# A laboratory experiment on the behavior of soil derived core and intact polar GDGTs in aquatic environments

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# 12 Abstract

13 We have performed incubation experiments in order to examine the behavior of soil-14 derived branched glycerol dialkyl glycerol tetraether (brGDGT) membrane lipids 15 upon entering an aquatic environment and to evaluate the processes that potentially 16 take place during their fluvial transport from land to sea. We incubated a soil from the 17 Rakaia River catchment on the South Island of New Zealand using Rakaia River 18 water and ocean water collected near the river mouth as inocula for a period of up to 19 152 days. The concentrations, as well as the relative distribution of brGDGTs derived 20 from intact polar ('living'; IPL) lipids and core ('fossil'; CL) lipids remained 21 unaltered over the course of the experiment. Although the stability of the brGDGTs may be a consequence of the higher than natural soil:water ratio used in the laboratory 22 23 experiment, the substantial increase (27-72%) in the total pool of isoprenoid GDGTs 24 (isoGDGTs) in all incubation setups, including the control using distilled water, 25 indicates that entering an aquatic environment does influence the behavior of soil-26 derived GDGTs. However, the availability of water appears to be more important than 27 its properties. As a consequence of increasing isoGDGT concentrations, a decrease in 28 Branched and Isoprenoid Tetraether (BIT) index values - a proxy for the relative input 29 of fluvially discharged soil material into a marine system - became evident after an 30 incubation period of 30 days, with a maximum final decrease of 0.88 to 0.74 in the

1 experiment with river water. The relative distribution within the isoGDGT pool shows 2 changes with time, suggesting that isoGDGT producers may either have different 3 rates of membrane adaptation or production/degradation, or that preferential release 4 from the soil matrix or a shift in source organism(s) may take place. While the 5 apparent stability of soil brGDGTs during this incubation experiment reinforces their 6 potential as tracers for land-sea transport of soil organic carbon and their use in 7 paleoclimate reconstructions, the distributional differences between GDGTs in river 8 water and nearby soil, as well as in river and ocean water, indicate that further 9 research is needed to pinpoint the sources of GDGTs that are ultimately discharged to 10 the oceans and are subsequently archived in continental margin sediments.

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#### 12 **1** Introduction

13 The global carbon cycle encompasses a myriad of biogeochemical processes that 14 influence our climate and link all carbon reservoirs on the Earth surface. Soils are 15 considered to play a very active and fundamental role in this cycle, as their 16 mobilization from land and subsequent deposition in marine sediments serves as a 17 long-term sink of atmospheric CO<sub>2</sub>. However, the exact magnitude and mechanisms 18 of soil organic carbon (OC) transfer from terrestrial source to marine sink remain 19 elusive (e.g. Cole et al., 2007; Weyhenmeyer et al., 2012), which partly stems from 20 the lack of a suitable tracer of the soil OC pool. Our current insights in land-sea soil 21 OC transport dynamics are primarily based on bulk properties of river suspended 22 particulate matter (SPM). However, next to soil OC, river SPM also comprises carbon 23 derived from aquatic production and 'fossil' carbon from rock erosion, each of which 24 influences the bulk properties of SPM (Blair et al., 2010).

25 Analytical advances in the field of organic biogeochemistry have yielded a growing 26 number of powerful tools with the potential to exclusively target components of 27 specific pools, including soil OC. In this context, branched glycerol dialkyl glycerol 28 tetraethers (brGDGTs; Fig. 1) have been put forward as tracers for soil-derived OC in 29 carbon transport studies (e.g. Hopmans et al., 2004; Kim et al., 2006). Since their 30 discovery (Sinninghe Damsté et al., 2000; Schouten et al., 2000), brGDGTs have been 31 found in soils and peats worldwide (Weijers et al., 2007b, 2009; Liu et al., 2010). The 32 exact organism(s) that produce(s) these compounds have yet to be identified, but 33 current evidence points towards an origin from heterotrophic soil bacteria (Pancost

