g.*, marine algae) is isotopically heavier than C3 terrigenous plants, and has  $\delta^{13}$ C values ranging between -25°/<sub>oo</sub> and -20°/<sub>oo</sub> (Meyers, 1994).



**Fig. 3.2.** Photosynthetic pathway of C4 plants, utilizing the enzyme phosphoenolpyruvate carboxylase (from Fogel and Cifuentes, 1993).

Nitrogen isotopes often supplement carbon isotope measurements for delineating source information, where the  $\delta^{15}N$  value varies as a result of the form of inorganic nitrogen assimilated. The two primary forms of nitrogen utilized are nitrate (*i.e.*, for marine organisms;  $\delta^{15}N_{\text{nitrate}} = 7^{\circ}/_{\text{oo}}$  to  $10^{\circ}/_{\text{oo}}$ ) and atmospheric nitrogen (*i.e.*, for terrigenous plants;  $\delta^{15}N_{atmos} = \sim 0^{\circ}/_{oo}$ ) (Peters *et al.*, 1978; Meyers and Lallier-Vergès, 1999; Meyers and Teranes, 2001; Owen and Lee, 2004). Nitrates derived from anthropogenic activities (*e.g.*, human and/or animal waste products) are enriched in <sup>15</sup>N where  $\delta^{15}$ N ranges from 10°/<sub>oo</sub> to 25°/<sub>oo</sub> (Teranes and Bernasconi, 2000; Meyers, 2003). These isotopic signatures can be traced in the environment where marine plankton or algae utilize NO<sub>3</sub> and terrigenous plants utilize atmospheric nitrogen as their nitrogen source (Muzuka et al., 1991; Meyers and Lallier-Vergès, 1999; Meyers and Teranes, 2001). Several factors, however, can affect the isotopic composition of nitrogen in organic matter, providing insights into nutrient cycling and processes such as nitrogen fixation or denitrification (Fogel and Cifuentes, 1993; Bickert, 2000).

# 3.2.4 Sewage Derived Carbon and Nitrogen in Marine Sediments

Nitrogen is an essential substrate needed for primary producer growth. The isotopic composition of the primary producers will be dependent on the isotopic composition of dissolved inorganic nitrogen components such as  $NO_3^-$ ,  $NH_4^+$ , or  $NO_2^-$  (Montoya, 1994). Potential sources of dissolved inorganic nitrogen in the aquatic environment include sewage and wastewater effluents. These sources of nitrogen can serve as nutrients for marine organisms. When nutrient overloading occurs (*e.g.*, via excessive sewage waste disposal), algal blooms can be stimulated leading to eutrophication. Eutrophication results in increasing anoxic conditions whereby excessive denitrification processes can occur, enriching the <sup>15</sup>N in residual dissolved inorganic nitrogen (Muzuka et al, 1991; Montoya, 1994; Teranes and Bernasconi, 2000; Bratton et al., 2003). Conversely, as the phytoplankton assimilate and metabolize the dissolved inorganic nitrogen, the phytoplankton biomass becomes depleted in <sup>15</sup>N relative to that of the growth substrate (Montoya, 1994). In the early stages of algal blooms, phytoplankton biomass is isotopically lighter than the dissolved inorganic nitrogen being used as the substrate. As <sup>14</sup>N is selectively removed, the residual dissolved inorganic nitrogen becomes isotopically heavier. Progressive assimilation of the isotopically heavier nitrogen will be reflected in the phytoplankton biomass (Montoya, 1994). Bacterial remineralization of settling particulate organic matter in the water column can also lead to progressive enrichment of <sup>15</sup>N in the residual organic matter (Altabet and McCarthy, 1985; Bickert, 2000).

Offshore disposal of sewage waste has been common practice since the early occurrence of population growth in near-coast environments with the presumption that minimal sewage particulates would reach or accumulate on the seafloor due to dispersion and dilution effects in surface waters. Various research groups have utilized elemental and bulk stable isotope compositions of carbon and nitrogen in sewage effluents, particulate organic matter, marine sediments, and/or phytoplankton-derived organic matter to study the effects of

sewage-derived organic matter on marine sediments. **Table 3.1** summarizes the results of studies from various localities around the world. Sewage effluents, sewage sludges, and sewage-derived organic matter sampled around coastal regions have been reported with the following bulk properties:  $\delta^{13}C_{org}$ =-28.5°/<sub>oo</sub> to -23.0°/<sub>oo</sub>;  $\delta^{15}N$ =1.8°/<sub>oo</sub> to 3.2°/<sub>oo</sub>; and C/N ratios=11.0 to 13.4 (Burnett and Schaeffer, 1980; Sweeney *et al.*, 1980; Gearing *et al.*, 1991; van Dover *et al.*, 1992; Hunt *et al.*, 1992; Thornton and McManus, 1994; Rogers, 2003).

Organic matter derived from sewage effluents have been differentiated from terrigenous derived organic matter (Rogers, 2003) and organic matter derived from marine sediments unaffected by sewage (Burnett and Schaeffer, 1980; Sweeney *et al.*, 1980) using bulk stable isotope values of carbon and nitrogen. The proportion of sewage contributions can be estimated using isotopic measurements of sewage contaminated sites and pristine sites (Rogers, 2003). Burnett and Schaeffer (1980), for example, used **equation 3.5** to estimate the relative amount of sewage affected sediments at their study sites, where:  $F_s=%$ sewage sludge;  $\delta^{13}C=\delta^{13}C$  value of  $C_{org}$  in the measured samples;  $\delta^{13}C_m=\delta^{13}C$ value of  $C_{org}$  in uncontaminated marine shelf sediments "normal" to the area;  $\delta^{13}C_s=\delta^{13}C$  value of the sewage sludge.

$$F_{s} = \frac{\delta^{13}C - \delta^{13}C_{m}}{\delta^{13}C_{s} - \delta^{13}C_{m}} \times 100$$
 (eq. 3.5)

Location	Sample	%C <sub>org</sub>	%N	C/N Ratio	δ <sup>13</sup> C (°/ <sub>00</sub> )	δ <sup>15</sup> N (°/ <sub>00</sub> )	Reference
Western Hong Kong (estuarine-derived)	Sediment	0.4 to 1.2	-	6.0 to 18.5	-25.3 to -21.3	5.0 to 12.5	Owen, 2005
Central Hong Kong	Sediment	0.5 to 1.1	-	6.0 to 20.0	-	-	Owen, 2005
Eastern Hong Kong (marine-derived)	Sediment	0.6 to 3.0	-	6.0 to 25.5	-26.7 to -19.0	10.0 to 18.5	Owen, 2005
Lake Biwa	Sediment	-	-	-	-24.7 to -21.5	5.7 to 7.8	Mishima et al., 1999
Yodo River	Sediment	-	-	-	-26.2 to -24.7	4.3 to 6.7	Mishima et al., 1999
Ane River	Sediment	-	-	-	-28.3 to -26.6	-0.9 to 2.6	Mishima et al., 1999
Tokyo Bay	POM	-	-	-	-15.0	-	Mishima et al., 1999
Otuchi Bay	ТОМ	-	-	-	-26.5	-	Mishima et al., 1999
S. California Bight	Marine PM	-	-	-	-21.0 to -19.0	8.0 to 12.0	Spies et al., 1989
S. California Bight (Whites Point, CA)	Sewage PM	-	-	-	-16.5	1.8	Spies et al., 1989
S. Calif. Coastal Area-1	Sewage Effluent	30.8	2.33	13.2*	-	3.0	Sweeney et al., 1980
S. Calif. Coastal Area-2	Sewage Effluent	31.7	2.38	13.3*	-	2.0	Sweeney et al., 1980
S. Calif. Coastal Area-3	Sewage Effluent	31.8	2.37	13.4*	-	2.4	Sweeney et al., 1980
S. Calif. Coastal Area-4 (Whites Point, CA)	Sewage Effluent	31.7	2.36	13.4*	-	2.5	Sweeney et al., 1980
Edinburgh (marine embayment)	Sewage Effluent	-	-	-	-25.2±0.9	10.7±0.7	Waldron <i>et al.</i> , 2001
Edinburgh (marine embayment)	Sediment	-	-	-	-22.9±0.2	6.1 to 6.7	Waldron et al., 2001
Cranston, Ri	Sewage Sludge	-	-	-	-23.5±0.4	-	Gearing et al., 1991
Los Angeles	Sewage Sludge	-	-	-	-23.5±0.5	-	Gearing et al., 1991
New York	Sewage Sludge	-	-	-	-26.0 & -25.7	-	Gearing et al., 1991
Moa Point, New Zealand	Sewage Effluent	-	-	-	-23.5	1.8 to 2.5	Rogers, 2003
Middlesex, NJ	Sewage derived OM	-	-	-	-24.7	-1.1	van Dover et al., 1992
Mergen, NJ	Sewage derived OM	-	-	-	-23.2	6.1	van Dover et al., 1992
Yonkers, NY	Sewage derived OM	-	-	-	-21.4	7.2	van Dover et al., 1992
Providence, RI	Sewage derived OM	-	-	-	-23.7	-	van Dover et al., 1992
Hunts Bay, Kingston Harbour, Jamaica	Sewage	-	-	11.0 to 13.0	-28.5 to -23.0	-	Andrews et al., 1998
New York Bight-Newton Creek Treatment Plant	Sewage Sludge	-	-	-	-25.7	-	Burnett & Schaeffer, 1980
New York Bight- Ward Island Treatment Plant	Sewage Sludge	-	-	-	-26.0	-	Burnett & Schaeffer, 1980
Mangrove Creek, Hong Kong	Seston	-	-	-	-27.16±0.44	10.48±0.21	Lee, 2000
Pearl River Estuary	Seston (phytoplankton dominated)	-	-	-	-25.22±0.48	-1.06±0.98	Lee, 2000
Shan Pui River, Hong Kong	POM (primarily anthropogenic)	-	-	-	-24.13	5.23	Lee, 2000
New York/New Jersey	Sewage-derived OM	-	-	-	-23.0	3.2	Hunt <i>et al.</i> , 1992
North Atlantic	Phytoplankton- derived POM	-	-	-	-21.7	6.1	van Dover et al., 1992
Western North Atlantic	Sed. POC (phytoplankton- derived)?	-	-	-	-21.6	-	Sayles & Curry, 1988; van Dover <i>et al.</i> , 1992
Pacific-Deepwater	PON	-	-	-	-	5.0 to 15.0	Saino & Hattori, 1987; van Dover <i>et al.</i> , 1992
Sargasso Sea	PON	-	-	-	-	5.0 to 7.0	Altabet, 1988; van Dover <i>et al.</i> , 1992
Invergowrie Bay, Tay Estuary, Scotland	Sewage Effluent	-	-	12.57	-26.7	2.3	Thorton & McManus, 1994

**Table 3.1.** Summary of elemental and bulk stable isotope measurements of carbon and nitrogen in marine sediments, sewage sludge/effluents, particulate organic matter, and plankton-derived particulate organic carbon.

#### 3.3 Results and Discussion

Bulk properties of sedimentary organic matter (*i.e.*, elemental and stable isotope compositions of carbon and nitrogen in sediments) have been measured in core MBH 54/2 to identify sources of organic matter, speculate on processes that affect carbon and nitrogen preservation and isotopic composition (*e.g.*, signatures resulting from microbial activity), and to infer processes that occurred as a result of raw sewage disposal in coastal environments. While the majority of organic matter (>90%) is remineralized during sedimentation, the C/N ratios and  $\delta^{13}C_{org}$  values of the initial organic matter that survives the water column is preserved (Meyers and Ishiwatari, 1993; Meyers, 2003; Sifeddine *et al.*, 2004). Results of elemental and bulk stable isotope measurements are summarized in **Table 3.2**.

#### 3.3.1 Elemental Analyses

Elemental analysis of organic carbon ( $\[mathcar{C}_{org}\]$ ) and total nitrogen ( $\[mathcar{M}\]$ N) content have been utilized to differentiate between terrigenous and aquatic sourced organic matter. The distribution of  $\[mathcar{C}_{org}\]$  has been used in various studies to study changes in organic matter input and preservation in sediment core sections. One possible application has been to use organic carbon distributions to reflect periods of enhanced anthropogenic waste input (*e.g.,* sewage wastes, oil spills, and/or industrial waste effluents). Bulk measurements

Sed. Rate (cm/y)*	Approx. Calendar Year	Depth (m)	%C <sub>org</sub>	%N	C/N (Weight %)	C/N (Atomic Mass)	δ <sup>13</sup> C <sub>org</sub> ( <sup>o</sup> / <sub>oo</sub> )	δ <sup>15</sup> N (°/ <sub>00</sub> )
3.5cm/y	1981	0.5	0.75	0.07	10.71	12.50	-22.60	
	1977	0.7	1.28	0.09	14.22	16.60	-23.86	6.19
4.4cm/y	1971	0.9	1.68	0.12	14.00	16.34	-28.59	2.57
	1965	1.2	2.74	0.20	13.70	15.99	-26.62	3.44
	1961	1.4	2.83	0.15	18.87	22.02	-26.30	2.14
	1954	1.6	1.62	0.11	14.73	17.19	-27.49	3.13
1.9cm/y	1943	1.8	0.88	0.09	9.78	11.41	-27.74	4.49
	1930	2.0	0.83	0.09	9.22	10.76	-27.99	3.05
	1428	2.2	0.59	0.07	8.43	9.84	-27.12	4.30
	928	2.3	0.62	0.07	8.86	10.34	-26.92	4.27
	604BC	2.6	0.94	0.06	15.67	18.28	-26.01	4.43
0.22mm/y	1604BC	2.8	0.73	0.06	12.17	14.20	-27.25	3.61
	3104BC	3.1	0.63	0.06	10.50	12.25	-27.23	4.60
	4104BC	3.3	0.88	0.06	14.67	17.12	-24.74	4.57
	5104BC	3.5	0.83	0.07	11.86	13.84	-27.50	4.33
M1	6104BC	3.7	0.86	0.05	17.20	20.07	-21.87	4.51
M2		3.9	0.42	0.04	10.50	12.25	-26.27	4.26
		4.0	0.22	0.03	7.33	8.56	-30.18	2.53
		4.1	0.39	0.03	13.00	15.17	-30.85	
		4.2	0.23	0.04	5.75	6.71	-33.17	

**Table 3.2.** Summary of elemental analyses and bulk stable isotope

 measurements of carbon and nitrogen in sediments from core MBH 54/2.

\* Sedimentation rates from 0.5m to 2.1m are based on <sup>210</sup>Pb measurements reported by Tanner *et al.* (2000). At depths below 2.1m, the sedimentation rate is based on the maximum Holocene age of 8100y BP, which is marked by the dessicated crust which represents the boundary between the M1 and M2 layers (Yim, 1994); -- Indicates that measurements were below the detection limit; values in "**bold**" indicate an average of replicate runs. C/N (atomic mass ratio) was calculated by multiplying the C/N weight ratio by 1.167 (Meyers and Teranes, 2001). See **Appendix I** for a complete list of measurements and standard deviations.



**Fig. 3.3.** Downcore profiles of  $%C_{org}$  and %N in sediment from core MBH 54/2, from Kowloon Bay, Victoria Harbour, Hong Kong SAR.

for several sediment samples were run in replicate and are summarized in **Appendix I**. The average  $%C_{org}$  had standard deviations ranging up to ±0.2; average %N up to ±0.01; average  $\delta^{13}C_{org}$  compositions as high as ±0.54; and average  $\delta^{15}N$  values within ±0.3. The downcore profile of organic carbon deposited in Kowloon Bay during the Holocene and possible late Pleistocene is shown in **Fig. 3.3a**. Three primary intervals can be observed, where the upper unit (0.7m to 1.6m) consists of the highest organic carbon content ranging between 1.28% and 2.83%. The maximum organic carbon content occurs between 1.2m and 1.4m, where  $C_{org}$  is 2.74% and 2.83%, respectively. The second unit (1.8m to 3.7m) is comprised of more intermediary organic carbon compositions ranging between 0.59% and 0.94%. Within this unit (1.8m to 3.7m) there appear to be five subgroups marked by slight changes to the organic carbon content. The first subgroup occurs between 1.8m and 2.0m, where  $C_{org}$  is 0.83% and 0.88%, respectively. The organic carbon content decreases to between 0.59% and 0.73% at depths of 2.2m to 2.3m, and 2.8m and 3.1m; a sharp spike occurs at 2.6m where  $C_{org}$  is 0.94%. A slight rise in organic carbon content to between 0.83% and 0.88% occurs between 3.3m and 3.7m. The third unit is at the base of the core (3.9m to 4.1m) where  $C_{org}$  ranges between 0.22% and 0.42%.

The total nitrogen content reported in this study is positively correlated with organic carbon (**Fig. 3.4**), where the best fit line gives a correlation coefficient (R<sup>2</sup>) of 0.8845. By extrapolating the organic carbon content to zero, the amount of inorganic nitrogen can be estimated (which was found to be ~0.02%). Thus, the total nitrogen content in sediments from MBH 54/2 can be assumed to be representative of organic nitrogen (Hedges *et al.*, 1986; Talbot and Johannessen, 1992; Andrews *et al.*, 1998; Talbot, 2001; Owen and Lee, 2004). The overall N content (**Fig. 3.3b**) is relatively low and ranges between 0.03% and 0.20%. In the same way as the organic carbon content, the nitrogen content can be divided into three major intervals. The upper unit (0.7m to 2.0m) has the highest N content ranging between 0.09% and 0.20% (maximum N occurs at 1.2m). The second unit extends from 2.2m to 3.7m and has a N composition ranging between 0.05% and 0.07%. The lowest N content occurs at

the base of the core (3.9m to 4.1m) with N values ranging between 0.03% and 0.04%.



**Fig. 3.4**. Graph of  $%C_{org}$  versus %N in sediments from core MBH 54/2, Kowloon Bay, Victoria Harbour. The trend line shows a positive correlation between organic carbon and total nitrogen.

# 3.3.2 Ratio of Organic Carbon to Total Nitrogen

The weight % ratio of organic carbon to total nitrogen (C/N) is highly variable in the sediment core (**Fig. 3.5**). Fluctuations in the downcore profile documents shifts in the proportion of terrigenous, algal, and/or anthropogenically derived organic matter deposited during the Holocene and late Pleistocene. Early diagenesis may result in alterations in the C/N ratio. For example, microbial remineralization in the water column can cause the C/N value to decrease, or

increased productivity with limited nitrogen availability can result in higher C/N values (Meyers, 1994). Despite these changes in the water column, Meyers and Ishiwatari (1993) and Meyers (1994) have demonstrated that the C/N ratios and  $\delta^{13}C_{org}$  signatures of organic matter do not undergo further diagenetic changes after burial, and that the overall source information is retained.



**Fig 3.5.** Downcore profile of the C/N (wt. % ratio) in core MBH 54/2, from Kowloon Bay, Victoria Harbour, Hong Kong SAR. "O" indicates possible flux of terrigenous-derived organic matter.
Four possible intervals are observed in **Fig. 3.5**. The first interval occurs between 0.7m and 1.6m with C/N ratios ranging between 13.7 and 14.7, except at 1.4m where the C/N ratio is 18.9. The relatively high C/N ratio between 0.7m and 1.6m occurs within a period known to have experienced a high influx of sewage waste. Higher disposal rates of sewage waste and nutrients lead to excessive algal blooms and this in turn could result in parts of Victoria Harbour becoming eutrophic. Phytoplankton and zooplankton typically have C/N ratios ranging between 4 and 10 (Meyers and Ishiwatari, 1993); however, the high C/N values observed in this interval indicates that other factors have affected this ratio. While the microbial denitrification of organic matter can result in the preferential loss of nitrogen, and increase the C/N values in sedimentary organic matter (Sarazin *et al.*, 1992; Sampei and Matsumoto, 2001; Owen and Lee, 2004), the high C/N values are likely due to contributions from sewage effluents.

The second interval occurs between 1.8m and 2.3m where the C/N ratio ranges between 8.4 and 9.8. The C/N ratio within this interval suggests that during this period Kowloon Bay received greater contributions of algal-derived organic matter compared to terrigenous-derived organic matter. These values are within the range for C/N ratios that have been used to implicate organic matter derived from phytoplankton, bacteria, and other single-celled organisms (Sigleo and Macko, 2002).

Intermediate C/N values are observed between 2.8m and 3.9m with values generally ranging between 10.5 and 12.2. Sedimentary organic matter in this region may reflect a mixture of both terrigenous- and algal-derived organic

matter. Spikes in the C/N ratio are observed at 2.6m (C/N=15.7), 3.3m (C/N=14.7), and 3.7m (C/N=17.2). The high C/N ratios at depths of 1.4m, 2.6m, 3.3m, and 3.7m (Fig. 3.5) resulted from higher organic carbon contents and were proposed to have been caused by the transport of terrigenous plant material by heavy monsoons or typhoons. At 1.4m, for example, a large spike in the C/N ratio (~18.9) was observed. Based on sedimentation rate data, sediments at this interval were deposited around 1961. Typhoons Mary (1960), Wanda (1962), Ruby (1964), and Dot (1964) passed through Hong Kong with Typhoon Wanda being the strongest typhoon to occur during this period (Yim, 1993; Huang and Yim, 1997). These events could have carried a greater abundance of terrigenous plant material into Kowloon Bay resulting in the spike in the C/N ratio observed at 1.4m. Owen (2005) reported spikes in the C/N ratio in two sediment core samples, corresponding to an event that occurred around 1910, from two locations in Tolo Harbour in northeastern Hong Kong. Heavy monsoons or typhoons were thought to have been responsible for the spike in the C/N ratio in his study. In core MBH 54/2, however, no spike in C/N ratio was observed in sediments deposited around depths corresponding to 1910 (*i.e.*, between 2.0m and 2.1m).

The fourth interval occurs at the base of the core, between 4.0m and 4.1m, where the C/N ratio is 7.3 and 5.8, respectively. A spike in the C/N ratio (13.0) is observed at 4.1m along with an elevated organic carbon content (0.39%). This may be indicative of an event where additional terrigenous organic matter was transported into Kowloon Bay.

# 3.3.3 Bulk Stable Isotope Composition of Organic Carbon and Total Nitrogen

Sewage affected marine sediments can be discriminated from unaffected sediments using  $\delta^{15}$ N signatures. Teranes and Bernasconi (2000) have reported isotopic values of sewage-derived nitrates to range between  $10.00^{\circ}/_{\circ\circ}$  and  $25.00^{\circ}/_{\circ\circ}$ . Bulk isotopic values for total nitrogen ranging between  $1.8^{\circ}/_{\circ\circ}$  and  $3.2^{\circ}/_{\circ\circ}$  have been reported in the literature for sewage effluent, sewage sludge, and sewage-derived organic matter (see **Table 3.1**). Bulk isotope values for organic carbon in these same samples were found to range between  $-28.5^{\circ}/_{\circ\circ}$  and  $-23.0^{\circ}/_{\circ\circ}$ .

Isotopically heavy organic carbon and nitrogen occur in sediments from the uppermost interval of core MBH 54/2 (0.5m to 0.7m), where  $\delta^{13}C_{org}$  ranges between -23.86°/<sub>oo</sub> to -22.60°/<sub>oo</sub>, and  $\delta^{15}N$  is 6.19°/<sub>oo</sub> (**Fig. 3.6**). Higher influxes of raw sewage and nutrients can result in intense algal blooms, which ultimately would lead to eutrophication in Kowloon Bay. Under conditions of eutrophication, severe denitrification processes can occur, leading to the isotopic enrichment of <sup>15</sup>N (Teranes and Bernasconi, 2000; Bratton *et al.*, 2003). The increased productivity can also result in organic matter becoming enriched in <sup>13</sup>C. As phytoplankton preferentially consume <sup>12</sup>C from dissolved inorganic carbon, the residual inorganic carbon will become isotopically heavier. Utilization of <sup>13</sup>C enriched inorganic carbon by phytoplankton will result in the production of isotopically heavier organic matter (Meyers, 2003).





The interval between 0.7m and 1.6m represents a period most affected by raw sewage waste. The estimated calendar years for this interval fall between 1954 and 1977, a period of rapid population growth and increased dumping of raw sewage (seawall-type sewage outfall). Bulk properties (*i.e.*,  $\delta^{13}C_{org}$ ,  $\delta^{15}N$ , and C/N ratio) of sediments in this interval are consistent with values reported in the literature for sewage contaminated sites, where sediments from MBH 54/2 have  $\delta^{13}C_{org}$  compositions between -28.59°/<sub>oo</sub> and -26.30°/<sub>oo</sub>;  $\delta^{15}N$  values range between 2.14°/<sub>oo</sub> and 3.44°/<sub>oo</sub>; and C/N ratio range between 13.7 and 14.7, except at 1.4m where the C/N ratio is 18.9.

The C/N ratio between 1.8m and 2.3m ranges between 8.4 and 9.8, suggesting that the sediments received greater contributions from algal-derived organic matter rather than terrigenous-derived organic matter during this time period. Bulk carbon isotope compositions, however, indicate values typical for C3 plants (-27.99°/<sub>00</sub> to -26.92°/<sub>00</sub>). The carbon isotopic values in this interval were expected to be heavier, with values more characteristic of aquatic-derived organic matter (*i.e.*, around -25°/<sub>00</sub> to -20°/<sub>00</sub>) since the C/N ratios indicated greater contributions from algal-derived organic matter.  $\delta^{15}$ N values ranged between 4.27°/<sub>00</sub> and 4.49°/<sub>00</sub>, except at 2.0m where  $\delta^{15}$ N was 3.05°/<sub>00</sub>. The relatively light  $\delta^{13}C_{org}$  and  $\delta^{15}$ N values may reflect the occurrence of excess bacterial biomass bound to the sediment (Bratton *et al.*, 2003).

The C/N ratio shifts between 10.5 and 15.7 at depths between 2.5m and 3.6m. Changes in the organic carbon content appear to be responsible for the variations observed in the C/N ratio. The nitrogen content is consistently around

0.06% between 2.5m and 3.6m. Spikes in C/N ratios are observed at 2.6m and 3.3m possibly reflecting a higher influx of terrigenous plant material transported into Kowloon Bay by strong storms.  $\delta^{13}C_{org}$  compositions at these intervals are -26.01°/<sub>oo</sub> and -24.73°/<sub>oo</sub>, respectively, which are normal values for C3 terrigenous plants. At other measured intervals within this unit  $\delta^{13}C_{org}$  values ranged between -27.50°/<sub>oo</sub> and -27.23°/<sub>oo</sub>.  $\delta^{15}N$  values were relatively consistent between 4.33°/<sub>oo</sub> and 4.60°/<sub>oo</sub>, except at 2.8m, where  $\delta^{15}N$  was 3.61°/<sub>oo</sub>. This may suggest that, for the most part, there was not much variability in organic matter source input within this interval.

At 3.7m, there is a large spike in the C/N ratio (*i.e.*, C/N=17.2) and the bulk carbon isotope composition is enriched in <sup>13</sup>C (-21.87°/<sub>oo</sub>). While a  $\delta^{13}C_{org}$  value of -21.87°/<sub>oo</sub> falls within the range typical for marine-derived organic matter, the sediments in this part of the core appears to be part of the desiccated crust observed throughout most of Hong Kong Harbour. The desiccated crust represents a period when marine sediments were subaerially exposed during a low sea-level stand and mark the boundary between Holocene marine sediments (*i.e.*, the M1 layer) and pre-Holocene marine sediments (*i.e.*, the M2 layer), described by Yim (1994). The heavier  $\delta^{13}C_{org}$  value observed at this interval may reflect contributions of C4 terrigenous plants (*e.g.*, C4 seagrasses) to the marine sediments. The  $\delta^{13}C_{org}$  values become progressively lighter from 3.7m towards the base of the core, where  $\delta^{13}C_{org}$  is -26.27°/<sub>oo</sub> at 3.9m, and ranges from - 33.17°/<sub>oo</sub> to -30.18°/<sub>oo</sub> between 4.0m and 4.1m. Isotopic values of organic carbon between 4.0m and 4.1m suggests predominant contributions of organic matter

from terrigenous plants.  $\delta^{15}$ N values at 3.7m and 3.9m do not deviate much from  $\delta^{15}$ N values measured in sediments deposited above these intervals, and range between  $4.26^{\circ}/_{\circ\circ}$  and  $4.51^{\circ}/_{\circ\circ}$ . At 4.0m, a sharp drop in  $\delta^{15}$ N (2.53°/<sub>00</sub>) is observed which may be reflective of terrigenous plants utilizing atmospheric nitrogen as their nitrogen source.

Crossplots of C/N (atomic mass ratio) to  $\delta^{13}C_{org}$  of plant and algal material, from lake and marine environments, have been utilized for distinguishing sources of sedimentary organic matter (**Fig. 3.7**; Meyers, 1994). The C/N atomic mass ratio is calculated by multiplying the C/N weight % ratio by 1.167 (Meyers and Teranes, 2001). The C/N (atomic mass ratios) and  $\delta^{13}C_{org}$  measurements from sediments in the Victoria Harbour core sample (MBH 54/2; **Fig. 3.8**) do not fall within the regions defined in **Fig. 3.7**. The majority of the samples have C/N (atomic mass ratios) less than 20, and have  $\delta^{13}C_{org}$  values that fall within the range for C3 terrigenous plants (**Figs. 3.8 and 3.9**). Sediments from the interval between 1.8m and 2.3m have C/N (atomic mass ratios) that fall within the range for algal-derived organic matter. The  $\delta^{13}C_{org}$  values within this interval, however, are much lighter than expected (-27.99°/<sub>oo</sub> to -26.92°/<sub>oo</sub>).

Uncontaminated sediments with isotopic compositions suggesting contributions from C3 terrigenous plants (*i.e.*,  $\delta^{13}C_{org}$  values ranging between -27.50°/<sub>oo</sub> and -24.74°/<sub>oo</sub>) fall into two identifiable groups (**Fig. 3.8**). One group has C/N ratios ranging between 12.2 and 14.2, and the second group ranges between 17.1 and 18.3. Sewage contaminated sediments in the uppermost part of the core (0.5m to 0.7m) may have been affected by denitrification processes,

resulting in the enrichment of <sup>13</sup>C (*i.e.*,  $\delta^{13}C_{org}$  values in this region range between -23.86°/<sub>oo</sub> and -22.60°/<sub>oo</sub>). In general,  $\delta^{13}C_{org}$  values of sewage contaminated sediments range from -28.59°/<sub>oo</sub> to -26.30°/<sub>oo</sub>, with C/N values between 16.0 and 17.2, except at 1.4m where there appears to have been a flux in organic matter input.

A plot of  $C_{org}$  versus atomic C/N mass ratio (**Fig. 3.10**) illustrates two envelopes separating those sediments affected or unaffected by sewage waste. Sediments in the upper 1.6m of the core are known to have been contaminated by sewage waste and contain the highest organic carbon content. Sediments contaminated



**Fig. 3.7.** Crossplot of atomic C/N mass ratio to  $\delta^{13}$ C of terrigenous plant and algal material, for differentiating sources of organic matter (from Meyers, 1994).



**Fig. 3.8.** Crossplot of C/N ratio to  $\delta^{13}C_{org}$  of sediments from core MBH 54/2.



**Fig. 3.9.** Overlay of the atomic C/N mass ratio vs  $\delta^{13}C_{org}$  of sediments from core MBH 54/2, and the Meyers (1994) plot for differentiating sources of organic matter.



Fig. 3.10. Discrimination of sewage contaminated sediments from uncontaminated sediments using the plot of  $C_{org}$  versus C/N ratio.

with sewage waste can not be differentiated from uncontaminated sediments using the plot of  $\delta^{15}N$  versus  $\delta^{13}C_{org}$  (**Fig. 3.11**). The envelopes drawn in **Fig. 3.11** are based on prior knowledge of sediments contaminated with sewage. The sewage affected sediments have  $\delta^{15}N$  values ranging between  $2.14^{\circ}/_{oo}$  and  $3.44^{\circ}/_{oo}$ , and  $\delta^{13}C_{org}$  values ranging between  $-28.59^{\circ}/_{oo}$  and  $-26.30^{\circ}/_{oo}$ . In the uppermost part of the core, the sewage contaminated sediment may have also been affected by denitrification processes, resulting in the enrichment of <sup>15</sup>N and <sup>13</sup>C.



**Fig. 3.11.** Crossplot of  $\delta^{15}$ N versus  $\delta^{13}C_{org}$  of contaminated and uncontaminated sediments from Kowloon Bay (core MBH 54/2).

## 3.4 Summary Remarks

Elemental analyses and bulk stable isotope measurements of organic carbon and nitrogen in sediments from core MBH 54/2, Kowloon Bay, provide clues for delineating sources of organic matter and provide a means for speculating on processes and events that have occurred in this region. Sediments in the upper 1.6m of the core are known to have been deposited during periods of rapid population growth and excessive disposal of raw sewage. The highest flux in organic carbon and nitrogen occur within this interval and  $\delta^{13}C_{org}$  and  $\delta^{15}N$  values (*i.e.*, -28.59°/<sub>oo</sub> to -26.30°/<sub>oo</sub> and 2.14°/<sub>oo</sub> to 3.44°/<sub>oo</sub>, respectively) are consistent with bulk isotopic compositions reported in the literature for sewage contaminated sites. The uppermost part of the core (0.5m to 0.7m) is more enriched in <sup>13</sup>C and <sup>15</sup>N, where  $\delta^{13}C_{org}$  ranged between -23.86°/<sub>oo</sub> to -22.60°/<sub>oo</sub>, and  $\delta^{15}N$  was  $6.19^{\circ}/_{oo}$ . It is speculated that this may be an indication that the organic matter was affected by denitrification processes as a result of eutrophication in Kowloon Bay. Enrichment of <sup>13</sup>C and <sup>15</sup>N in organic matter has been observed under conditions of eutrophication where increasing anoxicity results in intense denitrification processes.

Below the sewage contaminated sediments (1.8m to 2.3m), there is a shift in the C/N (weight % ratio) where organic carbon decreases relative to nitrogen. This may be an indication that the organic matter source has experienced a shift towards more aquatic-derived organic matter that would have a higher nitrogen content relative to Corg. Sediments deposited between 2.6m and 3.5m appear to be dominated by C3 terrigenous plant material with periodic shifts towards aquatic-derived organic matter (e.g., at 3.1m and 3.5m). C/N ratios and  $\delta^{13}C_{org}$ values within this interval range between 10.5 and 15.7, and  $-27.50^{\circ}/_{\circ\circ}$  and -24.74°/<sub>oo</sub>, respectively. The  $\delta^{13}C_{org}$  value is enriched in <sup>13</sup>C at 3.7m (-21.87°/<sub>oo</sub>), which is located around the desiccated crust indicating the M1-M2 boundary. The heavier  $\delta^{13}C_{org}$  value may represent contributions from C4 plants during the early Holocene/late Pleistocene, when sea level was approximately 130km south of Hong Kong. The isotopic signature at this depth may serve as a secondary parameter for identifying the boundary between Holocene marine sediments and pre-Holocene marine sediments (*i.e.*, the M1-M2 layer). Sediments at the base of the core (4.0m to 4.1m) are the most depleted in  $^{13}C$ , where  $\delta^{13}C_{\text{org}}$  ranges

between  $-33.17^{\circ}/_{\circ\circ}$  to  $-30.18^{\circ}/_{\circ\circ}$ .  $\delta^{15}$ N was only measurable at 4.0m with a value of  $2.53^{\circ}/_{\circ\circ}$ . Isotopic compositions of the sedimentary organic matter at these intervals suggest origins from C3 terrigenous plant material where atmospheric nitrogen is utilized via nitrogen fixation.

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#### **CHAPTER 4**

# Sources and Distribution of Extractable Organic Matter in Kowloon Bay Sediments, Hong Kong SAR, China

## 4.1 Introduction

Lipids are important constituents of sedimentary organic matter and can be used to delineate the sources of organic matter and transformation processes in Recent sediments. While lipids make up a minor fraction of the organic matter in Recent sediments (*i.e.*, proteins and carbohydrates are more abundant), they demonstrate a greater degree of resilience to environmental alteration (Killops and Killops, 1993; Wakeham et al., 1997; Smallwood and Wolff, 2000; Gogou and Stephanou, 2004; Muri et al., 2004). In general, lipids are defined as the fraction of organic matter extractable from biological material using organic solvents such as dichloromethane, methanol, toluene, ether, or hexane (Meyers) and Ishiwatari, 1993; Rullkötter, 2000). It should be noted that a significant portion of lipids in sedimentary organic matter occur in bound form (discussed in Chapter 5) and require harsher treatments (*e.g.*, alkaline and acid hydrolysis) in order to release ester- and amide-bound lipids (Goosens et al., 1986, 1989a,b; Cranwell, 1990; Fukushima et al., 1992a,b; Wakeham, 1999; Killops and Killops, 2005). Lipids extractable from marine sediments are comprised of sterols, fatty alcohols, fatty acids, and hydrocarbons. Each of these lipid groups provides diagnostic information that can be used to reconstruct the origin of sedimentary organic matter and transformation processes.

## 4.2 Review of Literature

## 4.2.1 Sterols in Marine Sediments

Sterols are well preserved in the environment and have unique structures, making them favorable compounds as biological markers. The basic structure and numbering scheme for sterols, using cholesterol as an example, is shown in **Fig. 4.1**.



**Fig. 4.1.** Cholesterol structure illustrating an example of the numbering scheme for sterols.

Sterols and compounds derived from sterols (*e.g.*, stanols, stenones, stanones, sterenes, and steranes; **Fig. 4.2**) have been used as proxies to determine the proportions of algal and terrigenously derived organic matter in marine sediments, study transformation processes, and to trace the origin of fecal material in the





environment (Gagosian *et al.*, 1980; Mackenzie *et al.*, 1982; Readman *et al.*, 1986; Volkman, 1986; Mudge *et al.*, 1999; Meyers, 2003).

Sterols from algal and zooplankton derived organic matter are typically dominated by cholesterol (Volkman, 1986; Santos *et al.*, 1994; Burns *et al.*, 2003; Meyers, 2003). The key sterols in vascular terrigenous plants are the C<sub>29</sub> sterols,  $\beta$ -sitosterol and stigmasterol (Huang and Meinschein, 1979; Volkman, 1986; Saliot *et al.*, 1991; Mudge and Norris, 1997). Relative contributions of aquatic versus terrigenous plant material in Recent sediments have been estimated using the C<sub>27</sub> to C<sub>29</sub> sterol ratio (Huang and Meinschein, 1979; Meyers, 2003). This ratio should be used with caution, since C<sub>29</sub> sterols have been observed to occur in certain marine organisms (Volkman, 1986; Santos *et al.*, 1994; Gogou and Stephanou, 2004). Other unique sterols include brassicasterol, a common marker for diatoms and prymnesiophytes; and dinosterol, an indicator for dinoflagellates (Gagosian *et al.*, 1980; Volkman, 1986; Santos *et al.*, 1994; Smallwood and Wolff, 2000).

Sewage contamination in the environment has been monitored and traced using sterols. Human sewage waste can be monitored in the environment using the fecal sterol, coprostanol (5 $\beta$ -cholestan-3 $\beta$ -ol; **Fig. 4.3a**). Coprostanol is a unique marker formed by the bacterial reduction of cholesterol (cholest-5-en-3 $\beta$ ol) in the human digestive tract, and released into the environment in human feces (**Fig. 4.3b**; Readman *et al.*, 1986; Nichols and Leeming, 1991; Mudge *et al.*, 1999). The process for transforming cholesterol to coprostanol is illustrated in





**Fig. 4.3.** Fecal sterol profiles from **(a)** sewage waste and **(b)** human feces (figures provided via personal communications with Dr. Rhys Leeming, CSIRO).

**Fig. 4.4**. During sewage treatment, coprostanol is transformed to its isomer epicoprostanol (5β-cholestan-3α-ol). To date, epicoprostanol is only known to be present in treated sewage sludge (McCalley *et al.*, 1981; Mudge *et al.*, 1999; and personal communication with Rhys Leeming, 2003). Leeming *et al.* (1994, 1996 and 1997) and Sinton *et al.* (1998) have demonstrated that fecal sterols can be used to distinguish human and animal feces. Herbivores (*e.g.*, sheep and cows) consume substantial amounts of plants enriched with C<sub>29</sub> sterols (*e.g.*, β-sitosterol and stigmasterol). Fecal matter released into the environment by herbivores are usually composed of considerable amounts of 24-ethylcoprostanol and 24-ethyl-epicoprostanol (Leeming *et al.*, 1997).



**Fig. 4.4.** Biohydrogenation of cholesterol to coprostanol in the human digestive tract (based on Björkhem and Gustafsson, 1971).

## 4.2.2 Fatty Alcohols in Marine Sediments

Fatty alcohols have been observed in marine sediments but have not been utilized to the same extent as fatty acids and aliphatic hydrocarbons (Mudge and Norris, 1997; Meyers, 2003). Organic matter derived from terrigenous or marine sources have been differentiated using fatty alcohols (Mudge and Norris, 1997; Mudge and Seguel, 1999; Meyer, 2003). In terrigenous plants, fatty alcohols ( $C_{22}$  to  $C_{30}$ ) occur as wax esters in epicuticular waxes with an even-over-odd carbon preference (Grimalt and Albaigés, 1990; Mudge and Norris, 1997; Mudge and Seguel, 1999; Meyers, 2003). Leaf waxes are generally dominated by  $C_{24}$ ,  $C_{26}$ , and  $C_{28}$  alcohols (Mudge and Norris, 1997; Fernandes *et al.*, 1999). Their primary function in terrigenous plant leaves is to retain water (Eglinton and Hamilton, 1967; Mudge and Norris, 1997; Mudge and Seguel, 1999).