1 and Sinninghe Damsté, 2003; Oppermann et al., 2010; Weijers et al., 2010) from the 2 phylum of Acidobacteria (Weijers et al., 2009; Peterse et al., 2010; Sinninghe Damsté 3 et al., 2011). BrGDGTs have been used as a proxy for the relative input of fluvially 4 transported soil OC in marine systems based on their abundance in coastal marine 5 sediments relative to that of crenarchaeol (Fig. 1), an isoprenoid GDGT (isoGDGTs) 6 produced by marine Thaumarchaeota (Sinninghe Damsté et al., 2002), parameterized 7 as the Branched and Isoprenoid Tetraether (BIT) index (Hopmans et al., 2004). 8 Moreover, subtle variations in the molecular distribution of the brGDGTs have been 9 found to relate to mean annual air temperature and soil pH (Weijers et al., 2007b), 10 indicating their potential for utility in paleoclimate studies. Indeed, using a 11 combination of the Methylation of Branched Tetraethers (MBT) and Cyclisation of 12 Branched Tetraether (CBT) indices (the so-called "MBT-CBT" proxy; Weijers et al., 2007b, recently revised as MBT'-CBT by Peterse et al., 2012), down-core variations 13 14 in brGDGT distribution in fluvially-dominated continental margin sediments have 15 been used to obtain an integrated climate history of the adjacent drainage basin (e.g. 16 Weijers et al., 2007a).

17 Although GDGT-based proxies are finding increase use, our understanding of 18 mobilization and transfer of soil OC and associated GDGT signals to fluvial 19 environments, and the processes acting upon these components during transport are 20 still poorly understood. There is emerging evidence, for example, for in situ 21 production of brGDGTs in aquatic environments including high-latitude fjord systems 22 (Peterse et al., 2009), open shelf sediments (Zhu et al., 2011), lakes (e.g. Sinninghe 23 Damsté et al., 2009; Tierney and Russell, 2009), and most recently, rivers (De Jonge 24 et al., 2014; Kim et al., 2012; Zell et al., 2013b; Zhang et al., 2012). The presence of 25 intact polar lipid (IPL) precursors of brGDGTs in SPM from the Amazon (Zell et al., 26 2013b), Pearl (Zhang et al., 2012), and Yenisei rivers (De Jonge et al., 2014) provides 27 strong evidence for aquatic brGDGT production, as the IPL head groups are thought 28 to be rapidly lost upon cell death (Harvey et al., 1986; White et al., 1979), although 29 recent studies suggest that IPLs with ether bound headgroups may actually have very 30 slow turnover rates in marine sediments (e.g. Logemann et al., 2011; Xie et al., 2013). 31 Regardless, the core lipids (CLs) that are released after IPL degradation are 32 considered to represent 'dead', or fossil material - the fraction that is considered to be 33 stored in sedimentary archives - and targeted for paleoclimate reconstruction. Hence,

during fluvial transport from land to sea, the initial soil-derived brGDGT distribution
 entering a river may be modified by the addition of aquatic produced brGDGTs with a
 potentially different signature, but also by microbial degradation and transformation
 of the soil brGDGTs.