Short chain fatty alcohols (<C<sub>20</sub>) are associated with organic matter derived from aquatic algae and/or bacterial sources. It has been suggested that wax esters are synthesized by marine animals to serve as energy reserves during periods of food shortages (Lee and Hirota, 1976; Mudge and Norris, 1997). Fatty alcohols in marine wax esters are normally dominated by saturated alcohols (*i.e.*, C<sub>14:0</sub> and C<sub>16:0</sub>) or monounsaturated alcohols (*e.g.*, C<sub>16:1</sub>, C<sub>18:1</sub>, C<sub>20:1</sub>, and C<sub>22:1</sub>) (Sargent *et al.*, 1977, 1981; Rajendran *et al.*, 1991; Mudge and Norris, 1997).

## 4.2.3 Fatty Acids in Marine Sediments

Fatty acids are ubiquitous in the environment and are common constituents in bacteria, microalgae, terrigenous plants, and marine plants (Volkman *et al.*, 1998; Mudge and Seguel, 1999; Meyers, 2003). They serve important roles in energy storage and are involved in the structure of cellular membranes (Ratledge and Wilkinson, 1988; Budge and Parrish, 1998). Two common applications of fatty acids include determination of the sources of organic matter and studying transformation processes (Budge and Parrish, 1998; Wakeham, 1999). Terrigenously derived fatty acids consist of long-chain monocarboxylic acids (> $C_{20:0}$ ; with an even-over-odd carbon preference), and are associated with epicuticular waxes of higher plants. Both n-alkanoic and nalkenoic acids < $C_{20}$  (with an even-over-odd preference) have been used as indicators for planktonic and bacterial input (Grimalt and Albaigés, 1990; Saliot *et al.*, 1991; Budge and Parrish, 1998; Volkman *et al.*, 1998; Meyers, 2003).

The diverse range of fatty acid structures allow certain fatty acids to be used as markers for specific organisms (Budge and Parrish, 1998). Branched fatty acids, in particular iso- (i-) and anteiso- (ai-)  $C_{15:0}$  and  $C_{17:0}$ , along with their corresponding unsaturated branched fatty acids are considered to be unique constituents of bacteria. Their occurrence in Recent sediments have been used as indications of bacterial activity (Cranwell, 1973; Saliot *et al.*, 1991; Budge and Parrish, 1998; Grimalt and Albaigés, 1990). Markers for planktonic input consist of mixtures of  $C_{14:0}$ ,  $C_{16:0}$ ,  $C_{16:1\omega7}$ ,  $C_{18:1\omega9}$ ,  $C_{18:0}$ ,  $C_{20:5\omega3}$ , and  $C_{22:6\omega3}/C_{20:5\omega3}$ 

ratio to distinguish between dinoflagellates and diatoms, where  $C_{22:6\omega3}/C_{20:5\omega3} > 1$ suggests the predominance of dinoflagellates and  $C_{22:6\omega3}/C_{20:5\omega3} << 1$  is indicative of diatoms. Additional indicators of diatoms include the occurrence of  $C_{16:4\omega1}$  (which is common in diatoms, but rare in other phytoplankton); an abundance of  $C_{16:1}$  relative to  $C_{16:0}$  (*e.g.*,  $C_{16:1}/C_{16:0} > 1.6$ ); and an abundance of  $C_{16}$  fatty acids relative to  $C_{18}$  fatty acids (*i.e.*,  $\Sigma C_{16(saturated + unsaturated)}/\Sigma C_{18(saturated + unsaturated)})$  (Saliot *et al.*, 1991; Mudge and Seguel, 1997; Budge and Parrish, 1998; Mudge and Seguel, 1999).

## 4.2.4 Hydrocarbons in Recent Sediments

Aliphatic hydrocarbons in Recent sediments are more resistant to microbial degradation than other types of organic matter and can be used to delineate source information. In studies, primarily around lakes, aliphatic hydrocarbons have been used to distinguish between sources of organic matter commonly found within lakes (*e.g.*, algae, bacteria, and vascular plants) and vascular plants surrounding the lakes (Meyers, 2003). Marine phytoplankton and bacteria have hydrocarbons that maximize around  $C_{17}$  (Cranwell *et al.*, 1987; Mudge and Seguel, 1999; Meyers, 2003). Vascular plants, on the other hand, are dominated by longer chain n-alkanes with an odd-over-even predominance pattern (Cranwell, 1978; Santos *et al.*, 1994). The type of vascular plant can be differentiated based on alkane distributions. Submerged and floating macrophytes in lakes have n-alkanes that maximize around  $C_{21}$ ,  $C_{23}$ , or  $C_{25}$ 

(Cranwell, 1984; Ficken *et al.*, 2000; Meyers, 2003). Terrestrially derived vascular plants are dominated by  $C_{27}$ ,  $C_{29}$ , and  $C_{31}$  n-alkanes, where n- $C_{27}$  and n- $C_{29}$  are indicators for trees and n- $C_{31}$  is dominant in grasses (Meyers, 2003).

### 4.3 Results and Discussion

Free lipids were extracted from sediment samples taken from core MBH 54/2, from Kowloon Bay, in Victoria Harbour, Hong Kong. Procedures used to recover non-saponifiable and saponifiable fractions are discussed in Chapter 2. Samples were derivatized to produce trimethylsilyl ethers (for sterols, stanols, and alcohols) and methyl esters (for fatty acids), prior to analysis by gas chromatography-mass spectrometry. Identification of lipid components was based on mass spectra interpretation, comparison to published spectra, and retention times. The extractable lipids provide information that can be used to determine the sources of sedimentary organic matter and to infer environmental conditions. Identified lipids are discussed in the following sections. Sterol structures, spectra, chromatograms, and tables summarizing lipid ratios are located in **Appendix II, III, and IV**. Examples of fatty acid and alcohol structures can be found in Chapter 5, Fig. 5.2.

## 4.3.1 Sterols and Stanols

Sterols and stanols were detected in sediments from core MBH 54/2. This area in Kowloon Bay has received significant amounts of raw sewage and is thought to be highly anoxic. The downcore distribution of sterols and stanols is illustrated in **Fig. 4.5**. Sediments from the uppermost part of the core (**Fig. 4.5a**) were deposited after the sewage outfall in Hong Kong was converted to a submarine-type outfall and diverted further into the channel of Victoria Harbour. No significant amount of fecal sterols were identified at 0.5m, possibly suggesting that there were improvements in reducing sewage contamination in this part of the harbour. Cholesterol, cholestanol, brassicasterol,  $\beta$ -sitosterol, and stigmastanol were also identified at 0.5m.

Sediments deposited during a period of rapid population growth can be recognized by the chromatograms shown in **Figs. 4.5 b and c**. At the time sediments were deposited at 1.1m and 1.6m, raw sewage was discharged into the harbour via a seawall-type sewage outfall. Significant amounts of fecal sterols were observed in the extractable lipid fraction. Fecal material derived from human sewage waste is indicated by the occurrence of coprostanol. The isomer of coprostanol, epicoprostanol, is commonly used as a ratio with coprostanol as an indicator for the degree of sewage treatment (*i.e.* a higher epicoprostanol to coprostanol ratio would indicate that the majority of sewage is treated before being released into the environment) (McCalley *et al.*, 1981; Mudge and Seguel, 1999).



**Fig. 4.5.** Distribution of sterols and stanols in sediment samples from core MBH 54/2. (1) coprostanol; (2) epicoprostanol; (3) cholesterol; (4) cholestanol; (5) brassicasterol; (6) 24-ethylcoprostanol; (7) 24-ethylepicoprostanol; (8) campesterol; (9) ergostanol (campestanol); (10)  $\beta$ -sitosterol; (11) stigmastanol

The occurrence of epicoprostanol throughout the core is peculiar since there were no sewage treatment plants in Hong Kong during these periods. Bacterial populations in the human digestive tract preferentially mediate cholesterol to coprostanol, but not to epicoprostanol or cholestanol. The epicoprostanol in core MBH 54/2 was most likely produced by the anaerobic bacterial community within the sediments. Both coprostanol and epicoprostanol are more abundant than cholesterol at 1.1m and 1.6m (**Fig. 4.6a**). Similar to the human fecal sterols, 24-ethylcoprostanol and 24-ethylepicoprostanol (*i.e.,* fecal markers for herbivores) are more abundant than cholesterol at 1.1m and 1.6m (**Fig. 4.1**).

Brassicasterol and campesterol were not detected at 1.1m or 1.6m (**Figs. 4.5 b, c**; and **Fig. 4.6b**). Brassicasterol is the major sterol in the algae Prymnesiophycean and is commonly used as a biomarker for diatoms (Volkman, 1986). Campesterol is widespread in vascular plants (Mudge and Seguel, 1999), but has also been observed in green algae and marine invertebrates (Goad and Akihisa, 1997). Both brassicasterol and campesterol were only observed at 0.5m, 2.3m, and 3.4m (**Figs. 4.5 b, d, and e**). Depth intervals around 2.3m and 3.4m represent periods when Victoria Harbour was not affected by sewage waste; around 0.5m the sewage was diverted away from the study site. The diminished sewage contamination appears to have improved conditions, allowing marine organisms to bloom.





Fig. 4.6. Stanol and sterol ratios in sediment samples from core MBH 54/2.

Cholestanol and campestanol increase downcore (between 1.0m and 2.0m) relative to their corresponding sterols. The ratios of cholestanol/cholesterol and campestanol/campesterol show similar patterns downcore (Fig. 4.7). These stanols are likely derived from the hydrogenation of their corresponding sterols (Gaskell and Eglinton, 1975; Nishumira and Koyama, 1977; Pinturier-Geiss et al., 2002). Gaskell and Eglinton (1975) have observed, and demonstrated experimentally with radiolabelled sterols, that sterols undergo rapid hydrogenation in anoxic Recent sediments. Gagosian et al. (1980) noted that while anaerobic conditions inhibit sterol degradation, sterol to stanol reduction is accelerated. At depth intervals between 1.0m and 2.0m, the study site received substantial amounts of raw sewage, which probably resulted in highly anoxic conditions. Between 2.0m and 3.0m, Kowloon Bay was an open bay and did not receive significant amounts of raw sewage. Conditions during this period may have been less anoxic resulting in the decrease in stanol/sterol ratios observed in **Fig. 4.7**. The slight increase in stanol/sterol ratio at 3.4m suggests a shift towards slightly more anoxic conditions.



**Fig. 4.7.** Hydrogenation of sterols to stanols in sediments deposited in Kowloon Bay.

## 4.3.2 Fatty Acids

Free n-alkanoic acids were identified in both saponifiable and nonsaponifiable fractions. In the saponifiable fraction (**Fig. 4.8**), n-alkanoic acids were distributed between  $C_{12:0}$  and  $C_{34:0}$ , with a distinct even-over-odd predominance pattern (average CPI=7.8). A slightly higher even preference is observed at depths of 1.1m (CPI=10.4) and 1.6m (CPI=9.5). Throughout the core, a bimodal distribution is observed, with short chain n-alkanoic acids predominating over long chain n-alkanoic acids. The short chain n-alkanoic acids maximize at n-C<sub>16:0</sub>, and long chain n-alkanoic acids at n-C<sub>30:0</sub> around 1.1m and n-C<sub>22:0</sub> or n-C<sub>24:0</sub> at all other depths. Short-chain n-alkanoic acids (<n-C<sub>20:0</sub>) are



**Fig. 4.8.** n-Alkanoic acids (as methyl esters) in the saponifiable fraction of extractable lipids from two depth intervals in core MBH 54/2.

attributed to planktonic and bacterial input, whereas longer chain alkanoic acids  $(>n-C_{20:0})$  originate from the cuticular waxes of terrigenous plant material (Grimalt and Albaigés, 1990).

A plot of the aquatic-to-terrigenous ratio (*i.e.,* short-chain to long-chain alkanoic acids; *e.g.,*  $\Sigma(C_{12:0}-C_{18:0})/\Sigma(C_{22:0}-C_{28:0})$ ) downcore illustrates a higher aquatic input from the surface down to 1.6m ( $\Sigma(C_{12:0}-C_{18:0})/\Sigma(C_{22:0}-C_{28:0})$ > 1.5). At depths below 2.3m, terrigenous plant material are more abundant than

aquatically-derived organic matter (**Fig. 4.9a**;  $\Sigma(C_{12:0}-C_{18:0})/\Sigma(C_{22:0}-C_{28:0}) \leq 1$ ). Iso- and anteiso- alkanoic acids ( $C_{13:0}$ ,  $C_{15:0}$ , and  $C_{17:0}$ ) have been used as indicators of bacterial activity. The most abundant branched acids are *i*- and *ai*- $C_{15:0}$ , where *i*- $C_{15:0}$  dominates over *ai*- $C_{15:0}$  downcore (avg *i*-/*ai*-  $C_{15:0}$  ratio =1.2). Aside from indicating the presence of bacteria, the branched fatty acids identified in the free lipid fraction provide limited information about the bacteria types.



**Fig. 4.9.** Aquatic-to-Terrigenous ratio for free n-alkanoic acids in the (a) saponifiable and (b) non-saponifiable fractions. Short-chain alkanoic acids are typically more abundant than long-chain alkanoic acids, and are attributed to planktonic and bacterial input. A greater abundance of short-chain acids is observed at 1.1m and 1.6m.

Alkenoic acids ( $C_{16:1}$  and  $C_{18:1}$ ) were only observed in the upper 1.6m, and are present in low abundance relative to their corresponding *n*-alkanoic acids. Two peaks in the chromatograms were identified as  $C_{18}$  alkenoic acids with 1-degree of unsaturation (the location of the double bond was not determined). The  $C_{16:1}/C_{16:0}$  and  $\Sigma C_{18:1}/C_{18:0}$  ratios are lowest at 1.1m (0.006 and 0.018, respectively), and highest at 1.6m ( $C_{16:1}/C_{16:0}$ =0.41) and 0.5m

( $\Sigma C_{18:1}/C_{18:0}=0.065$ ). The ratio of  $C_{16:1}/C_{16:0}$  has been used to measure the predominance of diatoms in sediments (Saliot *et al.* 1991; Santos *et al.*, 1994; Budge and Parrish, 1998; Mudge and Seguel, 1999; Azevedo, 2003; Burns *et al.*, 2003) whereas  $C_{18:1}$  is commonly used as an indicator for zooplankton (Santos *et al.*, 1994; Azevedo, 2003; Burns *et al.*, 2003).  $C_{16:1}$  and  $C_{18:1}$ , however, have also been observed in certain species of sulfate-reducing bacteria (Edlund *et al.*, 1985; Wilkinson, 1988), which may be the primary source of these acids in sediments from Kowloon Bay.

Free n-alkanoic acids in the non-saponifiable fraction had carbon number distributions between n-C<sub>14:0</sub> to n-C<sub>26:0</sub> with an even carbon number preference (**Fig. 4.10**). CPI typically ranged between 7.0 and 9.6, except at 1.6m, where a more pronounced even preference is observed (CPI=15.9). Short-chain alkanoic acids (<n-C<sub>20:0</sub>) are present at significantly higher abundance than long-chain alkanoic acids (>n-C<sub>20:0</sub>), and typically show a high aquatic-to-terrigenous ratio ( $\Sigma$ (C<sub>14:0</sub>-C<sub>18:0</sub>)/ $\Sigma$ (C<sub>20:0</sub>-C<sub>24:0</sub>) >3.5) downcore. Aquatic input is significantly higher between 1.1m and 1.6m, where the average aquatic/terrigenous ratio is 7.3 (**Fig. 4.9b**).


**Fig. 4.10.** n-Alkanoic acids (as trimethylsilyl ethers) in the non-saponifiable fraction of extractable lipids from two depth intervals in core MBH 54/2.

Monounsaturated fatty acids ( $C_{16:1}$  and  $C_{18:1}$ ) in the non-saponifiable fraction are more abundant than in the saponifiable fraction. Ratios of  $C_{16:1}/C_{16:0}$  and  $\Sigma C_{18:1}/C_{18:0}$  are significantly higher at a depth of 1.6m (0.07 and 0.3, respectively).  $C_{16:1}$  is not observed in the non-saponifiable fraction below 2.3m. *i*- and *ai*- $C_{15:0}$ and  $C_{17:0}$  are observed downcore with greater abundance of *i*- and *ai*- $C_{15:0}$ . At all depths, *ai*- $C_{15:0}$  is more abundant than *i*- $C_{15:0}$  (where *i*-*/ai*-  $C_{15:0}$  range between 0.5-0.8), except 2.3m (where *i*-*/ai*-  $C_{15:0}$ =1.1), possibly indicating the presence of the sulfate reducing bacteria *D. gigas* (Vainshtein *et al.*, 1992). The ratio of *i-/ai*- $C_{17:0}$  in the non-saponifiable fraction demonstrates a very similar pattern to that observed in the saponifiable fraction. In both fractions, *i-/ai*- $C_{17:0}$  ratio is about 1.1 at depths of 0.5m and 1.6m. At all other depths, the *i-/ai*- $C_{17:0}$  ratio ranges between 0.7-0.9. *i-* and *ai*- $C_{17:0}$  are markers for sulfate reducing bacteria and other anaerobic bacteria (Rajendran *et al.*, 1992a and b).

# 4.3.3 Alcohols

Free *n*-alcohols (C<sub>14</sub> to C<sub>32</sub>) in the non-saponifiable fraction (**Fig. 4.11**) have an even carbon-number preference downcore. Short-chain *n*-alcohols (C<sub>14</sub>-C<sub>20</sub>) represent a minor source of alcohols and may originate from zooplankton or other marine invertebrates (Grimalt and Albaigés, 1990). The long-chain *n*-alcohols (C<sub>22</sub>-C<sub>32</sub>) are more abundant and suggest cuticle waxes of terrigenous plant material (Grimalt and Albaigés, 1990; Santos *et al.*, 1994; Mudge and Norris, 1997; Mudge and Seguel, 1999) are more important sources of alcohols in these sediments. The aquatic-to-terrigenous ratio (C<sub>18</sub>/C<sub>28</sub> <1) supports the idea that terrestrial plants are the primary source of alcohols in Kowloon Bay. In general, the aquatic-to-terrigenous ratio ranged between 0.4 and 0.6, except at 1.1m where alcohols were dominated by C<sub>28</sub>, C<sub>30</sub>, and C<sub>32</sub> (**Fig. 4.12**). The C<sub>16</sub>-C<sub>26</sub> alcohols were only present as very small peaks, near the baseline of the chromatogram, in sediments around 1.1m. At 1.6m, the alcohols may have been degraded, although C<sub>16</sub>, C<sub>18</sub>, C<sub>28</sub>, and C<sub>30</sub> alcohols were still identifiable.



**Fig. 4.11.** Fatty alcohols (as trimethylsilyl ethers) in the non-saponifiable fraction of extractable lipids, from core MBH 54/2 (CPI~8).



**Fig. 4.12.** Aquatic-to-Terrigenous ratio for free fatty alcohols. Downcore distribution indicates cuticle waxes of higher plants are the primary source for alcohols.

# 4.3.4 Hydrocarbons

A distinct odd-over-even preference is observed for n-alkanes (n-C<sub>18</sub> to n-C<sub>37</sub>) at all depths (Fig. 4.13). CPI values for n-alkanes ranged between 2.2 and 3.0, except at 1.1m and 1.6m where CPI was about 1.2. The distinct odd predominance suggests that the free n-alkanes were most likely derived from biotic hydrocarbons rather than petroleum hydrocarbons. The aquatic-toterrigenous ratio for free *n*-alkanes  $(n-C_{19}/n-C_{31})$  ranged between 0.1 to 0.2, except at 1.1m and 1.6m where  $n-C_{19}/n-C_{31}$  was about 0.6 (Fig. 4.14a). n-Alkanes in sediments around 1.1m and 1.6m were partially degraded, as was observed for n-alcohols at these depths. Cuticle waxes of higher plants appear to be the primary contributors of n-alkanes and can be further subdivided into those derived from terrigenous plants (maximum at n-C<sub>29</sub> and n-C<sub>31</sub>) and macrophyte plant material (n-C<sub>23</sub> and n-C<sub>25</sub>) (Ficken et al., 2000; Filley et al., 2001; Silliman and Schelske, 2003). Proxy ratios based on mid-chain ( $C_{23}$ ,  $C_{25}$ ) to long chain  $(C_{29}, C_{31})$  n-alkanes have been used to distinguish between macrophyte and terrigenous plant input (Ficken et al., 2000; Filley et al., 2001; Silliman and Schelske, 2003). The downcore profile of this proxy shows a slight increase at 1.1m and 1.6m (0.8 and 1.0, respectively). All other depths appear to be dominated by free n-alkanes derived from terrigenous plants based on the  $(C_{23}+C_{25})/(C_{29}+C_{31})$  ratio between 0.3 to 0.6 (**Fig. 4.14b**). Fisher *et al.* (2003) have used the mean carbon number (MC#) parameter  $(\Sigma([C_i]*C_i)/\Sigma[C_i], where [C_i])$ is the amount of *n*-alkane with carbon number  $C_i$ , and  $C_i$  ranges between  $C_{27}$ - $C_{31}$ ) to reflect slight changes in vegetation. Grasses are dominated by  $C_{31}$  n-alkanes,



**Fig. 4.13.** n-Alkane distribution in the extractable lipid fraction from core MBH 54/2, illustrating a pronounce odd-over-even predominance pattern (CPI ~ 3.0).