5 In order to explore the latter process, i.e. the behavior of soil brGDGTs upon entering 6 an aquatic system, we performed a laboratory experiment in which we incubated soil 7 using river water or ocean water from near the mouth of the same river as microbial 8 inocula. In the frame of a larger study on fluvial transport of soil organic carbon from 9 land to sea on the South Island of New Zealand, we used soil and river water from the 10 Rakaia River catchment for the experiment. In addition, the braided character of the 11 Rakaia River may reduce the potential contribution of in situ produced brGDGTs due 12 to the generally harsh and unstable conditions in such river systems. This should 13 enable the monitoring in a controlled setting of what could also be an actual soil-14 derived brGDGT signal in the river. For that matter, concentration and distributional 15 changes in IPL-derived and CL-brGDGTs were determined at different time intervals 16 during the incubation experiment. Although isoGDGTs (Fig. 1) are primarily 17 associated with marine archaea, they are also produced by soil Thaumarchaeota 18 (Sinninghe Damsté et al., 2012), albeit in small amounts relative to the brGDGTs 19 (Weijers et al., 2006b). In the marine realm, changes in isoGDGT distributions form 20 the basis for the  $TEX_{86}$  index, which is used as a proxy for sea surface temperature 21 (SST; Schouten et al., 2002; Kim et al., 2010). Large contributions from soil 22 isoGDGTs to the total isoGDGTs pool in marine sediments may consequently 23 complicate the interpretation of TEX<sub>86</sub>-derived SST records from near continental 24 margins (Weijers et al., 2006b). Therefore, all soil-derived GDGTs, i.e. both branched and isoprenoidal, were monitored during the experiment. 25

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# 27 2 Materials and methods

28 **2.1** Sample collection and incubation setup

The Rakaia River is one of the largest braided rivers on the South Island of New Zealand and has a mean annual discharge of 203  $m^3/s$ . The Rakaia is estimated to contribute 4.15 Mt of sediment a year, accounting for approximately 5% of the total South Island sediment yield to the adjacent continental shelf (Hicks et al., 2011). The

1 river originates in the Southern Alps and flows through the Canterbury Plains before 2 reaching the Pacific Ocean approximately 150 km from its source. Fresh water from the river was sampled in jerry cans close to the town Rakaia (S 43°44'57.9", E 3 172°01'52.9"), about 20 km upstream from the river mouth and outside of the tidal 4 5 influence. The ocean water used as inoculum was collected from the shoreline (S 43°54'15.1", E 172°11'41.9"), close to the river mouth (Fig. 2). Based on pH 6 7 measurements, this water (pH 7.6) was strongly fluvially influenced (pH 7.5). Surface 8 soil (0-10 cm; pH 5.8) was collected on the Canterbury Plains within the Rakaia River 9 catchment (S 43°52'00.8", E 172°09'10.8"). All soil material and incubation water 10 was collected in January 2012 and directly transported to the University of Otago, 11 where the incubation experiment was set up the next day. SPM was collected by 12 filtration of river and ocean water (100L onto a 0.2 µm polyethersulfone membrane 13 and 10L onto a pre-combusted 0.7 µm GF/F, respectively) in the field, after which the 14 filters were stored frozen.

15 In the laboratory at the University of Otago, the soil was sieved over a 2 mm mesh to 16 remove roots and homogenize the sample, after which two 10 g subsamples of the soil 17 were directly frozen at -20°C for subsequent determination of initial GDGT 18 composition. For the incubations, twenty-eight 250 ml bottles were filled with 10 g 19 soil subsamples, after which 100 ml of river water or ocean water, untreated to 20 preserve the natural microbial community, was added to twelve bottles each. Distilled 21 water was added to the remaining four bottles to serve as control setup. The 22 incubation bottles were loosely plugged with cotton wool, creating quasi-aerobic 23 conditions, and placed on a shaker table from January to June 2012. Based on IPL 24 degradation rates observed in previous studies (Harvey et al., 1986; Logemann et al., 25 2011), two bottles with river water and two bottles with ocean water were taken from 26 the shaker table and immediately placed into a freezer (-20°C) at 1, 7, 14, 30, 91, and 27 152 days after the start of the experiment. Duplicate control samples were frozen after 28 91 and 152 days. At the end of the experiment all bottles were shipped on ice to ETH Zürich, Switzerland, where they were briefly thawed and the supernatant was pipetted 29 30 off. The soils were then freeze dried, and stored frozen at -20°C until subsequent 31 sample work-up and analysis.