**Fig. 4.14. (a)** Aquatic-to-Terrigenous ratio for free n-alkanes; **(b)** Downcore profile illustrating shifts in predominance of macrophyte versus terrigenous plant material; **(c)** Downcore shifts in vegetation type, based on the mean carbon number ranging between  $C_{27}$ - $C_{31}$ .

while trees tend to contribute leaves with significant amounts of  $C_{27}$  and  $C_{29}$  nalkanes (Cranwell, 1973; Fisher *et al.*, 2003; Meyers, 2003). A plot of the MC# downcore shows a shift from 29.4 to about 28.7 between 1.1m and 1.6m (**Fig. 4.14c**). The shift towards a MC# of 28.7 is due to a decrease in the C<sub>31</sub> n-alkane compared to C<sub>27</sub> and C<sub>29</sub>. Less grass and possibly more terrestrial plant leaves were transported into Kowloon Bay between 1.1m and 1.6m. This may support the idea of storms transporting higher plant material into this area, indicated by a spike in the C/N ratio around 1.4m (discussed in Chapter 3).

## 4.4 Summary Remarks

Extractable lipids in core MBH 54/2, Kowloon Bay, are comprised of sterols, fatty alcohols, fatty acids, and hydrocarbons. Each of these lipid groups (*i.e.,* the lipid composition and profile) provide information that allow the sources of sedimentary organic matter to be determined, and to also infer conditions and transformation processes that may have occurred. Stanols were more abundant than their corresponding sterols suggesting that the sterols had been hydrogenated to stanols under anaerobic conditions.

Significant amounts of fecal sterols (*i.e.*, coprostanol, epicoprostanol, 24ethylcoprostanol, and 24-ethylepicoprostanol) were identified in the upper 2m of the core, maximizing at 1.1m and 1.6m. These depth intervals correspond to periods of rapid population growth in Hong Kong and also during a time when

raw sewage was disposed directly over the sample site via seawall-type sewage outfall. Of the fecal sterols extracted, coprostanol (a marker for human feces) was most abundant. 24-Ethylcoprostanol, a lipid marker for herbivores (*e.g.*, cows and sheep), was also very abundant in the sediment. The occurrence of epicoprostanol in the sediments was unusual since it has only been observed in treated sewage. Microbes in the human intestine are not known to be able to hydrogenate cholesterol to epicoprostanol and no direct evidence of bacteria mediating cholesterol to epicoprostanol has been documented in the literature. Since sewage treatment plants were not operating in Hong Kong when epicoprostanol was detected in the sediments, it is proposed that the microbes in the anaerobic sediments were probably responsible for transforming cholesterol to epicoprostanol.

n-Alkanoic acids were identified in both saponifiable and non-saponifiable fractions of extractable lipids. A bimodal distribution of even carbon numbered nalkanoic acids ( $C_{12:0}$  to  $C_{34:0}$ ) were detected in the saponifiable lipid fraction. The short chain components ( $<C_{20:0}$ ) are attributed to bacterial and/or planktonic input, and the long chain components ( $>C_{20:0}$ ) are associated with cuticular waxes of terrigenous plants. Iso- and anteiso-alkanoic acids ( $C_{13:0}$ ,  $C_{15:0}$ , and  $C_{17:0}$ ) were identified in the sediments and are used as general markers for bacteria. In the non-saponifiable fraction, n-alkanoic acids ranged between  $C_{14:0}$  to  $C_{26:0}$ . However,  $C_{16:0}$  and  $C_{18:0}$  were dominant in each of the samples.

n-Alcohols ranged between  $C_{14}$  and  $C_{32}$ , with an even carbon preference. The longer chain n-alcohols ( $C_{22}$  to  $C_{32}$ ) were more abundant throughout the core,

suggesting that terrigenous plants are the primary source of alcohols in Kowloon

Bay. n-Alkanes ranged between  $C_{18}$  and  $C_{37}$ , with distinct odd carbon preference.

The n-alkanes are derived from a biotic source where cuticular waxes of

terrigenous plants (*e.g.*, terrestrial plant leaves and grasses) are the likely

contributors.

# 4.5 References

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### **CHAPTER 5**

The Occurrence and Distribution of Microbial Markers in Ester- and Amide-Bound Lipid Fractions from Kowloon Bay, Hong Kong SAR, China

## 5.1 Introduction

Ester- and amide-bound lipids in Recent sediments are well preserved in sedimentary organic matter and provide a record of different sources of input in the biogeochemical record. While various research groups have suggested that bound lipids can provide more detailed information than freely extractable lipids for characterizing sedimentary organic matter, bound lipids have not been widely utilized (Wakeham, 1999; Stefanova and Disnar, 2000; Zegouagh et al., 2000; Garcette-Lepecq et al., 2004). Recovery of bound lipids from Recent sediments requires harsher treatments than conventional solvent extraction. Ester-bound lipids are freed from solvent extracted sediments via alkaline hydrolysis followed by solvent extraction. The amide-bound lipids are freed by acid hydrolysis and subsequent solvent extraction (Goosens et al., 1986, 1989a,b; Cranwell, 1990; Fukushima et al., 1992a,b; Wakeham 1999; Garcette-Lepecq et al., 2004). Unlike free lipids, bound lipids are sterically protected and resilient to chemical and diagenetic degradation (Kawamura and Ishiwatari, 1984; Kawamura et al., 1986; Zegouagh et al., 2000). Preserved organic material in sediments can be utilized to reconstruct the origin and diagenetic pathways from which the organic matter was derived.

A key interest in identifying the types of bacteria present in sediments from core MBH 54/2 is that significant amounts of methane have been released from sediments in Victoria Harbour during dredging activities. Since no methane was detected in sediments from core MBH 54/2, markers identifying the bacterial community may provide a better understanding of organic matter remineralization and possible methane generation. The general sequence of organic matter remineralization processes is summarized in **Fig. 5.1**, starting with aerobic respiration. Once  $O_2$  has been consumed, denitrification occurs followed by manganese and iron respiration. Methane generation via methanogenesis will occur after the sulfate oxygen has been consumed by sulfate reduction (Froelich *et al.*, 1979; Jørgensen, 2000).

Pathy	vays of organic matter remineralization:
Oxic respiration:	$CH_2O + O_2 \rightarrow CO_2 + H_2O$
Denitrification:	$5CH_2O + 4NO_3^- \rightarrow 2N_2 + 4HCO_3^- + CO_2 + 3H_2O$
Mn(IV) reduction:	$CH_{2}O + 3CO_{2} + H_{2}O + 2MnO_{2} \rightarrow 2Mn^{2+} + 4HCO_{3}^{-}$
Fe(III) reduction:	$CH_{2}O + 7CO_{2} + 4Fe(OH)_{3} \rightarrow 4Fe^{2+} + 8HCO_{3}^{-} + 3H_{2}O$
Sulfate reduction:	$2CH_{2}O + SO_{4}^{2-} \rightarrow H_{2}S + 2HCO_{3}^{-}$ $4H_{2} + SO_{4}^{2-} + H^{+} \rightarrow HS^{-} + 4H_{2}O$ $CH_{3}COO^{-} + SO_{4}^{2-} + 2H^{+} \rightarrow 2CO_{2} + HS^{-} + 2H_{2}O$
Methanogenesis:	$4H_{2} + HCO_{3}^{-} + H^{+} \rightarrow CH_{4} + 3H_{2}O$ $CH_{3}COO^{-} + H^{+} \rightarrow CH_{4} + CO_{2}$
Methane Oxidation:	$CH_4 + SO_4 \rightarrow HCO_3^- + HS^- + H_2O$

**Fig. 5.1**. Generic reactions summarizing steps in the remineralization of organic matter (from Jørgensen, 2000).

Sulfate reducing and methanogenic bacteria often coexist in anoxic marine sediments (Martens and Berner, 1974; Barnes and Goldberg, 1976) where the sulfate reducing bacteria play two important roles. First, sulfate reducing bacteria are important for removing sulfate from the system, allowing methanogenesis to proceed (Martens and Berner, 1974; Barnes and Goldberg, 1976; Martens and Klump, 1984). Secondly, sulfate reducing bacteria (along with other methanogens) can also reduce the flux of methane from marine sediments by anaerobic methane oxidation (Barnes and Goldberg, 1976; Reeburgh, 1980; Valentine and Reeburgh, 2000). This chapter discusses the occurrence and distribution of ester- and amide-bound lipids recovered from sediments in core MBH 54/2, from Kowloon Bay, in the Hong Kong Special Administrative Region of China. Lipid groups in the ester- and amide-bound lipid fractions may provide insights into the types of bacterial communities that were present and possible roles they may have played in the remineralization of organic matter.

### 5.2 Literature Review

### 5.2.1 Ester- and Amide-Bound Lipids in Sedimentary Organic Matter

Bound lipids in sedimentary organic matter are typically dominated by bacterial lipids. Compounds observed in ester- and amide-bound lipid fractions include carboxylic acids, alcohols,  $\beta$ -hydroxy fatty acids,  $\alpha$ -hydroxy fatty acids,  $\omega$ hydroxy fatty acids, ( $\omega$ -1)-hydroxy fatty acids, and  $\alpha$ , $\omega$ -dicarboxylic fatty acids (**Fig. 5.2**; Kawamura and Ishiwatari, 1984; Goosens *et al.*, 1989b; Fukushima *et* 

*al.*, 1992a; Skerratt *et al.*, 1992; Wakeham, 1999). Bacterial fatty acid profiles have been described by Boon *et al.* (1977), Edlund *et al.* (1985), Dowling *et al.* (1986), Ratledge and Wilkinson (1988), Vainshtein *et al.* (1992), and others. However, limited information has been documented for bacterial fatty acid profiles isolated from





sediments (Zelles *et al.*, 1995). Other sources of bound lipids that have been recognized include those derived from terrigenous plant material, algae, and formed as intermediaries during oxidative transformations (Cardoso *et al.*, 1977; Albaigés *et al.*, 1984; Cranwell, 1984; Kawamura and Ishiwatari, 1984; Fukushima *et al.*, 1992a,b; Wakeham, 1999; Garcette-Lepecq *et al.*, 2004).

## 5.2.2 Carboxylic Acids

Carboxylic acids are ubiquitous in the environment and occur in free, ester-, and amide-bound forms. In free form, the low molecular weight monocarboxylic acids ( $<C_{20:0}$ ) are more susceptible to degradation and can appear to be less abundant than the high molecular weight monocarboxylic acids (>C<sub>20:0</sub>; Wünsche et al., 1988). Carboxylic acids in the ester- or amide-bound forms are more resilient to degradation, and when utilized with free carboxylic acids can provide a better assessment of the original source material (Farrington and Quinn, 1973; Cranwell, 1984; Goosens et al., 1989b; Garcette-Lepecq et al., 2004). General source information can be obtained by the distribution and abundance of monocarboxylic acids in the ester- and amide-bound forms. Low molecular weight monocarboxylic acids ( $C_{12:0}$  to  $C_{20:0}$ ) have been used as indicators for bacterial and algal source material, whereas iso- and anteisocarboxylic acids (e.g., i- and ai- $C_{13:0}$ ,  $C_{15:0}$ , and  $C_{17:0}$ ) have been utilized as evidence for bacterial input (Cranwell, 1978; Kawamura and Ishiwatari, 1984). As was the case for free lipids, high molecular weight monocarboxylic acids ( $C_{20:0}$  to

 $C_{30:0}$ ) are derived from terrigenous sources (Cranwell, 1978; Kawamura and Ishiwatari, 1984).

# 5.2.3 $\alpha$ - and $\beta$ -Hydroxy Fatty Acids

α- and β-Hydroxy fatty acids (**Fig. 5.2**) are more abundant in sediments in ester- and amide-bound form than as free lipids (Kawamura and Ishiwatari, 1984; Fukushima *et al.*, 1992b; Wakeham, 1999). In free lipids, the occurrence of αand β-hydroxy fatty acids is typically attributed to intermediates formed during the oxidative degradation of fatty acids (Goosens *et al.*, 1986; Wakeham, 1999). Hydroxy acids, formed as intermediates in the oxidative degradation of fatty acids, preferentially follow the β-oxidation pathway rather than the α-oxidation pathway (Lehninger, 1975; Wakeham, 1999; Garcette-Lepecq, 2004). Distribution patterns of α- and β-hydroxy fatty acids, formed as intermediates, will parallel those of their precursor monocarboxylic fatty acids (Wakeham, 1999).

Significant amounts of  $\alpha$ - and  $\beta$ -hydroxy fatty acids bound by esterlinkages are released by saponification of solvent extracted sediments (Cranwell, 1978; Kawamura and Ishiwatari, 1984; Goosens *et al.*, 1986; Garcette-Lepecq *et al.*, 2004). Ester-bound hydroxy acids are thought to originate from biotic sources, where short-chain  $\alpha$ - and  $\beta$ -hydroxy fatty acids are representative of bacterial sources and the longer chain homologues are characteristic of terrigenous sources (Kawamura and Ishiwatari, 1984; Wünsche *et al.*, 1987). Further acid hydrolysis of residual sediments releases amide-bound lipid

moieties (Klok *et al.*, 1984; Goosens *et al.*, 1986, 1989a,b; Mendoza *et al.*, 1987; Fukushima *et al.*, 1992a; Garcette-Lepecq *et al.*, 2004). In general,  $\beta$ -hydroxy fatty acids in sediments are more abundant in the amide-bound form than as ester-linked compounds, and are thought to be intact cellular remains of bacteria (Kawamura and Ishiwatari, 1982; Goosens *et al.*, 1986; Wakeham, 1999).

## 5.2.4 $\omega$ - and ( $\omega$ -1)-Hydroxy Fatty Acids

Fatty acids hydroxylated at the  $\omega$ - and ( $\omega$ -1)- positions (**Fig. 5.2**) have been observed in the ester- or amide-bound lipid fractions in sediments. The  $\omega$ hydroxy fatty acids are common constituents of cutin and suberin, and have been used as indicators for terrigenous plant material (Kawamura and Ishiwatari, 1984; Wünsche *et al.*, 1987; Fukushima *et al.*, 1992a,b; Garcette-Lepecq *et al.*, 2004). C<sub>16</sub> and C<sub>18</sub>  $\omega$ -hydroxy fatty acids are the most common acids in cutin, whereas the longer chain  $\omega$ -hydroxy fatty acids (*e.g.*, C<sub>22</sub> and C<sub>24</sub>) are more prevalent in suberin (Cardoso *et al.*, 1977). An alternative source of  $\omega$ -hydroxy fatty acids is the microbial oxidation of n-carboxylic acids at the terminal end.  $\omega$ -Hydroxy fatty acids derived from microbial oxidation reactions can be recognized by distribution patterns analogous to the distribution patterns of their precursor n-carboxylic acids (Wakeham, 1999; Garcette-Lepecq *et al.*, 2004).

Even numbered ( $\omega$ -1)-hydroxy fatty acids in sediments are thought to be derived directly from methanotrophic bacteria or indirectly through the microbial hydroxylation of monocarboxylic acids at the carbon adjacent to the terminal end (typically via some type of aerobic microorganism; Skerratt *et al.*, 1992; Wakeham, 1999). Seagrasses and cuticles of bryophytes have also been speculated as possible sources of ( $\omega$ -1)-hydroxy fatty acids (Skerratt *et al.*, 1992).

## 5.2.5 $\alpha$ , $\omega$ -Dicarboxylic Acids

α,ω-Dicarboxylic acids (**Fig. 5.2**) with carbon numbers distributed between  $C_{10}$  and  $C_{30}$  have been observed in sediments (Cranwell, 1977; Wakeham, 1999; Stefanova and Disnar, 2000). Reported origins for α,ω-dicarboxylic acids include biosynthesis in seagrasses and higher plants, or as oxidation products of ω-hydroxy fatty acids and monocarboxylic acids (Cranwell, 1977; Wakeham, 1999; Stefanova and Disnar, 2000). In higher plants, α,ω-dicarboxylic acids typically occur as constituents of cuticular waxes, or as components in cutin and suberin (Cranwell, 1977; Wakeham, 1999). Formation of α,ω-dicarboxylic acids, via oxidative processes, will result in profiles similar to their precursor acids (Cranwell, 1977; Wakeham, 1999).

## 5.2.6 Ester- and Amide-Bound Lipids in Bacteria

Bacterial fatty acids typically occur in bound form as phospholipids, glycolipids, lipoproteins, lipopolysaccharides, and lipoteichoic acids (O'Leary, 1962; Zelles, 1999). The predominant components of bacteria, useful for characterizing microbial communities in sediments, are phospholipid fatty acids and lipopolysaccharides (Zelles, 1999; Rütters et al., 2002). The ester- and amide-bound lipids can be utilized as unique markers for bacteria in the environment. The phospholipid fatty acids are predominantly located in the inner cellular membrane but also occur in the outer cellular membrane (Fig. 5.3). The illustration in Fig. 5.4 represents the components of phospholipids in bacterial cellular membranes, where the head of the phospholipid is hydrophilic (polar end) and the tail is hydrophobic (non-polar end). Lipopolysaccharides make-up a significant portion of the outer cell-membrane of Gram-negative bacteria, where the most abundant fatty acids are the  $\beta$ -hydroxy fatty acids (Kawamura and Ishiwatari, 1984; Zelles, 1999). The  $\beta$ -hydroxy fatty acids are unique markers exclusive to a bacterial origin (Kawamura and Ishiwatari, 1984; Wakeham, 1999; Garette-Lepecg et al., 2004). In the outer cell-membrane, n-carboxylic acids and  $\beta$ -hydroxy fatty acids occur as substituted constituents (via ester- and amidelinkages, respectively) on the phosphate-sugar backbone of Lipid-A of lipopolysaccharides (Fig. 5.5; Bhat and Carlson, 1992).



**Fig. 5.3.** Cellular structure of a Gram-negative bacterium (from http://www.bmb.leeds.ac.uk/mbiology/ug/ugteach/icu8/introduction/bacteria.html# cell\_wall).







**Fig. 5.5.** Structure of Lipid-A in Gram-negative bacteria (from http://www.cyberlipid.org/glycolip/glyl0005.htm)

# 5.2.7 Carbon Isotopic Composition of Fatty Acids

Sources of fatty acids (e.g., terrigenous, marine plankton, or bacteria) can

be elucidated using their compound-specific carbon isotope compositions (e.g.,

Monson and Hayes, 1982; Abrajano et al., 1994; Naraoka et al., 1995; Duan et

al., 1997; Naraoka and Ishiwatari, 1999, 2000; Tolosa et al., 1999). Terrigenous

plants utilize different photosynthetic pathways for carbon fixation (*i.e.*, C3, C4, or

CAM pathways; see Chapter 3, Section 3.1.3). C3 plants are depleted in <sup>13</sup>C, C4 plants are more enriched in <sup>13</sup>C, and CAM plants (which can utilize both C3 and C4 pathways) have intermediate  $\delta^{13}$ C values (Collister *et al.*, 1994). The  $\delta^{13}$ C values of C3 plants range around  $-26^{\circ}/_{00}$ , C4 plants around  $-13^{\circ}/_{00}$ , and  $\delta^{13}$ C values for CAM plants cover the range typical of C3 and C4 plants (Deines, 1980). Lipids, however, are commonly depleted in <sup>13</sup>C relative to the biomass by  $3^{\circ}/_{oo}$  to  $12^{\circ}/_{oo}$  (Deines, 1980; Monson and Hayes, 1982; Collister *et al.*, 1994; Abraham et al., 1998; Naraoka and Ishiwatari, 1999). In a study by Naraoka and Ishiwatari (2000), fatty acids ( $C_{20:0}$  to  $C_{30:0}$ ) from leaves of C3 terrigenous plants ranged between  $-35^{\circ}/_{00}$  to  $-30^{\circ}/_{00}$ , whereas marine derived fatty acids (e.g., C<sub>14:0</sub>,  $C_{16:0}$ , and  $C_{18:0}$ ) had an average  $\delta^{13}C$  value of -23.8±1.1<sup>o</sup>/<sub>00</sub>. Duan *et al.* (1997) measured the isotopic composition of fatty acids (C<sub>16:0</sub> to C<sub>28:0</sub>) in sediment samples from Ruoergai marsh, China. The average  $\delta^{13}$ C composition for individual fatty acids was  $-33.7^{\circ}/_{\circ\circ}$ , which was about  $7.3^{\circ}/_{\circ\circ}$  lighter than the average  $\delta^{13}$ C composition of C3 plants (*i.e.*, -26.4°/<sub>00</sub>) around the marsh. The  $7.3^{\circ}/_{\circ\circ}$  difference is consistent with levels of depletion observed between lipids and their associated biomass. Knowledge of end-member isotopic compositions of fatty acids would allow the proportion of source contributors to be estimated in marine sediments (Naraoka and Ishiwatari, 2000).

More recently, carbon pathways within microbial communities have been investigated using compound-specific carbon isotope analysis of fatty acids derived from cellular membranes of bacteria (Boschker *et al.*, 1998, 1999; Abraham *et al.*, 1998; Boschker and Middleburg, 2002; Petsch *et al.*, 2003;

Londry and Des Marais, 2003; Londry *et al.*, 2004). Carbon isotopic fractionation of bacterial lipids is affected by the mode of growth (*i.e.*, heterotrophic or autotrophic growth in the environment), growth substrate (*e.g.*, acetate, mannose, lactose, or glycerol), and metabolic pathway (*e.g.*, the acetyl-CoA pathway or tricarboxylic acid cycle) (Abraham *et al.*, 1998; Boschker *et al.*, 1998; Londry and Des Marais, 2003; Londry *et al.*, 2004). However, it has been demonstrated that the isotopic composition of fatty acids is not dependent on the growth stage of the bacteria (Summons *et al.*, 1994; Abraham *et al.*, 1998; and Londry and Des Marais, 2003).

Sulfate-reducing bacteria, for example, are ubiquitous in coastal sediments and can mineralize as much as 50% of the total organic carbon (Jørgensen, 1982; Boschker *et al.*, 1998). The primary substrate used by sulfate-reducing bacteria, during the anaerobic decomposition of organic matter, is acetate (Parkes *et al.*, 1989; Boschker *et al.*, 1998). In a study by Boschker *et al.* (1998), the isotopic composition of fatty acids was measured for sulfate-reducing bacteria, using <sup>13</sup>C-labelled acetate as the substrate. The isotopic composition of key phospholipid fatty acids associated with the Gram-positive bacteria *Desulfotomaculum acetoxidans*, indicated that these bacteria consumed a significant portion of the <sup>13</sup>C-labelled acetate. <sup>13</sup>C-Labelled acetate, however, was not significantly utilized by the Gram-negative bacteria of the *Desulfobacter* species (Boschker *et al.*, 1998).

Methanotrophs (*i.e.,* methane-oxidizing bacteria) in anaerobic sediments utilize methane as their carbon and energy source. They play an important role in

limiting the amount of methane released from anaerobic environments (Boschker *et al.*, 1998). Biogenic methane has isotopic compositions depleted in <sup>13</sup>C in the range of  $-110^{\circ}/_{\circ\circ}$  to  $-60^{\circ}/_{\circ\circ}$  (Hunt, 1996). Bacteria utilizing biogenic methane will reflect biomass depleted in <sup>13</sup>C, by as much as  $20^{\circ}/_{\circ\circ}$  (Boschker *et al.*, 1998; Boschker and Middelburg, 2002).

# 5.3 Results and Discussion

In the following sections, compound classes identified in the ester- and amide-bound lipid fractions from core MBH 54/2 will be discussed. Ester- and amide-bound lipids were isolated by the saponification and acid hydrolysis of solvent extracted sediments. Techniques used to release and extract ester- and amide-bound lipids are discussed in more detail in Chapter 2. Lipids were identified based on mass spectra, retention time, and comparison to literature data (see **Appendix III** for representative spectra from identified lipid classes). Ester-bound lipids in sediments from core MBH 54/2 include carboxylic acids,  $\alpha$ -hydroxy fatty acids,  $\beta$ -hydroxy fatty acids,  $\omega$ -hydroxy fatty acids, and n-alcohols, and are summarized in **Table 5.1**. The amide-bound lipid fraction consists of carboxylic acids and  $\beta$ -hydroxy fatty acids. The quantitative data for the lipid constituents from ester- and amide-bound lipid fractions are summarized in **Appendix V**.

Ś						
(m Dep	t,	Carboxylic acids	β-FAOH	α-FAOH	ω-FAOH	n-alcohols
0	5.	12,i13,ai13,13,i14,14,i15, ai15,15,i16,16,i17,ai17,17, 18,19,20-30	10,i12,12,13,ai13,14,14 115,ai15,15,i16,16,i17,ai17,17 18,19,20,21,22,23	16,17,18,20,22,24	16,22,24	16,18,20, 22,24,26
0	2.0	12,i13,ai13,13,i14,14,i15, ai15,15,i16,16:1,16,i17,ai17, 17,18:1ª,18:1 <sup>6</sup> ,18,19,20-30	i13,i14,14,i15,ai15,15,i16,16, i17,ai17,17,18	16.18.20.22	I	ł
	1.2	12,i13,ai13,13,i14,14,i15, ai15,15,i16,16:1,16,i17,ai17, 17,18:1ª,18:1 <sup>6</sup> ,18,19,20-30	10,i12,12,i15,ai15,15,i16,16, i17,ai17,17,18	16,18,20,22,23,24	16	16,18,20, 22,24,26
	1.4	12,i13,ai13,13,i14,14,i15, ai15,15,i16,16, <b>10Me16:0</b> , i17, ai17, <b>cy17</b> ,18:1 <sup>ª</sup> ,18,19,20-30	10,12,i13,ai13,13,i14,14,i15, ai15,15,i16,16,i17,ai17,17	16	1	ł
	1.6	12,i13,ai13,13,i14,14,i15,ai15, 15,i16,16:1,16, <b>10Me16:0</b> ,i17, ai17, <b>cy17</b> ,17,18:1 <sup>a</sup> ,18,19,20-32	10,11,i12,12,i13,ai13,13,i14, 14,i15,ai15,15,16,16,16,i17,ai17, 17,18	16,18,20,22,23,24, 24:1,25,26	16,22,24,26	16,18,20, 22,24,26
	2.3	i13,ai13,13,i14,14,i15,ai15, 15,i16,i17,ai17,17,18,19,20-30	12,i13,ai13,13,i14,14,i15,ai15, 15,i16,16,i17,ai17,17,18,20	16,17,18,20	16,22	16,18,24
	3.0	12,i13,ai13,13,i14,14,i15, ai15,15,i16,16,i17,ai17,17, 18,19,20-30	10,12,i13,ai13,13,114,14, i15,ai15,15,i16,16,i17,ai17, 17,18,20	16,17,18	I	16,18,22,24
	3.3	i14,14,15,ai15,15,i16,16, i17,ai17,17,18,19,20-32	12,i13,ai13,13,i14,14,i15,ai15, 15,i16,16,i17,ai17,17,18,19, 20,21,22	16,18,20,22,24,26	16,18,20, 22,24,26	16,18,20, 22,24,26
	3.5	12,i13,ai13,13,i14,14,i15, ai15,15,i16,16,16,16:1 <b>,10Me16:0</b> , i17,ai17,17,18,19,20-30	10,11,12,i13,ai13,13,i14,14, i15,ai15,15,i16,16,17,ai17,17, 18,19,20,21,22	16,18,20	16,22	16,18,20, 22,24,26
	3.9	i14,14,15,ai15,15,i16,16, i17,ai17,17,18,19,20-32	12,i13,ai13,13,i14,14,i15,ai15, 15,i16,16,i17,ai17,17,18,19, 20,21,22	16,18,22	16,18,20, 22,26	16,18,20, 22,24,26

ester bound fraction in core MBH 5//2 ocition in the mon binit form ( 2 Tahle 5.1 Sum

Roman numerals indicate carbon chain-length.

### 5.3.1 n-Carboxylic Acids in the Ester-Bound Lipid Fraction

Carboxylic acids comprised of straight chain, branched, and monounsaturated fatty acids dominated the ester-bound lipid fraction in core MBH 54/2. Throughout the core, a bimodal distribution is observed for carboxylic acids (**Fig. 5.6**). Short-chain fatty acids ( $C_{12:0}$  to  $C_{20:0}$ ) are significantly more abundant than long-chain fatty acids ( $C_{21:0}$  to  $C_{30:0}$ ). On average, the short-chain fatty acids comprise ~78% of the ester-bound alkanoic acids and the long-chain fatty acids the remaining ~22%. The ester-bound short-chain fatty acids ( $C_{12:0}$  to  $C_{20:0}$ ) are most abundant between 0.7m and 2.3m, with amounts ranging between 81% and 89%. In general, short-chain alkanoic acids ( $<C_{20:0}$ ) are associated with bacterial and planktonic input, whereas long-chain alkanoic acids ( $>C_{20:0}$ ) are derived from cuticular waxes of terrigenous plants.

The concentrations of short- and long-chain fatty acids are plotted in **Fig. 5.7**. The long-chain alkanoic acids do not demonstrate much variation down the core (concentration ranges between 2-4  $\mu$ g/g dry sediment weight in the upper 3.5m). A slight enrichment, to about 6  $\mu$ g/g dry sediment weight, is observed at 1.2m and 1.6m; below 3.5m the concentration of long-chain alkanoic acids falls below 1  $\mu$ g/g dry sediment weight.

The degree of variation in fatty acid composition and shifts in source contributions preserved in core MBH 54/2 can be illustrated by plotting the aquatic-to-terrigenous ratio (*i.e.*,  $\Sigma(C_{12:0}-C_{18:0})/\Sigma(C_{22:0}-C_{28:0})$ ; **Fig. 5.8**). The highest influx of aquatic derived organic matter is observed between 0.7m and 1.4m. A higher influx of terrigenously derived organic matter is observed







**Fig. 5.7.** Downcore distribution illustrating the abundance of short- and longchain n-alkanoic acids in the ester-bound lipid fraction.



**Fig. 5.8.** Ratio of short to long chain ester-bound fatty acids as methyl esters. Downcore variations and shifts in contributions of organic source material are illustrated in this diagram.

at 1.6m, followed by increased contributions from aquatically derived organic matter around 2.3m. Between 2.3m and 3.3m, more abundant contributions from terrigenously derived organic matter are recorded in the sediment. A slight increase in aquatically derived organic matter is observed at 3.5m and 3.9m. Results of the C/N ratio (discussed in Chapter 3) support these observations where a higher C/N ratio at 1.6m shifts towards lower C/N ratios around 2.3m, followed by more intermediate C/N ratios between 2.3m and 3.3m (indicating a mixture of both terrigenous and aquatic derived organic matter).

The carbon preference index (CPI) was originally used to estimate the thermal maturity of source rocks and crude oils, based on the odd-over-even preference of n-alkanes (Bray and Evans, 1961). n-Alkanes with CPI values greater than or less than 1.0 suggest low thermal maturity, whereas CPI values closer to 1.0 indicate higher thermal maturity. In Recent sediments, which are immature, the CPI has been applied to n-alkanoic acids (even-over-odd preference) as an indicator for the degree of diagenetic alterations (Matsuda and Koyama, 1977; Meyers and Ishiwatari, 1993). CPI values of n-alkanoic acids that approach 1.0 suggest higher degrees of diagenetic alterations. In ester-bound lipid fractions from core MBH 54/2, an even-over-odd predominance pattern is observed over the total carbon number range ( $C_{12:0}$  to  $C_{30:0}$ ; **Fig. 5.9**), with an average CPI ~ 6.5 (**eq. 5.1**). The average CPI for short-chain fatty acids ( $C_{12:0}$  to  $C_{20:0}$ ) is 6.8 (**eq. 5.2**); long-chain fatty acids ( $C_{20:0}$  and  $C_{30:0}$ ) have an average CPI



**Fig. 5.9.** Carbon preference index for ester-bound fatty acids as methyl esters, relative to depth, in core MBH 54/2.

$$CPI_{C_{12:0}-C_{30:0}} = \frac{\left(C_{12:0} + 2*\left(C_{14:0} + C_{16:0} + \dots + C_{26:0} + C_{28:0}\right) + C_{30:0}\right)}{2*\left(C_{13:0} + C_{15:0} + \dots + C_{27:0} + C_{29:0}\right)} \quad (eq. 5.1)$$

$$CPI_{C_{12:0}-C_{20:0}} = \frac{\left(C_{12:0} + 2*\left(C_{14:0} + C_{16:0} + C_{18:0}\right) + C_{20:0}\right)}{2*\left(C_{13:0} + C_{15:0} + C_{17:0} + C_{19:0}\right)} \quad (eq. 5.2)$$

$$CPI_{C_{20:0}-C_{30:0}} = \frac{\left(C_{20:0} + 2*\left(C_{22:0} + C_{24:0} + C_{26:0} + C_{28:0}\right) + C_{30:0}\right)}{2*\left(C_{21:0} + C_{23:0} + C_{25:0} + C_{27:0} + C_{29:0}\right)} \quad (eq. 5.3)$$

of 5.5 (**eq. 5.3**). In the short chain fatty acid fraction, the highest CPI values are observed between 0.7m and 1.4m, where CPI values range between 8.3 and 9.6. Small spikes in the CPI values for short-chain fatty acids are observed at depths of 2.3m and 3.3m, where CPI values are 6.9 and 6.1, respectively. The long-chain fatty acids appear to demonstrate more variability in CPI values down the core, with CPI values shifting between 4.1 and 6.6. Peak CPI values occur at 0.7m, 1.6m, and 3.3m (where CPI values are 6.6, 6.2, and 6.5, respectively). The downcore plot of CPI values calculated over the total carbon number range closely parallel CPI values calculated for the short-chain fatty acids.

At depth intervals between 0.7m and 1.4m, CPI values for short-chain fatty acids are significantly greater than CPI values for long-chain fatty acids. Cellular fatty acids in bacteria are commonly dominated by short-chain even carbon numbers. The high CPI values of short-chain fatty acids between 0.7m and 1.4m may reflect an increase in bacteria in the region due to the excessive discharge of sewage into the harbour. The long-chain fatty acids, on the other hand, had much lower CPI values between 0.7m and 1.4m. Lower CPI values may reflect bacterial reworking of sedimentary organic matter. Around 2.3m, there is another slight rise in the CPI value for short-chain fatty acids, with a corresponding lower CPI value for long-chain fatty acids. Again, this may reflect a higher flux of bacteria in the sediment and increased reworking of the longer chain fatty acids. CPI values of long-chain fatty acids demonstrate a slight increase at 1.6m and 3.3m, and correspond to fluxes observed in the concentration of  $\omega$ -hydroxy fatty acids and n-alcohols (discussed in **section 5.3.5**).

#### 5.3.2 Branched-Carboxylic Acids in the Ester-Bound Lipid Fraction

Branched-carboxylic acids in sediments (*i.e.*, primarily i- and ai-C<sub>13:0</sub>, C<sub>15:0</sub>, and C<sub>17:0</sub>) have been used as lipid markers providing evidence for bacterial input (Cranwell, 1973, 1978; Saliot *et al.*, 1991; Guezennec and Fiala-Medioni, 1996; Budge and Parrish, 1998). The proportion of bacteria in a sedimentary environment can be represented by the relative abundance of i-C<sub>15:0</sub> and ai-C<sub>15:0</sub> to C<sub>16:0</sub>, since the C<sub>16:0</sub> fatty acid is ubiquitous in most organisms (Nichols *et al.*, 1987; Mancuso *et al.*, 1990; and Rajendran *et al.*, 1992a). The ratio of i-C<sub>15:0</sub> plus ai-C<sub>15:0</sub> to C<sub>16:0</sub> in the ester-bound lipid fraction in core MBH 54/2 is plotted relative to depth in **Fig. 5.10**. At depths between 2.0m and 4.0m, the (i-C<sub>15:0</sub> + ai-C<sub>15:0</sub>)/C<sub>16:0</sub> ratio ranges from 0.06 to 0.14. In the upper 2.0m, the (i-C<sub>15:0</sub> + ai-C<sub>15:0</sub>)/C<sub>16:0</sub> ratio ranges from 0.23 to 0.68. A significant increase in the proportion of bacterial components is observed in the uppermost section of the core (*i.e.*, peak occurrences at 0.7m, 1.2m, and 1.6m).

Throughout core MBH 54/2, branched fatty acids have been identified in the  $C_{12:0}$  to  $C_{20:0}$  range, comprising between 7.6% to 29.3% of the total short chain fatty acids (**Fig. 5.11**). Branched fatty acids are most abundant in the upper 2.0m of the core (*i.e.*, between 16.8% and 29.3%), with a maximum abundance at 1.6m. Between 2.3m and 3.9m, branched fatty acids become less abundant and consist of 7.6% to 12.7% of the short chain fatty acid fraction. Various groups have attempted to identify and classify bacteria types based on the carbon chain length of branched fatty acids and on the location of branching points (*e.g.*, Edlund *et al.*, 1985; Rajendran *et al.*, 1992c; Vainshtein *et al.*, 1992).


**Fig. 5.10.** Plot illustrating the relative proportion of bacterial components in the ester-bound fraction of sediments from core MBH 54/2, using the ratio of  $(i-C_{15:0} + ai-C_{15:0})/C_{16:0}$  vs depth.



Fig. 5.11. Percent composition of branched chain fatty acids within the  $C_{12:0}$  to  $C_{20:0}$  range of ester-bound lipids.

The ester-bound branched fatty acids identified in sediments from core MBH 54/2 include i-C<sub>13:0</sub>, ai-C<sub>13:0</sub>, i-C<sub>14:0</sub>, i-C<sub>15:0</sub>, ai-C<sub>15:0</sub>, i-C<sub>16:0</sub>, 10Me16:0, i-C<sub>17:0</sub>, ai-C<sub>17:0</sub>, i- $C_{18:0}$ , i- $C_{19:0}$ , and ai- $C_{19:0}$  (as illustrated in **Fig. 5.12**). A broader distribution of branched fatty acids is present in the ester-bound form, compared to free lipids, and has more specificity enabling the identification of bacteria types. Branched fatty acids, with carbon chain lengths ranging between  $C_{14:0}$  and  $C_{16:0}$  are commonly associated with Gram-positive and anaerobic bacteria. Branched fatty acids in the  $C_{16:0}$  to  $C_{19:0}$  range are attributed to sulfate reducing and anaerobic bacteria (Rajendran et al. 1992c). Each of these branched fatty acids was identified throughout core MBH 54/2. Vainshtein et al. (1992) observed that i-C<sub>15:0</sub> fatty acids are predominant in most *Desulfovibrio* species of sulfate reducing bacteria (e.g., Desulfomicrobium and Desulfomomas). i-C<sub>15:0</sub> Fatty acids were observed to be less abundant than ai- $C_{15:0}$  fatty acids in *D. sulfodismutans*, D. alocoholvorans, D. carbinolicus, D. fructosovorans, D. giganteus, and D. gigas (Vainshtein et al., 1992). Throughout the core section from Kowloon Bay, the i- $C_{15:0}$  acid is the most abundant branched fatty acid. The iso-/anteiso-  $C_{15:0}$  ratio (Fig. 5.13) demonstrates that the iso- acid is more abundant than the anteisoacid, with the greatest difference occurring below 3.0m. The Desulfovibrio species of bacteria were likely present throughout the core. In studies of fatty acid profiles of Desulfovibrio species of sulfate reducing bacteria, Edlund et al. (1985) observed i- $C_{15:0}$ ,  $C_{16:0}$ , and i- $C_{17:0}$  saturated fatty acids to be major constituents of D. desulfuricans, D. vulgaris, and D. africanus. These







**Fig. 5.13.** Iso-/Anteiso-ratio of  $C_{15:0}$  and  $C_{17:0}$  fatty acids in the ester-linked fraction of sediments from core MBH 54/2.

bacteria also contain significant amounts of monounsaturated fatty acids (e.g.,  $C_{16:1(n-7)c}$ , i- $C_{17:1(n-7)c}$ , and  $C_{18:1(n-7)c}$ ), and were only observed between 0.7m to 1.6m, and at 3.5m. Major components of *D. gigas* include i- $C_{15:0}$ , ai- $C_{15:0}$ ,  $C_{16:0}$ , and i- $C_{17:0}$  (Edlund *et al.*, 1985). The branched fatty acid, 10Me16:0 (*i.e.*, 10-methyl-hexadecanoic acid), has been identified as a signature compound for *Desulfobacter* species of sulfate reducing bacteria (Rajendran *et al.*, 1992b;

Rajendran *et al.*, 1992c; Vainshtein *et al.*, 1992). The marker for *Desulfobacter* species of sulfate reducing bacteria was only identified at 1.4m, 1.6m, and 3.5m. Each of these depths corresponds to periods with the greatest stanol/sterol ratio (discussed in Chapter 4).