#### 32 2.2 Total organic carbon and nitrogen analysis

The freeze-dried, homogenized soils were weighed in silver capsules and fumigated with 1M HCl for three days and neutralized with NaOH for two days in a desiccator to remove carbonates prior to total organic carbon (TOC) and total nitrogen (TN) analysis. The measurements were performed on a vario MICRO elemental analyzer at ETH Zürich.

#### 6 2.3 GDGT extraction and analysis

7 Freeze-dried soil samples (3-4 g) were solvent-extracted three times using a Bligh and 8 Dyer technique modified from Sturt et al. (2004). In short, a solvent mixture of 9 MeOH:dichloromethane (DCM): phosphate buffer at pH 7.4 (2:1:0.8, v/v/v) was 10 added to the soils and ultrasonically extracted for 10 min. The extract was collected 11 each time after centrifuging at 1000 rpm for 5 mins. DCM and phosphate buffer were 12 added to the combined extracts to a volume ratio of 1:1:0.9 to obtain phase separation. 13 The DCM phase, containing the GDGTs, was collected after centrifuging, and the 14 remaining solvent was rinsed twice with DCM. The combined DCM phases were 15 dried under N<sub>2</sub> and passed over a silica column (deactivated with 1 weight% water) to 16 separate CLs and IPLs according to (Pitcher et al., 2009), with the exception that 17 hexane:ethyl acetate 1:1 (v/v) was used to elute the CLs. An aliquot of the IPL 18 fraction was analyzed directly for CLs in order to assess potential carry over. The 19 remainder of IPL fraction was dissolved in 6N HCl in MeOH and heated at 100°C for 20 at least 3h to release IPL-bound CLs. A known amount of C46 GDGT standard 21 (Huguet et al., 2006) was added to all fractions prior to analysis.

22 The filters were freeze dried and extracted with a MARS Xpress microwave 23 extraction system, using DCM:MeOH 9:1 (v/v). After centrifugation (5 min at 400 24 rpm), the total lipid extract (TLE) was pipetted off and the residues were rinsed twice 25 with DCM:MeOH 9:1 (v/v). The combined extracts were dried under N<sub>2</sub> with a 26 known amount of C<sub>46</sub> GDGT standard, after which the TLEs were separated into an 27 apolar and a polar (GDGT) fraction by passing them over a silica (1% water 28 deactivated) column using hexane:DCM 9:1 (v/v) and DCM:MeOH 1:1 (v/v), 29 respectively.

30 All GDGT-fractions were dissolved in hexane:isopropanol 99:1 (v/v), filtered over a 31 0.45  $\mu$ m PTFE filter, and analyzed using high performance liquid 32 chromatography/atmospheric pressure chemical ionization – mass spectrometry

1 (HPLC/APCI-MS) with an Agilent 1260 Infinity series LC/MS at ETH Zürich 2 according to Schouten et al. (2007a). Separation of the GDGTs was achieved with a 3 Grace Prevail Cyano column (3µm, 150×2,1mm) after passing through a guard 4 column of the same material  $(5\mu m, 7,5\times 2,1mm)$  with hexane: isopropanol (99:1, v/v)as an eluent at a flow rate of 0.2 ml/min. The GDGT-fractions eluted isocratically 5 6 with 90% A and 10% B for 5 min, and then with a linear gradient to 18%B for 34 min, where A=hexane and B=hexane:isopropanol 9:1 (v/v). Selective ion monitoring 7 8 of the [M+H]<sup>+</sup> was used to detect the different GDGTs. Although brGDGTs appear to 9 have a higher response factor than isoGDGTs (Schouten et al., 2013a), quantification 10 was done assuming similar response factors for all GDGTs and the internal standard. 11 Hence, reported brGDGT concentrations are likely overestimated compared to those 12 of the isoGDGTs. The amounts of all GDGTs are given as the average and the range 13 of variation of the duplicate incubation samples. The amount of GDGTs in each 14 sample has been corrected for carry over of CLs into the IPL fraction, which was on 15 average <8% for brGDGTs and <1% for isoGDGTs.