## 5.3.3 Unsaturated-Carboxylic Acids in the Ester-Bound Lipid Fraction

Unsaturated fatty acids were not detected in significant amounts in the ester-bound lipid fraction of core MBH 54/2, and only C<sub>16:1</sub>, C<sub>18:1</sub>, and C<sub>18:2</sub> unsaturated fatty acids were identified in relatively low concentrations. While monounsaturated fatty acids have been used as markers for aerobic bacteria and polyunsaturated fatty acids as markers for eukaryotes (Findlay et al., 1990; Rajendran et al., 1992b,c), certain unsaturated fatty acids have been detected in anaerobic bacteria (Perry et al., 1979; Edlund et al., 1985; Dowling et al., 1986; Rajendran et al., 1992b; Vainshtein et al., 1992). In particular, C<sub>16:1w7</sub> and C<sub>18:1w7</sub> fatty acids have been observed as predominant components in anaerobic bacteria (Perry et al., 1979; Summit et al., 2000), and i- and ai-C<sub>15:1</sub> and C<sub>17:1</sub> fatty acids have been attributed to sulfate reducing bacteria and other anaerobic bacteria (Edlund et al., 1985; Rajendran et al., 1992b; Vainshtein et al., 1992; Summit et al., 2000). No branched unsaturated fatty acids (*i.e.*, *i*- and ai-C<sub>15:1</sub> and  $C_{17:1}$ ) were identified in core MBH 54/2. If any branched unsaturated fatty acids had been present in older sediments of Victoria Harbour, they may have been

hydrogenated to their saturated counterparts (*e.g.*, i- and ai- $C_{15:0}$  and  $C_{17:0}$ ) due to the highly anoxic conditions around the study site.

#### 5.3.4 $\alpha$ - and $\beta$ -Hydroxy Fatty Acids in the Ester-Bound Lipid Fraction

α-Hydroxy fatty acids were less abundant in the ester-bound lipid fraction than β-hydroxy fatty acids. Only straight-chain, even numbered α-hydroxy fatty acids ( $C_{16}$  to  $C_{26}$ ), with variable distributions, were detected downcore (**Table 5.1**). Possible sources of α-hydroxy fatty acids ( $C_{16}$  to  $C_{26}$ ) include phytoplankton, bacteria, and cyanobacteria (Wakeham, 1999; Smallwood and Wolff, 2000).

Substantial amounts of  $\beta$ -hydroxy fatty acids (C<sub>10</sub> to C<sub>20</sub>) were detected and identified in the ester-bound lipid fraction (**Fig. 5.14**). n- $\beta$ -Hydroxy fatty acids comprised ~48% to ~75% of the total  $\beta$ -hydroxy fatty acids; whereas branched  $\beta$ hydroxy fatty acids constituted ~25% to ~52% of the total  $\beta$ -hydroxy fatty acids. The total straight chain and branched  $\beta$ -hydroxy fatty acid contents (in µg/g dry sediment weight) are plotted relative to depth in **Fig. 5.15**. Significant contributions of these compounds to the ester-bound lipid fraction of the sediments are observed between 0.8m and 2.0m, maximizing at 1.2m and 1.6m. A slight secondary flux of  $\beta$ -hydroxy fatty acids is observed at 3.5m. Ester-linked  $\beta$ -hydroxy fatty acids are unique markers for bacteria (Edlund *et al.*, 1985; Mendoza *et al.*, 1987; Skerratt *et al.*, 1992; Wakeham, 1999). They are important cellular components in lipopolysaccharides of Gram-negative bacteria, and are linked by ester- or amide-bonds (Weckesser and Drews, 1979; Edlund *et al.*,





2.3m



**Fig. 5.15.** Downcore profile of straight chain and branched  $\beta$ -hydroxy fatty acids (µg/g dry sediment weight), in the ester-bound lipid fraction of core MBH 54/2.

1985; Mendoza *et al.*, 1987). The occurrence of β-hydroxy fatty acids in sediments is thought to reflect intact cellular remains of bacteria in the sediment (Klok et al., 1988). The depth interval where a large flux in β-hydroxy fatty acids occurs (*i.e.*, 0.8m to 2.0m), corresponds to a period of rapid population growth in regions surrounding the study site. The rapid population growth, in turn, resulted in excess sewage waste discharged into Kowloon Bay. The increase in abundance of bacterially derived β-hydroxy fatty acids suggests that bacterial communities thrived during this period (between ~1933AD and 1974AD). The secondary flux at 3.5m probably indicates an event which led to an abundance of bacterial remnants in sediment. Similar fluxes suggesting the occurrence of bacterial remnants in sediments were observed in **Fig. 5.10** and **Fig. 5.11**, using branched-carboxylic acids as bacterial markers.

## 5.3.5 Ester-bound ω-Hydroxy Fatty Acids and n-Alcohols

Small amounts of even numbered  $\omega$ -hydroxy fatty acids and n-alcohols were detected in the ester-bound lipid fraction of core MBH 54/2 (**Table 5.1**). Variable distributions of even numbered  $\omega$ -hydroxy fatty acids (C<sub>16</sub> to C<sub>26</sub>) were identified in the ester-bound lipid fraction. The presence of  $\omega$ -hydroxy fatty acids reflect contributions from vascular plant material, where C<sub>16</sub> and C<sub>18</sub>  $\omega$ -hydroxy fatty acids are derived from cutin and C<sub>20+</sub>  $\omega$ -hydroxy fatty acids are likely derived from suberin (Cardoso *et al.*, 1977; Kawamura and Ishiwatari, 1984; Wünsche *et al.*, 1987; Fukushima *et al.*, 1992a,b). The downcore distribution of the total  $\omega$ -

hydroxy fatty acid content in **Fig. 5.16** shows that there were at least two periods of high influx of terrigenous plant material (*i.e.*, around 1.6m and 3.3m).

The n-alcohols consisted of variable distributions of even numbered alcohols in the range C<sub>16</sub> to C<sub>26</sub>. Short chain alcohols (*e.g.*, C<sub>12</sub>, C<sub>14</sub>, and C<sub>16</sub>) are typically used as indicators of marine organisms, whereas long chain alcohols  $(C_{20+})$  originate from cuticular waxes of terrigenous plant material (Mudge and Norris, 1997; Mudge and Seguel, 1999). Terrigenous plant material may be the primary source of alcohols detected in the ester-bound lipid fraction. The downcore distribution pattern of the total n-alcohol content (Fig. 5.17) follows the pattern observed for the downcore distribution of the total  $\omega$ -hydroxy fatty acid content. At least two periods of high influx of terrigenous plant material were seen at 1.6m and 3.3m, analogous to the  $\omega$ -hydroxy fatty acids. The two periods with a high influx of terrigenous plant material may indicate periods of strong storms, which transported excess terrigenous plant material into the study site. The first spike occurred around 1.6m, which corresponds to a calendar date around 1954. Around this time period at least 7 typhoons were documented in this area (http://www.hko.gov.hk/informtc/historical tc/no10track.htm).



Fig. 5.16. Downcore distribution of  $\omega$ -hydroxy fatty acids in the ester-bound lipid fraction of core MBH 54/2.



**Fig. 5.17.** Downcore distribution of n-alcohols in the ester-bound lipid fraction of core MBH 54/2.

### 5.3.6 Amide-Bound Carboxylic Acids

Amide-bound lipids consist of n-alkanoic acids and  $\beta$ -hydroxy fatty acids. The n-alkanoic acids are bimodally distributed between C<sub>12:0</sub> and C<sub>30:0</sub> (**Fig. 5.18**), and have an even-over-odd predominance pattern with an average CPI = 6.1. The short chain fatty acids (C<sub>12:0</sub> to C<sub>20:0</sub>) have an average CPI = 7.0, and the long chain fatty acids (C<sub>20:0</sub> to C<sub>30:0</sub>) have an average CPI = 4.9 (**Fig. 5.19**). In the uppermost interval of core MBH 54/2 (0.5m to 0.8m), the CPI over the total carbon number range (C<sub>12:0</sub> to C<sub>30:0</sub>) falls between 5.3 and 6.0. Between 1.0m and 2.0m, the CPI (C<sub>12:0</sub> to C<sub>30:0</sub>) increases to 10.2, decreases to 4.0 to 4.8 (between 3.0m and 3.8m), and increases slightly to 6.7 at 3.9m. The relatively high CPI values indicate that the n-alkanoic acids have not been significantly reworked by bacteria. The slight decrease in CPI values between 3.0m and 3.8m may indicate that slight alterations (*e.g.*, bacterial degradation) may have occurred at these depths. An increase in the presence of bacterial lipids within this depth range supports alterations due to bacteria.

Analogous to the ester-bound lipid fraction, the short chain fatty acids  $(C_{12:0} \text{ to } C_{20:0})$  are more abundant than the long chain fatty acids  $(C_{21:0} \text{ to } C_{30:0})$ . The downcore plot of the aquatic-to-terrigenous ratio (*i.e.*,  $\Sigma(C_{12:0}-C_{18:0})/\Sigma(C_{22:0}-C_{28:0})$ ; **Fig. 5.20**) illustrates the change in contributions of organic source material. A high influx of aquatically derived organic matter is observed between 0.8m and 2.3m, and at 3.9m. The aquatic-to-terrigenous ratio then decreases between 2.6m and 3.3m, possibly due to a slight increase in contributions from terrigenous plant material.







Fig. 5.19. Average CPI of fatty acids in the amide-bound lipid fraction, relative to depth. CPIs were calculated using eq. 5.1, eq. 5.2, eq. 5.3.



**Fig. 5.20.** Aquatic-to-Terrigenous ratio using n-alkanoic acids in the amide-bound lipid fraction.

## 5.3.7 Amide-Linked β-Hydroxy Fatty Acids

β-Hydroxy fatty acids are more abundant in the amide-bound lipid fraction, than in the ester-bound lipid fraction (**Fig. 5.21**). The β-hydroxy fatty acid profiles (C<sub>10</sub> to C<sub>20</sub>) consists of significant amounts of iso- and anteiso-β-hydroxy fatty acids (**Fig. 5.22**). β-Hydroxy fatty acids in the amide-bound lipid fraction are unique to bacteria and are widespread in Gram-negative bacteria (Edlund *et al.*, 1985; Mendoza *et al.*, 1987; Klok *et al.*, 1988; Kaneda, 1991; Bhat and Carlson, 1992; Wakeham, 1999; Garcette-Lepecq *et al.*, 2004).



**Fig. 5.21.** Downcore abundance of ester- and amide-bound  $\beta$ -hydroxy fatty acids in core MBH 54/2.





0.5m

# 5.3.8 Compound-Specific Carbon Isotope Composition of Ester- and Amide-Bound Fatty Acids

The carbon isotopic values for ester-bound fatty acids (C<sub>14:0</sub> to C<sub>30:0</sub>), at depths between 0.8m and 2.6m, are generally more depleted in <sup>13</sup>C than corresponding values at 3.3m and 3.9m (**Fig. 5.23**). The  $\delta^{13}$ C compositions of fatty acids at 3.7m, however, are slightly isotopically lighter than at adjacent depth intervals (*i.e.*, 3.3m and 3.9m). In general, the isotopic composition of C<sub>15:0</sub> and C<sub>18:0+</sub> fatty acids at 3.7m are still enriched in <sup>13</sup>C relative to the average isotopic compositions measured between 0.8m and 2.6m. Downcore distributions of  $\delta^{13}$ C compositions for individual ester-bound fatty acids (*i.e.*, C<sub>14:0</sub> to C<sub>30:0</sub>) are illustrated in **Figures 5.24a-c**. At depths between 0.8m and 3.1m,  $\delta^{13}$ C values ranged between -26°/<sub>oo</sub> and -33°/<sub>oo</sub>. Each of the ester-bound fatty acids (C<sub>14:0</sub> to C<sub>30:0</sub>) demonstrated a general overall enrichment in <sup>13</sup>C at depths below 3.1m.



**Fig. 5.23.** Average carbon isotopic composition of ester-bound fatty acids at depths ranging from 0.8m to 2.6m, and the average isotopic composition at 3.3m and 3.9m.



Fig. 5.24a-c. Downcore carbon isotopic composition of ester-bound fatty acids.

Throughout the core,  $C_{28:0}$  and  $C_{30:0}$  ester-bound fatty acids are more depleted in <sup>13</sup>C, compared to ester-bound fatty acids  $< C_{28:0}$ . The isotopic composition of  $C_{28:0}$  fatty acids range between  $-29.76^{\circ}/_{00}$  and  $-32.49^{\circ}/_{00}$ ; whereas  $C_{30:0}$  fatty acids are isotopically lighter, with  $\delta^{13}$ C values ranging between  $-32.35^{\circ}/_{00}$  and  $-33.19^{\circ}/_{00}$ . The  $C_{28:0}$  and  $C_{30:0}$  fatty acids are likely derived from C3 terrigenous plants. Slight enrichments in <sup>13</sup>C were observed at depths of 3.3m and 3.9m, where  $\delta^{13}$ C values for  $C_{28:0}$  fatty acids are  $-26.92^{\circ}/_{00}$  and  $-24.76^{\circ}/_{00}$ , respectively; and  $\delta^{13}$ C values for  $C_{30:0}$  fatty acids are  $-29.04^{\circ}/_{00}$  and  $-28.74^{\circ}/_{00}$ , respectively. The slight enrichment in <sup>13</sup>C observed at 3.3m and 3.9m may be due to mixing with C4 terrigenous plants.

Branched fatty acids (*e.g.*, i- and ai-C<sub>15:0</sub> and C<sub>17:0</sub>) are typically used as markers for bacteria. In the ester-bound lipid fraction, i- and ai-C<sub>15:0</sub> have  $\delta^{13}$ C values ranging between -28.11°/<sub>00</sub> to -30.59°/<sub>00</sub> (in the upper 2.6m), and are isotopically heavier at 3.3m and 3.9m with an average isotopic composition of -23.75°/<sub>00</sub> (**Fig. 5.25**). Similarly, i- and ai-C<sub>17:0</sub> are isotopically lighter at shallower depths (*i.e.*, upper 1.6m, where  $\delta^{13}$ C ranges between -27.08°/<sub>00</sub> and -28.02°/<sub>00</sub>) and isotopically heavier deeper in the core (*i.e.*, at 3.9m,  $\delta^{13}$ C = -24.66°/<sub>00</sub>; **Fig. 5.25**). In general, however, each of the ester-bound fatty acids appear to be isotopically lighter in the upper half of the core, then systematically become enriched in <sup>13</sup>C with increasing depth.



**Fig. 5.25.** Carbon isotopic composition of i- and ai- $C_{15:0}$  and  $C_{17:0}$  ester-bound fatty acids in core MBH 54/2.

The isotopic compositions of amide-bound fatty acids demonstrate more variability downcore (**Fig. 5.26a-c**).  $C_{14:0}$  and  $C_{16:0}$  amide-bound fatty acids have  $\delta^{13}$ C values ranging between -26.26°/<sub>00</sub> to -29.34°/<sub>00</sub>, and -25.50°/<sub>00</sub> to -29.12°/<sub>00</sub>, respectively; where  $C_{14:0}$  fatty acids are generally slightly lighter than  $C_{16:0}$  fatty acids. Amide-bound fatty acids ( $C_{18:0}$  to  $C_{30:0}$ ) exhibit more shifts in isotopic compositions downcore, where  $\delta^{13}$ C values at 0.5m are more enriched in <sup>13</sup>C compared to all other depths (**Fig. 5.27**). The isotopic composition of  $C_{16:0}$  to  $C_{28:0}$ 



Fig. 5.26a-c. Carbon isotopic composition of amide-bound fatty acids in core MBH 54/2.



**Fig. 5.27.** Isotopic composition of amide-bound fatty acids, at increasing depth intervals in core MBH 54/2.

fatty acids are more depleted in <sup>13</sup>C at 1.4m, than at other depth intervals.  $C_{18:0}$ ,  $C_{24:0}$ , and  $C_{26:0}$  fatty acids demonstrate enrichments in <sup>13</sup>C at 2.3m, 2.6m, 3.3m, and 3.9m (**Fig. 5.26a-c**). At depths of 1.4m and 3.7m,  $\delta^{13}$ C values are slightly lighter than isotopic compositions measured at adjacent depth intervals.

Limited information is currently available on the carbon isotopic composition of fatty acids in bacteria. Various groups (*e.g.,* Abraham *et al.*, 1998; Boschker *et al.*, 1999; Londry *et al.*, 2001; Boschker and Middelburg, 2002;

Londry and Des Marais, 2003) have proposed that the isotopic composition of bacterial fatty acids can be used to evaluate the substrate utilized or to determine if the bacteria were autotrophic or heterotrophic. In a study by Londry et al. (2004) the isotopic composition of fatty acids were measured for four types of sulfate reducing bacteria, grown autotrophically, heterotrophically, or mixotrophically, on substrates of known isotopic composition. No consistent fractionation pattern was observed between the isotopic composition of fatty acids relative to the biomass or substrate, where differences were dependent on growth mode or bacteria type. For example, Londry et al. (2004) reported that Desulfovibrio desulfuricans grown heterotrophically with lactate  $(-29.1^{\circ}/_{00})$  resulted in fatty acids  $(-41.0^{\circ}/_{00})$  that were  $11.7^{\circ}/_{\circ\circ}$  lighter than their biomass (-29.3°/ $_{\circ\circ}$ ) and  $11.9^{\circ}/_{\circ\circ}$  lighter than the substrate. When grown mixotrophically with acetate  $(-34.2^{\circ})_{\infty}$ , fatty acids  $(-36.2^{\circ})_{\circ\circ}$  were  $4.1^{\circ}$  lighter than biomass  $(-32.1^{\circ})_{\circ\circ}$  and  $2.0^{\circ}$  lighter than substrate. Desulfobacter hydrogenophilus grown autotrophically produced fatty acids  $(-52.2^{\circ}/_{00})$  that were  $11.8^{\circ}/_{00}$  lighter than biomass  $(-40.4^{\circ}/_{00})$  and  $24.4^{\circ}/_{00}$ lighter than substrate. Desulfobacter hydrogenophilus was also grown heterotrophically with acetate  $(-34.2^{\circ}/_{00})$ , producing fatty acids  $(-48.1^{\circ}/_{00})$  that were  $13.3^{\circ}/_{\circ\circ}$  lighter than biomass (-34.9 $^{\circ}/_{\circ\circ}$ ) and  $13.9^{\circ}/_{\circ\circ}$  lighter than substrate. Abraham et al. (1998) also compared the isotopic compositions of fatty acids in a set of bacteria grown with different substrates (*e.g.*, glycerol, glucose, mannose, lactose, and a complex medium). They found that the isotopic composition of fatty acids varied between bacterial strains based on the substrates being utilized. It is uncertain whether compound specific carbon isotopes can be used to

differentiate bacterial strains, substrates being utilized, or growth modes being followed, by measuring the isotopic composition of lipids in marine sediments. Under natural environmental settings, the situation is much more complex where multiple bacterial strains and substrates are present, and where each of the bacteria follows a different growth mode and competes for available substrate. This will require further investigation.

There have been reports of the isotopic composition of bacterial fatty acids measured around methane seeps (Hinrichs et al., 2000; Orphan et al., 2001; Zhang et al., 2002; Pancost and Sinninghe Damsté, 2003). The isotopic compositions of lipids from bacteria, involved in anaerobic methane oxidation, were measured in these studies. In general, the  $\delta^{13}$ C composition of fatty acids was depleted in <sup>13</sup>C, and had isotopic values within the range of the methane seeps. Pancost and Sinninghe Damsté (2003) used i- and ai-C<sub>15:0</sub> and C<sub>17:0</sub> fatty acids as markers for sulfate reducing bacteria. The isotopic composition of these branched fatty acids ranged between -60 and -90°/<sub>oo</sub>, demonstrating that the sulfate reducing bacteria were involved in anaerobic methane oxidation. Fatty acids in sediments from core MBH 54/2, in Kowloon Bay, did not have isotopic compositions that were significantly depleted in <sup>13</sup>C. Isotopic values of bacterial fatty acids ranged between -22 and -34°/<sub>oo</sub>, possibly suggesting that biogenic methane was not a significant carbon source or that the bacteria did not play a significant role in anaerobic methane oxidation.

The compound-specific carbon isotope measurement of ester- and amidebound fatty acids will require more in-depth evaluation for potential future

applications. There are currently many uncertainties with potential fractionation effects that may occur with the cleaving of ester- and amide-linkages in bound fatty acids. Ester- and amide-bound fatty acids are likely derived from cellular membranes of a broad variety of bacteria, which follow different metabolic pathways, and metabolize carbon sources from different substrates. Each of these variables further complicates the interpretation of the compound-specific carbon isotope composition of bound fatty acids. Further work should be conducted to determine if fractionation occurs during the isolation of bound lipids, and to better understand possible fractionation effects due to the type of bacteria, metabolic pathways, and type of substrate utilized.