#### 16 **2.4 GDGT-based index calculations**

17 The CBT and MBT' indices based on brGDGTs were calculated according to18 (Weijers et al., 2007b):

19 
$$CBT = -\log ((Ib + IIb) / (Ia + IIa))$$
 (Eq. 1)

20 and (Peterse et al., 2012):

21 MBT' = 
$$(Ia + Ib + Ic) / (Ia + Ib + Ic + IIa + IIb + IIc + IIIa)$$
 (Eq. 2)

22 The BIT index was calculated following (Hopmans et al., 2004):

23 
$$BIT = (Ia + IIa + IIIa) / (cren + Ia + IIa + IIIa)$$
 (Eq. 3)

and for the TEX<sub>86</sub> index based on isoGDGTs the equation of (Schouten et al., 2002)
was used:

26 
$$TEX_{86} = (GDGT-2 + GDGT-3 + cren') / (GDGT-1 + GDGT-2 + GDGT-3 + cren')$$
  
27 (Eq. 4)

Roman numerals and GDGT names refer to the molecular structures in Fig. 1.
Regular reruns of selected samples on the HPLC-MS at ETH show that the analytical
error on the indices is <0.01.</li>

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# 2 3 Results and discussion

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## 3.1 TOC and TN concentrations over time

The TOC concentration in the soil is 1.6-1.7% at t=0, and varies in a range from 1.2-1.7%, whereas the TN concentration is 0.15% at t=0 and remains practically stable (0.12-0.15) over the course of the experiment (Fig. 3). Given the limited range of variation in TOC and TN concentrations, no direct influence of TOC or TN on the GDGTs in this experiment is assumed during the following discussion.

#### 9 **3.2 BrGDGT concentrations over time**

10 The total brGDGT pool (IPLs + CLs) initially present in the soil is  $239\pm13$  ng/g. In 11 the different incubation experiments these concentrations varied between 187-325 12 ng/g (river water), 186-322 ng/g (ocean water), and 252-407 ng/g (distilled water 13 control) (Fig. 4). Although the abundance of CL-brGDGTs seems to increase with time in the control setup, the trend is only weak and not significant ( $r^2 = 0.34$ , p =14 0.122) due to the large spread in concentration between the two replicate samples at 15 16 t=152 days (Fig. 4). Overall, the CL concentrations are within the same range 17 between different experimental setups and remain essentially constant through time. 18 Since the soils were incubated under natural light conditions, this suggests that soil-19 derived brGDGTs might not be sensitive to photodegradation during the incubation.

20 The contribution of IPLs to the total amount of brGDGTs in the soil at t=0 is 21  $12.4\pm0.3\%$  (Fig. 5), which is in the same range as has previously been reported for 22 temperate soils from the Netherlands, Scotland, and the UK (Peterse et al., 2010; 23 Weijers et al., 2011). The contribution of IPL-derived brGDGTs to the total pool 24 varies mostly within the same range in the river water (7-16%), ocean water (9-15%), 25 and control (9-17%) setups, and like the total amount of CL-brGDGTs, IPL-brGDGT concentrations show no strong trends or changes over time (river water:  $r^2 = 0.34$ , p =26 0.014; ocean water:  $r^2 = 0.00$ , p = 0.944; distilled water:  $r^2 = 0.20$ , p = 0.268). This 27 28 suggests that brGDGTs are either produced at the same rate that they are degraded, or 29 that production and/or degradation of brGDGTs occurs at such low rates that these 30 processes are not detected within the timeframe of this experiment. The apparent 31 stability of the IPL-derived, or 'living' brGDGT signature (Fig. 4) during the