#### 5.4 Summary Remarks

Ester- and amide-bound lipids in sediment core MBH 54/2 from Kowloon Bay are well preserved and provide a record of the sources of organic matter. Ester-bound lipids were dominated by n-alkanoic acids and smaller amounts of hydroxy fatty acids and n-alcohols. The n-alkanoic acids had bimodal distributions where the first modal ranged between  $C_{12:0}$  and  $C_{20:0}$ , maximizing at  $C_{16:0}$ ; the second modal was distributed between  $C_{20:0}$  and  $C_{30:0}$ , maximizing at either  $C_{22:0}$ ,  $C_{24:0}$ , or  $C_{26:0}$ . Relatively high CPI values (average CPI = 6.5) suggest that the n-alkanoic acids are well preserved and have not undergone significant diagenetic alteration. While short chain fatty acids were typically more abundant than long-chain fatty acids throughout the core, this was more evident

at depths between 0.7m and 1.4m. This depth interval corresponds to a period of rapid population growth in Hong Kong, which led to the substantial discharge of sewage waste into Kowloon Bay.

Branched fatty acids were very abundant, constituting between 7.6% to 29.3% of the total short-chain fatty acid fraction. Branched fatty acids in the carbon range  $C_{14:0}$  to  $C_{16:0}$  may have been derived from Gram-positive bacteria or some type of anaerobic bacteria, while branched fatty acids in the  $C_{16:0}$  to  $C_{19:0}$  range may have originated from either sulfate-reducing bacteria or some type of anaerobic bacteria. Iso- and anteiso- $C_{15:0}$  fatty acids are abundant in various species of *Desulfovibrio* type sulfate reducing bacteria. *Desulfobacter* species of sulfate reducing bacteria were identified at depths of 1.4m, 1.6m, and 3.5m, using the signature lipid marker 10Me16:0 fatty acid. Monounsaturated and polyunsaturated fatty acids were not detected in any significant amounts. Only  $C_{16:1}$ ,  $C_{18:1}$ , and  $C_{18:2}$  were identified in the ester-bound lipid fraction.

β-Hydroxy fatty acids (C<sub>10</sub> and C<sub>20</sub>) were identified throughout the core, with significant amounts of iso- and anteiso-β-hydroxy fatty acids. The β-hydroxy fatty acids are derived from bacterial sources and demonstrate two periods of higher influx (*i.e.*, between 0.8m and 2.0m, and around 3.5m). The substantial influx of bacterial material (0.8m to 2.0m) occurs during the period of rapid population growth and excess discharge of sewage waste.

 $\omega$ -Hydroxy fatty acids and n-alcohols were detected in minor amounts and are likely derived from cuticle waxes of terrigenous plant material. At least two intervals in the downcore profile indicate significant contributions from

terrigenous plant material. The two fluxes may be indications for the occurrence of strong storms in the area which could transport excess terrigenous plant material into the area.

Amide-bound lipids were dominated by  $\beta$ -hydroxy fatty acids and smaller amounts of n-alkanoic fatty acids. The  $\beta$ -hydroxy fatty acids are derived directly from the outer cellular membrane of bacteria and have carbon number distributions between C<sub>10</sub> and C<sub>20</sub>. n-Alkanoic fatty acids in the amide-bound lipid fraction, like the ester-bound fatty acids, were bimodally distributed between C<sub>12:0</sub> and C<sub>30:0</sub>. The short-chain fatty acids <C<sub>20:0</sub> were likely derived from bacterial sources, while the long-chain fatty acids >C<sub>20:0</sub> originated from terrigenous plant material. More detailed profiles of  $\beta$ -hydroxy fatty acids from various strains of sulfate reducing bacteria and other types of anaerobic bacteria may provide a means to better delineate the types of bacteria in marine sediments.

Carbon isotopic compositions were measured for ester- and amide-bound fatty acids. Fatty acids in the ester-bound lipid fraction were isotopically lighter in the upper 3m of the core ( $\delta^{13}$ C values ranged between about -26°/<sub>oo</sub> to -33°/<sub>oo</sub>) and became enriched in <sup>13</sup>C at 3.3m and 3.9m. Enrichment in <sup>13</sup>C at 3.3m and 3.9m may have been due to the presence of significant amounts of C4-type terrigenous plants. A slight decrease in  $\delta^{13}$ C composition was observed at 1.4m and 3.7m, which corresponds to spikes observed in the C/N ratio (discussed in Chapter 3). This depletion in  $\delta^{13}$ C composition may reflect the influx of C3-type terrigenous plant material thought to have been carried into the study site via strong storms. The isotopic composition of amide-bound fatty acids

demonstrated more varied distributions downcore. However, similar to the esterbound fatty acids, a slight decrease in  $\delta^{13}$ C composition was also observed at 1.4m and 3.7m. It is still uncertain how the isotopic composition might be affected when ester- and amide-bound fatty acids are freed from sediment samples. Since the short-chain fatty acids are thought to be derived from bacterial sources, the interpretation of isotopic compositions becomes more challenging. The fatty acids most likely represent a broad range of bacterial strains, feeding on different substrates, and which follow different metabolic pathways. Further work is necessary to address each of these uncertainties.

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#### **CHAPTER 6**

#### **Conclusions and Recommendations for Future Work**

### 6.1 Concluding Remarks

Core MBH 54/2 consists of two sediment units, a dark greenish gray to black Holocene mud unit and a desiccated crust unit representing sediments from the upper unit of the late Pleistocene. Very little has been done on the detailed organic geochemical characterization of sediment cores around Hong Kong, which could provide a glimpse of changes to environmental conditions, changes in organic matter source contributions, and changes in bacterial communities throughout the late Quaternary.

Recent dredging activities around Hong Kong have resulted in the release of methane from the Recent sediments, initiating interest in the study of potential sources of methane in Victoria Harbour. The initial goal was to study the downcore variations in isotopic composition of methane in a 4m core section (MBH 54/2) from Kowloon Bay, in Victoria Harbour, Hong Kong. However, methane was not detected in the core sediments. Hence, a lipid marker approach was undertaken to determine the sources of organic matter, evaluate changes in organic matter composition and environmental conditions during the Quaternary, and to ascertain whether remnants of bacterial communities might be present to enlighten our understanding of processes that may have contributed to methane generation. Organic geochemistry tools such as bulk properties (*e.g.*, %C<sub>org</sub>, %N,  $\delta^{13}C_{org}$ , and  $\delta^{15}N$ ), lipid composition and profiles were applied to delineate and

map changes in organic matter sources deposited in the Kowloon Bay area of Victoria Harbour during the late Quaternary.

#### 6.1.1 Summary of Bulk Parameter Measurements

Bulk properties (*i.e.*,  $%C_{ora}$ , %N,  $\delta^{13}C_{ora}$ , and  $\delta^{15}N$ ) of sedimentary organic matter were measured in core MBH 54/2. Fluctuations in the sources of organic matter derived from terrigenous and aquatic sources can be assessed using the C/N ratio. Although the Holocene unit in core MBH 54/2 appears as a thick unit of dark greenish gray to black mud, the C/N ratio demonstrates significant changes in sedimentary organic matter sources at different periods in Hong Kong's history. Higher C/N ratios are observed between 0.7m and 1.6m (C/N ratios generally fall between 13.7 and 14.7). Sediments deposited at this depth interval correspond to calendar dates between ~1954AD and ~1977AD, a period after significant portions of Kowloon Bay had been reclaimed and when untreated sewage was discharged into the study site. Below this depth range (*i.e.*, 1.8m to 2.3m), Kowloon Bay was an open bay and the area did not receive significant amounts of raw sewage. A sharp shift towards lower C/N ratios (*i.e.*, between 8.4 and 9.8) is observed at these depths, indicating that there was a higher input of aquatically-derived organic matter. C/N ratios were more variable between 2.6m and 3.5m, with values shifting between 10.5 and 15.7. The frequent fluctuations suggested that these sediments received a mixture of aquatically and terrigenously derived organic matter. Spikes in the C/N ratio were observed at

1.4m, 2.6m, 3.3m, and 3.7m (where C/N ratios were 18.9, 15.7, 14.7, and 17.2, respectively). The sharp spikes in the C/N ratio may reflect periods of strong storms, carrying excess terrigenously derived organic matter into the area.

Bulk stable isotope compositions (*i.e.*,  $\delta^{13}C_{org}$ , and  $\delta^{15}N$ ) between 0.7m and 1.6m ranged between -28.59°/<sub>oo</sub> to -26.30°/<sub>oo</sub> and 2.14°/<sub>oo</sub> to 3.44°/<sub>oo</sub>, respectively. These isotopic values are consistent with isotopic compositions reported in the literature for sewage contaminated sites. A shift towards isotopically heavier  $\delta^{13}C_{org}$  values (-21.87°/<sub>oo</sub>) was observed at 3.7m in the desiccated crust. The enrichment in <sup>13</sup>C may reflect higher contributions from C4type plants (*e.g.*, C4 seagrasses) thought to have once been present around Hong Kong. The base of the core (4.0m to 4.1m) had  $\delta^{13}C_{org}$  values ranging between

 $-33.17^{\circ}/_{\circ\circ}$  and  $-30.18^{\circ}/_{\circ\circ}$ , and a nitrogen isotope composition of  $2.53^{\circ}/_{\circ\circ}$  (at 4.0m). These isotopic compositions suggest contributions from C3-type terrigenous plants, fixating atmospheric nitrogen as their nitrogen source.

### 6.1.2 Summary of Free Lipid Composition and Profiles

Free lipids in sediments from core MBH 54/2 consisted of sterols, nalcohols, fatty acids, and hydrocarbons. The lipids can be used to delineate sources of sedimentary organic matter and to provide a record of past environmental conditions. The relative abundance of stanols-to-sterols suggested that conditions in Kowloon Bay were anoxic. The stanols were significantly more abundant than sterols between 1.1m and 1.6m (~1967AD and ~1954AD, respectively), and again around 3.4m (~4604BC). These two depth intervals represent periods when Victoria Harbour was highly anoxic. Lower stanol-tosterol ratios were observed at 0.5m and between 2.0m and 3.0m, indicating that conditions were less anoxic. At 0.5m, the seawall-type sewage outfall was diverted further out into the channel of Victoria Harbour via a submarine-type sewage outfall. Between 2m and 3m, Kowloon Bay was an open bay and did not receive significant contributions of sewage waste. Conditions appear to have been more favorable for aquatic organisms during these periods. Sterols common to aquatic organisms (e.g., brassicasterol and campesterol) were observed at these depths; however, they were not detected at 1.1m and 1.6m when conditions were more anoxic. Fecal sterols (*e.g.*, coprostanol, epicoprostanol, 24-ethylcoprostanol, and 24-ethylepicoprostanol) were identified at relatively high concentrations compared to cholesterol at 1.1m and 1.6m. These depths corresponded to periods of rapid population growth in Hong Kong, and periods of high sewage disposal into Kowloon Bay.

Free lipids commonly used to distinguish between aquatic and terrigenous sources include n-alcohols, fatty acids, and n-alkanes. Short chain n-alcohols and fatty acids ( $<C_{20}$ ) denote an aquatic source, whereas long chain n-alcohols and fatty acids ( $>C_{20}$ ) indicate a terrigenous source. The n-alcohols were dominated by the longer chain constituents, indicating that they were derived from cuticular waxes of terrigenous plant material. Fatty acids demonstrated bimodal distributions where short chain fatty acids were more prevalent at 1.1m

and 1.6m. More terrigenously derived organic matter appears to be more abundant than aquatically derived organic matter in the bottom half of the core. Branched fatty acids (*i.e.*, iso- and anteiso- $C_{13:0}$ ,  $C_{15:0}$ , and  $C_{17:0}$ ) indicate a bacterial source. The n-alkanes in core MBH 54/2 had pronounced odd-overeven preference patterns at all depths. It is likely that these hydrocarbons originated from a biotic source, probably cuticular waxes of terrigenous plant material.

### 6.1.3 Summary of Ester- and Amide-Bound Lipids

Ester- and amide-bound lipids in Recent sediments have not been widely utilized but are well preserved and can provide a record of sources of organic matter. Ester-bound lipids in sediments from core MBH54/2 were dominated by carboxylic acids and  $\beta$ -hydroxy fatty acids, with smaller amounts of n-alcohols and  $\omega$ -hydroxy fatty acids. While the carboxylic acids had a bimodal distribution, the short chain fatty acids (C<sub>12:0</sub> to C<sub>20:0</sub>) predominated over long chain fatty acids throughout the core. Branched fatty acids made up a significant fraction (7.6% to 29.3%) of the total short chain fatty acids and other types of anaerobic bacteria. At depths of 1.4m, 1.6m, and 3.5m, the signature marker 10Me16:0 fatty acid was identified. The occurrence of 10Me16:0 fatty acids in the sediments indicate the presence of *Desulfobacter* species of sulfate-reducing bacteria.

The  $\beta$ -hydroxy fatty acids (C<sub>10</sub> to C<sub>20</sub>) are unique to bacteria and can be found at all depths of core MBH 54/2. At least two periods of high influx of  $\beta$ hydroxy fatty acids occur between 0.8m and 2.0m, and around 3.5m. The interval between 0.8m and 2.0m was a period when excess raw sewage was discharged into Kowloon Bay, and conditions appear to have been highly anoxic. Around 3.5m, the event that occurred is not known, but environmental conditions around that depth appear to have been more anoxic.

Lipid markers for terrigenous plants were also identified in the ester-bound lipid fraction. These include  $\omega$ -hydroxy fatty acids and n-alcohols, which are common components of cuticular waxes of land plant material. There were at least two periods of high influx of terrigenous plants, denoted by greater contributions of vascular plant material (*i.e.*, indicated by higher abundance of  $\omega$ -hydroxy fatty acids and n-alcohols) at depths around 1.6m and 3.3m. Sharp fluxes in the C/N ratio were also observed around these depths. The higher flux of  $\omega$ -hydroxy fatty acids and n-alcohols may support the idea that spikes in the C/N ratio are indications of the occurrence of strong storms in the area, which can carry in excess terrigenous plant material.

Amide-bound lipids are comprised of  $\beta$ -hydroxy fatty acids and n-alkanoic acids. The  $\beta$ -hydroxy fatty acids are the dominant compound group in the amidebound form and are thought to be derived directly from the outer cellular membrane of bacteria. More detailed  $\beta$ -hydroxy fatty acid profiles of various strains of sulfate-reducing bacteria and other types of bacteria may provide a means to differentiate bacterial communities in marine sediments.

#### 6.2 Recommendations for Future Work

Organic geochemical characterization of sediments from core MBH 54/2 demonstrated that bulk properties (such as the C/N ratio,  $\delta^{13}C_{org}$  and  $\delta^{15}N$  compositions), lipid marker composition, and profiles can be used to reconstruct past contributions of sedimentary organic matter and to infer environmental conditions as the organic matter was being deposited. This study looked at the deposition of sedimentary organic matter in an area of Kowloon Bay that has undergone significant environmental transformations (*e.g.*, reclamation activities, which in turn have increased sedimentation rates), experienced extreme conditions (*e.g.*, excessive raw sewage disposal directly into the study site), organic matter transported into the area from the Pearl River, and strong storms (which are capable of carrying in excess terrigenous plant material).

In order to gain a broader perspective and understanding of organic matter deposition throughout Hong Kong during the late Quaternary, detailed organic geochemical characterization should be carried out on cores throughout Victoria Harbour and in regions surrounding Hong Kong. A sediment core from the western border of Victoria Harbour might show a greater influence of sedimentary organic matter from the Pearl River, whereas sediment cores from the eastern rim may show a greater marine influence. Core sites should be selected in areas throughout Victoria Harbour that have experienced variable sedimentation rates, proximity to raw sewage disposal sites, and areas that have undergone changes in oxic/anoxic conditions. Each of these factors will have an

impact of the amount and type of sedimentary organic matter deposited and accumulated around Hong Kong.

The occurrence and identification of bacterial remnants in Victoria Harbour sediments should be studied further. If bacterial strains native to Victoria Harbour can be isolated and identified, their lipid profiles may allow us to gain better insights into the types of bacterial communities that were active in Victoria Harbour sediments (*i.e.*, to try and link bacterial lipid profiles to lipid profiles preserved in ester- and amide-bound forms in sediment samples). This may help us understand the roles of bacteria in organic matter remineralization and their role in the consumption and/or generation of carbon dioxide or methane.

Compound-specific carbon isotopes can be measured on ester- and amide-bound fatty acids in sediments. However, it is uncertain if isotopic fractionations occur during the cleavage of ester- or amide-linkages. Further work should be conducted to investigate potential fractionation effects on synthesized ester- and amide-bound fatty acids of known isotopic composition. This would provide better confidence in the possible application of compound-specific isotope compositions of bound fatty acids. The ester- and amide-bound fatty acids in sediments are likely derived from cellular membranes of bacteria. Recent groups (*e.g.*, Boschker *et al.*, 1998, 1999; Abraham *et al.*, 1998; Boschker and Middleburg, 2002; Petsch *et al.*, 2003; Londry and Des Marais, 2003; Londry *et al.*, 2004) have begun evaluating the compound-specific carbon isotope analysis of fatty acids in bacteria. This is a complicated task with many variables to consider in interpreting the isotopic composition of bound fatty acids (*e.g.*, type of

bacteria, metabolic pathway and carbon sources being metabolized). Each of these uncertainties will need to be addressed in order to better utilize and incorporate compound-specific isotope measurements of bound lipids as a tool to better understand processes recorded in Recent sediments.

# 6.3 References

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## **APPENDIX I**

Depth (m)	%C <sub>ora</sub>	Avg %C <sub>org</sub> ± stdev	%N Avg %N ± stdev
0.5	0.75		0.07 0.07 0.00
	0.75	$= 0.75 \pm 0.00$	$0.07 \qquad \qquad$
0.7	1.28	<b>~</b>	0.09
0.9	1.51		0.12
	1.90	≻ 1.68 ± 0.20	$0.13 > 0.12 \pm 0.01$
	1.64	J	0.12
1.2	2.74		0.20
1.4	2.83		0.15
1.6	1.62		0.11
1.8	0.88		0.09
2.1	0.75	]	0.08
	0.85	$> 0.83 \pm 0.07$	$0.09 > 0.09 \pm 0.00$
	0.90	J	0.09
2.2	0.59		0.07
2.4	0.62		0.07
2.6	1.03	$\int 0.04 \pm 0.12$	0.06 0.06 + 0.00
	0.85	$\int 0.94 \pm 0.13$	0.06 5 0.00 ± 0.00
2.8	0.77	$10.72 \pm 0.05$	0.06 0.06 + 0.00
	0.70	0.73 ± 0.05	0.05
3.1	0.67	٦	0.06
	0.65		0.06
	0.65	$> 0.63 \pm 0.05$	$0.06 > 0.06 \pm 0.00$
	<del>0.50</del>		0.06
	0.56	J	0.06
3.4	0.89	0.88 + 0.01	0.06 0.06 + 0.00
	0.88	0.00 ± 0.01	0.06
3.5	0.87	0.83 + 0.06	$0.07  0.07 \pm 0.00$
	0.79	0.03 ± 0.00	0.07
3.7	<del>0.87</del>	]	<del>0.05</del> )
	0.89	$-0.86 \pm 0.04$	0.05 > 0.05 + 0.00
	0.81	0.00 ± 0.01	0.04
	0.89	J	0.05 _
3.9	0.48	$> 0.42 \pm 0.09$	$0.04 \rightarrow 0.04 \pm 0.00$
	0.36		0.04
4.0	0.26	$\succ 0.22 \pm 0.05$	$0.03 \succ 0.03 \pm 0.00$
	0.19		0.03
4.1	0.45		0.03
	0.35		0.03
	0.29	$> 0.39 \pm 0.06$	$0.03 \rightarrow 0.03 \pm 0.00$
	0.42		0.03
	0.42		0.03
	0.42	/	0.03 2
4.2	0.27	≻ 0.23 ± 0.06	$0.04 > 0.04 \pm 0.00$
	0.19	J	0.04

**A1.1.** Bulk measurements of sediments from core MBH 54/2: total organic carbon ( $%C_{org}$ ) and total nitrogen (%N).