1 experiment is surprising, given the perceived lability of IPLs in general and their 2 susceptibility to loss of their headgroups within a few days after cell lysis (Harvey et 3 al., 1986; White et al., 1979). However, previous studies have indicated that the type 4 of headgroup and the bond through which it is attached to a core lipid may influence 5 the rate of degradation, offering a potential explanation for the absence of changes in 6 IPL-brGDGT concentration in our experiment. For example, after a 96 hr aerobic 7 incubation of a mixture of IPLs with different headgroups in beach sediment, 70% of 8 the phospholipids had degraded, compared to only 3% of the glycolipids (Harvey et 9 al., 1986). Although a primarily glycosidic headgroup composition would be expected 10 based on these former results, the majority of IPL-brGDGTs in soils, peat, and lake 11 sediments are thus far identified as phospholipids (Weijers et al., 2011; Peterse et al., 12 2011; Tierney et al., 2012), implying that they should be sensitive to degradation, 13 counter to our observations. That this may not be the case may potentially be 14 explained by the findings of Logemann et al. (2011), who recently showed that the 15 bond type within the CL, rather than the type of headgroup and its connection to that 16 CL, determines IPL sensitivity to degradation. In their experiment, membrane lipids 17 in which the side chains were ester-bound, as is common for bacteria, started to 18 rapidly degrade within the first 5 days of the incubation period, whereas the 19 abundance of lipids with ether-bound chains remained invariant during the entire 97-20 day experiment. BrGDGTs are thought to be produced by bacteria, but they also 21 possess archaeal traits, including their tetraether structure (Weijers et al., 2006a). As a 22 consequence, the presence of at least four ether bonds in each brGDGT may thus not 23 only protect the IPL, but also the CL-brGDGTs from degradation during our 24 experiment, even when the IPLs possess a phospho head group.

25 Weijers et al. (2010) determined a turnover rate for the total pool of CL brGDGTs in a 26 soil of about two decades, which implies that any changes in brGDGT concentration 27 or distribution may indeed not yet be detectable after 152 days of incubation. On the 28 other hand, a recent study by Huguet et al. (2013) indicated that the brGDGT 29 signature in a French peat bog had completely adapted to the 2°C maximal daytime 30 temperature increase induced by the placement of open top mini-greenhouses within a 31 period of less than 26 months. Although these results suggest that brGDGTs are 32 turned over at a substantially faster rate, significant changes were only observed in the 33 latter phase of the warming experiment, and were primarily distributional, as their concentration remained the same (Huguet et al., 2013). The unaltered IPL-brGDGT concentrations in our experiment may thus be an indication that the time frame of our incubation has been too short to reveal any degradation processes based on IPLbrGDGT abundance. However, since the type of headgroup attached to the GDGTs has not been identified, our data does not allow to fully exclude equal production and degradation rates as possible explanation for the apparently stable IPL-brGDGT concentrations during the experiment.

8 While nutrient conditions in the river and ocean water added to the soil are 9 presumably different, this does not seem to have had any effect on brGDGT 10 concentrations. Moreover, given the soil:water ratio (10 g in 100 ml) used in this 11 experiment, the amounts of nutrients that are released from the soil after the addition 12 of water is likely substantially higher than the concentrations initially present in the 13 river and ocean water. Nevertheless, the effect of nutrient availability on brGDGTs in 14 soils has not yet received much attention, although recently was shown that variations 15 in the amount and distribution of brGDGTs in lake sediments did not relate to water 16 column nutrient concentrations (Loomis et al., 2014).