<sup>a</sup>bad value – atmospheric nitrogen present <sup>b</sup>bad value – inorganic carbon present

Depth (m)	δ <sup>13</sup> C <sub>org</sub>	Avg δ <sup>13</sup> C <sub>org</sub> ± stdev	δ <sup>15</sup> N	Avg δ <sup>15</sup> N ± stdev
0.5	-22.50	$2260 \pm 0.14$		
	-22.70	-22.00 ± 0.14		
0.7	-23.86		6.19	
0.9	-28.93	]	2.66	]
	-28.38	≻ -28.59 ± 0.30	2.51	≻ 2.57 ± 0.08
	-28.46	J	2.55	<u> </u>
1.2	-26.62		3.44	
1.4	-26.30		2.14	
1.6	-27.49		3.13	
1.8	-27.74		4.49	
2.1	-28.20		3.19	
	-27.92	≻ -27.99 ± 0.18	3.20	$> 3.05 \pm 0.26$
	-27.87	J	2.75	J
2.2	-27.12		4.30	
2.4	-26.92		4.27	
2.6	-26.40	$-26.01 \pm 0.54$		
	-25.63	۲	4.43	
2.8	-27.22	$-27.25 \pm 0.05$		
	-27.29	J	3.61	
3.1	-26.84			
	-27.25			
	-27.20	$-27.23 \pm 0.29$	4.0508	
	- <u>20.03</u> 27.56		<del>1.900</del> 4.60	
2 4	-27.30	)	4.00	
3.4	-24.71	-24.74 ± 0.04	4 57	
2 5	-24.70		4.57	
5.5	-27.41	-27.50 ± 0.13	1 33	
3 7	16 24 <sup>9</sup>	٠	4.55	٠
5.7	-21.96		4 51	
	-21.00	$> -21.87 \pm 0.08$	4.81	≻ 4.51 ± 0.30
	-21.86		4.01	
3.9	-26 10	1		·····
0.0	-26.43	$> -26.27 \pm 0.24$	4.26	
4.0	-30.56			
	-29.79	$> -30.18 \pm 0.54$	2.53	
4.1	-31.03	<u>)</u>		
	-31.09	$-31.85 \pm 0.45$		
	-31.02			
	-31.03	(if delete -29.94,then		
	-31.03	avg= -31.04 ±0.03)		
	<del>-29.9</del> 4	J		
4.2	-33.10	$-33.17 \pm 0.10$		
	-33.24	$\int_{-33.17 \pm 0.10}$		

**A1.2.** Bulk measurements of sediments from core MBH 54/2:  $\delta^{13}C_{org}$  (°/<sub>oo</sub>) and  $\delta^{15}N$  (°/<sub>oo</sub>).

<sup>a</sup>bad value – atmospheric nitrogen present <sup>b</sup>bad value – inorganic carbon present

A2.1. Sterol structures, common names, IUPAC names, and chemical formula.



## **APPENDIX III**

**A3.1.** Representative mass spectra for carboxylic acids,  $\beta$ -hydroxy fatty acids,  $\alpha$ -hydroxy fatty acids,  $\omega$ -hydroxy fatty acids, sterols, and stanols.



























 $C_{14:0}$   $\beta$ -hydroxy fatty acid methyl ester-trimethylsilyl ether





 $C_{16:0} \beta$ -hydroxy fatty acid methyl ester-trimethylsilyl ether































## **APPENDIX IV**

Depth (m)	brassicasterol cholesterol	<u>sitosterol</u> cholesterol	<u>campesterol</u> cholesterol	<u>coprostanol</u> cholesterol	<u>epicoprostanol</u> cholesterol	<u>cholestanol</u> cholesterol
0.5	0.58	0.80	0.44		0.21	0.78
1.1		0.83		3.07	1.17	3.13
1.6		0.70		1.49	0.78	1.76
2.3	0.58	1.09	0.44		0.27	1.17
3.4	0.57	0.90	0.53		0.49	1.90

A.4.1. Summary of sterol ratios in free lipids in core MBH 54/2.

A.4.2. Summary of aquatic/terrigenous ratios for free lipids in core MBH 54/2.

Depth (m)	Saponifiable Fatty Acids Σ(12:0-18:0)/Σ(22:0-28:0)	Non-Saponifiable Fatty Acids Σ(14:0-18:0)/Σ(20:0-24:0)	Alcohol C <sub>18</sub> /C <sub>28</sub>	n-alkanes C <sub>19</sub> /C <sub>31</sub>
0.5	1.50	3.71	0.60	0.25
1.1	2.32	7.25	0.01	0.63
1.6	3.28	6.45	0.41	0.69
2.3	1.00	4.41	0.54	0.20
3.4	0.72	5.72	0.38	0.07

Depth (m)	Saponifiable Fatty Acids CPI <sub>(12:0-34:0)</sub>	Non-Saponifiable Fatty Acids CPI <sub>(14:0-26:0)</sub>	n-Alcohols CPI <sub>(C22-C32)</sub>	n-Alkanes CPI <sub>(C19-C35)</sub>
0.5	7.4	7.0	6.49	2.58
1.1	10.4	9.1		1.24
1.6	9.5	15.9		1.22
2.3	7.1	9.6	7.99	2.24
3.4	5.0	7.5	9.70	3.01

Saponifiable fatty acids: 
$$CPI_{(12:0-34:0)} = \frac{[C_{12:0} + 2*(C_{14:0} + C_{16:0} + \dots + C_{30:0} + C_{32:0}) + C_{34:0}]}{[2*(C_{13:0} + C_{15:0} + \dots + C_{31:0} + C_{33:0})]}$$

Non-saponifiable fatty acids:  $CPI_{(14:0-26:0)} = \frac{[C_{14:0} + 2*(C_{16:0} + C_{18:0} + .... + C_{22:0} + C_{24:0}) + C_{26:0}]}{[2*(C_{15:0} + C_{17:0} + .... + C_{23:0} + C_{25:0})]}$ 

**n-Alcohols:**  $CPI_{(C22-C32)} = \frac{[C_{22} + 2*(C_{24} + C_{26} + \dots + C_{28} + C_{30}) + C_{32}]}{[2*(C_{23} + C_{25} + \dots + C_{29} + C_{31})]}$ 

**n-Alkanes:**  $CPI_{(C19-C35)} = \frac{[C_{19} + 2*(C_{21} + C_{23} + .... + C_{31} + C_{33}) + C_{35}]}{[2*(C_{20} + C_{22} + .... + C_{32} + C_{34})]}$ 



A4.4. Chromatograms of free fatty acids in the saponifiable lipid fraction.



A4.5. Chromatograms of free fatty acids in the non-saponifiable lipid fraction.


A4.6. Chromatograms of free alcohols in the non-saponifiable lipid fraction.



**A4.7.** Chromatograms of free n-alkanes in the non-saponifiable lipid fraction.

**APPENDIX V** 

A.5.1. Ester-bound fatty acid methyl esters (µg/g sediment dry weight)

, 5	12:0	i13:0 (	ai13:0 (	13:0	i14:0 (	14:0	i15:0 (	ai15:0 (	15:0 (	i16:0 (	16:1	16:0	10Me16:0	i17:0 (	ai17:0 (	cy17:0	17:0 (	18:1	18:1	18:0	19:0	20:0	21:0	22:0	23:0 (	24:0	25:0 (	26:0	27:0 (	28:0	29:0	30:0	31:0
0.5m	0.168	0.039	0.020	0.087	0.085	1.105	0.291	0.159	0.336	0.155		2.574		0.155	0.085		0.240			1.089	0.099	0.480	0.096	0.487	0.105	0.738	0.082	0.589	0.051	0.191	0.047	0.142	
0.7m	0.047	0.029	0.014	0.045	0.126	1.037	0.559	0.435	0.332	0.277	0.071	4.035		0.181	0.156		0.281	0.340	0.182	2.689	0.085	0.431	0.058	0.534	0.837	0.528	0.061	0.268	0.040	0.212	0.036	0.148	
1.2m	1.376	0.345	0.111	0.300	0.665	3.952	2.304	1.576	1.157	0.895	0.292	14.63		0.449	0.465		0.867	1.262	0.597	8.324	0.233	1.260	0.185	1.780	0.314	1.194	0.142	0.549	0.118	0.578	0.065	0.451	
1.4m	0.127	0.085	0.017	0.067	0.153	0.994	0.697	0.458	0.432	0.285		5.950	0.442	0.185	0.158	0.015	0.313	0.087		2.801	0.096	0.389	0.098	0.579	0.152	0.445	0.065	0.180	0.042	0.299	0.035	0.152	
1.6m	3.285	0.123	0.042	0.135	0.388	2.525	1.819	0.972	1.050	0.631		8.702	0.940	0.403	0.319	0.123	0.759	0.826	0.595	5.490	0.172	0.876	0.172	1.277	0.228	1.170	0.178	0.674	0.133	1.062	0.111	0.967	0.055
2.3m		0.011	0.005	0.029	0.034	0.476	0.128	0.073	0.195	0.083		1.755		0.075	0.046		0.155			0.916	0.064	0.241	0.057	0.261	0.060	0.313	0.036	0.174	0.019	0.059	0.019	0.036	
3.0m	0.051	0.012	0.005	0.046	0.031	0.497	0.110	0.039	0.190	0.051		1.684		0.050	0.028		0.143			0.593	0.071	0.217	0.070	0.304	0.070	0.440	0.042	0.300	0.027	0.092	0.022	0.061	
3.3m					0.007	0.158	0.043	0.014	0.093	0.027		0.936		0.024	0.012		0.088			0.355	0.045	0.153	0.046	0.253	0.050	0.371	0.038	0.411	0.032	0.158	0.035	0.126	0.024
3.5m	0.003	0.001	0.0004	0.004	0.003	0.044	0.017	0.005	0.021	0.007	0.002	0.188	0.005	0.006	0.003		0.016			0.074	0.007	0.021	0.007	0.028	0.006	0.038	0.004	0.026	0.002	0.010	0.002	0.006	
3.9m					0.003	0.060	0.017	0.007	0.042	0.010		0.345		0.010	0.005		0.038			0.164	0.015	0.045	0.015	0.054	0.014	0.068	0.011	0.054	0.010	0.031	0.007	0.023	0.00

3.9m			2 08	1.37	1.54	1 22	1.27	7.26	2.25	4.37	1.85	0.81	9.83	2.63	2.08	1.36	3.41	1.92
3.5m	14 20		24.35	02.6	5.28	8.14	8.69	49.87	16.76	20.53	18.28	5 07	67.44	28.80	15.74	7.49	26.37	12.71
3.3m			4.75	4.32	1.60	3.37	3.94	18.31	6.84	10.13	4.90	2.45	22.93	7.50	9.40	5.50	10.97	5.55
3.0m	7.72		14.79	4.77	2.09	4.25	5.25	35.09	9.29	17.47	16.36	4.60	51.87	13.62	14.14	7.45	14.57	19.60
2.3m			12.31	20.44				35.90	14.17	17.37	6.56		40.31	24.62	13.94	5.09	13.30	6.76
1.6m	81.26	31.74	185.64	68.89	19.85	27.97	24.43	208.08	89.44	37.04	38.35	23.39	170.45	148.81	32.87	51.15	57.15	
1.4m	54.34		78.82	10.80	5.95	32.29	8.29	95.57	32.95	15.95	53.58	7.96	168.51	58.32	27.85	14.68		
1.2m	86.33	26.12	132.49	121.36	16.48	4876	47.72	239.18	128.05	46.75	74.24	36.86	244.67	212.71	54.55	105.12	60.09	
0.8m	40.38		69.07	53.45	26.40	8.59	165.82	72.81	45.18	24.27	143.36		72.80	24.71	42.93	60.59		
0.7m				31.19			5.69	47.92	22.54	14.96	7.98	7.21	44.73	38.05	25.82	13.60	21.68	
0.5m	8.08	1.93	23.05	23.57	4.17	6.93	11.14	50.65	17.37	25.06	9.04	6.10	57.25	23.54	21.36	10.11	21.71	13.45
#	10β	i12B	12β	i13B	ai13ß	13β	i14ß	14β	i15β	ai158	15β	i16β	16β	i17ß	ai17ß	178	18β	20ß

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2. Ester-bound β-hydroxy	
5.2. Ester-bound β-hydroxy	
<b>A.5.2.</b> Ester-bound β-hydroxy	

	3.9m	<u>9.90</u>	4.28	2.36	7.20	3.14			3.9m	17 89	34,19	12.78	17.51	8.65	1.96
	3.5m	37.25			16.29				3.5m	70.68	48.79	33.1	49.29	10.15	1.56
	3.3m	33.36	27.62	9.38	53.75	37.41			3.3m	40.53	30.92	41.0	68.32	39.28	12.63
lt)	2.3m	2.93			3.86				3.0m	38.18	12.31		9.85	6.43	
ent dry weigh	Ę	3.8			3.9	1.4	c.2	ght)	2.3m	24.5	11.0			3.34	
(ng/g sedim	1.6	27:			22	15	721	ment dry wei	1.6m	214.5	170.7	109.1	160.3	110.5	73.7
<u> vy fatty acids</u>	1.2m	31.02						ls (ng/g sedir	1.2m	183.8	110.1	64.3	65.0	19.9	10.0
ound w-hydro	0.5m	14.40			26.23	17.43		und n-alcoho	0.5m	41.0	29.2	11.6	27.0	13.8	2.90
A.5.3. Ester-bo	C#	16w	18w	200	220	24W 264	moz	<b>A.5.4</b> . Ester-bo	C#	16	18	20	22	24	26

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i14:0	29.98			63.52	88 q1		110-0
14:0	135.33	114.76	248.24	274.45	142.27	39.00	38.20
i15:0	56.79	49.05	122.06	147.47	119.80	18.04	14.98
ai15:0	23.12	30.14	86.02	59.55	3.64	5.54	10.29
15:0	64.23	77.37	141.92	204.47	144.34	30.04	38.31
i16:0	41.00	42.29	150.70	95.26	90.06	22.51	6.53
16:0	555.55	684.02	1358.15	2058.00	1729.11	192.65	225.19
i17:0	107.22	69.46	211.34	206.81	161.95	33.98	20.57
ai17:0	58.01	46.07	75.88	108.28	90.96	31.19	4.26
17:0	134.09	73.34	309.15	139.26	158.13	38.99	35.98
18:0	402.07	382.06	780.08	1183.41	1331.86	56.17	111.08
19:0	42.58	25.22	138.47	68.92	60.33	9.97	7.87
20:0	177.62	<u> 06.90</u>	272.45	363.06	295.95	40.01	22.28
21:0	22.21	21.86	59.72	68.22	44.59	29.30	13.07
22:0	268.19	185.54	358.65	782.71	510.31	59.63	49.36
23:0	56.97	44.75	58.86	97.13	70.77	15.85	15.40
24:0	393.15	155.88	358.60	498.08	451.48	74	39.53
25:0	27.08	32.98	94.18	59.20	38.32	87	19.32
26:0	130.88	104.19	113.59	161.61	135.22	12.93	21.11
27:0	16.50				31.86	41.74	
28:0	47.86	36.17	159.01	105.92	102.46		10.89
29:0	17.51				23.34	21.54	
30:0	29.54			76.87	95.01		8.77
31:0	10.72				1 8 8		
32:0					56.81		

A.5.5. Amide-bound fatty acid methyl esters (ng/g sediment dry weight)

3.9m	18.98	20.64	5.14	4.74	31.87	77.57	35.02	63.32	24.55	6.66	75.18	13.84	11.38	4.57	6.70		11.47
3.3m	34.48	13.78	3.73	2.46	29.42	86.66	28.97	69.53	17.84	19.38	56.10	1315	12.33	5.36	24.72		15.48
1.6m	202.31	82.02	24.39	57.50	86.93	608.38	206.21	113.91	66.65	82.88	563.43	420.69	81.89	146.79	164.08	14.47	48.41
1.2m	235.61	81.18	9.36	78.12	59.26	545.37	231.90	110.88	127.32	144.10	543.54	406.71	66.33	105.99	96.44		
0.8m	159.89	67.26	23.52	36.43	88.47	456.14	123.64	151.74	95.97	138.55	383.04	192.25	79.71	57.52	132.73		63.01
0.5m	74.98	58.38	20.39	24.93	49.13	220.97	75.42	168.66	63.92		172.27	52.34	54.54		236.82		
₿	12β	i13B	ai13ß	13β	i14ß	14β	i15β	ai15β	158	i16β	16β	i17B	ai17ß	17B	18ß	19 <b>β</b>	20ß

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Depth (m)	CPI <sub>(12:0-30:0)</sub>	CPI <sub>(12:0-20:0)</sub>	CPI <sub>(20:0-30:0)</sub>	$\frac{\sum (C_{12:0} - C_{18:0})}{\sum (C_{22:0} - C_{28:0})}$	$\frac{(iC_{15:0} + aiC_{15:0})}{C_{16:0}}$
0.5	6.48	6.68	6.09	2.46	0.18
0.7	9.63	10.78	6.56	5.07	0.25
1.2	9.82	11.04	6.02	6.90	0.27
1.4	9.06	11.01	4.54	6.57	0.19
1.6	8.14	8.88	6.21	4.78	0.32
2.3	6.64	7.38	4.93	3.90	0.11
3.0	6.15	6.46	5.54	2.49	0.09
3.3	6.72	6.75	6.67	1.21	0.06
3.5	6.31	6.71	5.43	3.04	0.12
3.9	5.51	6.25	4.26	2.76	0.07

**A.5.7.** Summary of carbon preference indices, aquatic-to-terrigenous ratio, and the  $(i-C_{15:0} + ai-C_{15:0})/C_{16:0}$  ratio for ester-bound fatty acids.

**A.5.8.** Summary of carbon preference indices, and the aquatic-to-terrigenous ratio for amide-bound fatty acids.

Depth (m)	<b>CPI</b> <sub>(12:0-30:0)</sub>	<b>CPI</b> <sub>(12:0-20:0)</sub>	CPI <sub>(20:0-30:0)</sub>	$\frac{\sum (C_{120} - C_{180})}{\sum (C_{220} - C_{280})}$
0.5	6.00	5.91	6.15	1.59
0.8	5.33	5.35	5.29	3.23
1.2	7.86	8.85	6.04	2.93
1.4	8.34	10.16	5.72	2.87
1.6	7.93	8.93	6.18	2.86
2.3	5.88	7.33	3.79	3.10
2.6	5.71	6.54	4.59	2.06
3.1	4.84	5.22	4.39	1.77
3.3	4.51	5.18	3.53	2.39
3.7	4.04	5.57	2.75	1.82
3.9	6.72	7.52	5.05	3.43

**APPENDIX VI** 

A.6.1.	δ <sup>13</sup> C (°,	/ <sub>00</sub> ) corr	positior	of este	sr-bound	d fatty a	icids in	core MI	3H 54/2							
Depth (m)	14:0	i15:0	ai15:0	15:0	i16:0	16:0	i17:0	ai17:0	17:0	18:0	20:0	22:0	24:0	26:0	28:0	30:0
0.8	-27.66	-28.48	-28.83	-27.33	-30.36	-27.54	-27.40	-28.02	-27.15	-26.78		-29.24	-28.44	-28.77	-32.30	
1.2		-28.11	-29.15	-27.97		-27.20	-27.08	-27.11	-26.20	-26.66	-27.80	-28.58	-27.99	-29.26	-29.76	
1.4	-28.98	-29.26	-30.48	-28.08		-29.59				-27.37	-28.61	-29.97	-29.34			-32.68
1.6	-28.14	-28.67	-29.26	-27.92		-28.85	-27.45	-27.11	-26.09	-26.07		-28.06	-27.97	-29.26	-32.49	-32.35
2.3	-28.94	-29.13	-30.59	-28.81		-28.00			-26.18	-27.92	-26.82	-28.43	-28.29	-29.20	-30.08	-32.52
2.6	-29.12	-29.97	-29.01	-28.09		-29.05			-26.22	-26.68	-25.68	-26.95	-28.12	-29.33	-29.88	-33.19
3.1	-30.43			-26.19		-26.79				-26.24	-26.29	-28.06	-30.30	-31.30	-30.97	-32.90
3.3	-26.70	-23.69		-22.94		-25.27			-23.15	-23.91		-24.25	-26.51	-27.38	-26.92	-29.04
3.7	-28.17			-26.33		-28.22				-24.61	-25.52	-24.68	-26.66	-28.00	-28.89	-30.41
3.9	-25.80	-23.80		-22.65		-24.42	-24.66		-24.61	-23.53	-27.48	-24.20	-23.79	-25.62	-24.76	-28.74

Depth (m)	14:0	16:0	18:0	20:0	22:0	24:0	26:0	28:0	30:0
0.5	-27.90	-26.52	-23.85	-25.64	-22.89	-23.10	-22.95	-24.41	-27.24
0.8		-25.50	-26.92		-25.82	-23.41			
1.2		-26.46	-26.90		-28.86	-27.31			
1.4		-29.12	-27.68	-29.98	-30.27	-27.55	-29.60	-34.84	
1.6	-28.56	-26.40	-27.23	-29.64	-28.37	-26.55	-29.53	-32.89	-33.8(
2.3	-28.86	-27.66	-25.69	-27.96	-29.02	-25.16			
2.6	-29.34	-26.85	-24.69	-28.25	-27.52	-25.27	-24.87	-29.34	-32.1
3.1	-29.03	-28.17	-27.35	-27.19	-26.91	-26.18	-27.71		
3.3	-26.26	-25.95	-24.44	-27.46	-26.58	-24.41	-24.99	-27.63	-32.0
3.7	-27.06	-28.30	-26.82	-31.76		-30.35	-31.12		
3.9	-28.83	-27.05	-26.43	-28.41	-28.70	-24.79	-23.73	-29.24	-33.6

d fatty acids in core	
i of amide-boun	
) composition	
0/00)	
. δ <sup>13</sup> C	
6.2	