# 17 **3.3** IsoGDGT concentrations over time

18 Crenarchaeol is the most abundant isoGDGT in the New Zealand soil at the start of 19 the incubation, although its total amount (CLs + IPLs;  $63\pm7$  ng/g), as well as that of 20 all other isoGDGTs (50±5 ng/g) is low compared to the total pool of brGDGTs 21 (239±13 ng/g). In contrast, the average fraction of isoGDGTs initially present in the 22 soil as IPL is 54% for crenarchaeol to 76% for isoGDGT-2, which is substantially 23 higher than proportion of brGDGTs occurring as IPLs (12%) at the commencement of 24 the experiment. Similar proportions of IPL-derived crenarchaeol have been found in 25 soils from the Amazon (~50%; Zell et al., 2013b). The total concentration (IPLs + 26 CLs) of all isoGDGTs and increases substantially during the experiment (Fig. 4), and that of isoGDGT-0 in the ocean water setup has even doubled by the end of the 27 incubation (24 $\pm$ 2 to 48 $\pm$ 5 ng/g, r<sup>2</sup> = 0.58, p = 0.000). 28

The concentrations of the individual IPL-derived isoGDGTs vary between the different incubation setups, but are overall highest in ocean water (Fig. 4; Supplementary Table). The difference in IPL-isoGDGT concentration between fresh and saline water incubations may potentially be a result of the addition of varying

1 amounts of aquatic isoGDGTs to the soil-derived isoGDGT pool. However, 100 ml of 2 ocean water only contained ~0.5 ng isoGDGTs, which is less than 1% of their initial 3 concentration in the soil. Alternatively, the isoGDGTs could be increasingly released 4 from the soil matrix upon mixing with (saline) water, although the absence of a 5 simultaneous increase in brGDGTs suggests that this is not the most likely scenario. 6 Finally, since *Thaumarchaeota* play an important role in the nitrogen cycle in soils 7 and the marine environment (Leininger et al., 2006; Wuchter et al., 2006; Zhang et al., 8 2010), nutrient availability may provide another explanation for the concentration 9 differences between experiments. So far, crenarchaeol has been found in all 10 (enrichment) cultures of ammonia-oxidizing Thaumarchaeota, and is thus considered 11 as a biomarker lipid for this group (de la Torre et al., 2008; Pitcher et al., 2010, 2011; 12 Sinninghe Damsté et al., 2012), although a recent study suggested that Marine Group 13 II Euryarchaeota may also produce crenarchaeol (Lincoln et al., 2014). Nevertheless, 14 the relatively high crenarchaeol concentration in the incubation setup using ocean 15 water could then be a result of ammonium release due to the higher abundance of exchangeable cations (e.g.  $Na^+$ ,  $Mg^+$ ) in ocean water compared to fresh water. This is 16 in agreement with the common method to determine available NH<sub>4</sub><sup>+</sup> in soils, which 17 18 involves shaking soil in a KCl or Na<sub>2</sub>SO<sub>4</sub> solution to extract ammonium from the soil 19 matrix (e.g. Mehlich, 1953). The concentration profiles of the other isoGDGTs are 20 comparable to that of crenarchaeol (Fig. 4), suggesting that they are most likely also 21 primarily derived from soil Thaumarchaeota.

22 Despite the significant increase of the total pools of crenarchaeol and isoGDGTs 0-3 in the river ( $r^2 = 0.74$ , p = 0.000 for crenarchaeol;  $r^2 = 0.56$ , p = 0.000 for isoGDGTs), 23 ocean water ( $r^2 = 0.30$ , p = 0.014 for crenarchaeol;  $r^2 = 0.40$ , p = 0.003 for 24 isoGDGTs), as well as control setups ( $r^2 = 0.52$ , p = 0.034 for crenarchaeol; the 25 increase of isoGDGTs is not significant  $r^2 = 0.24$ , p = 0.215), there is no statistically 26 27 significant increase in the concentration of individual IPL-derived isoGDGTs during the course of the incubation experiment (e.g. max.  $r^2 = 0.18$  for IPL-derived 28 crenarchaeol in ocean water; Fig. 4). We assume therefore that the production rate of 29 30 IPL-isoGDGTs is approximately constant during this time. The proportional decrease 31 of IPL-isoGDGTs with time (Fig. 5) subsequently indicates that the increase in the 32 overall isoGDGT pool is a result of CL accumulation. This in turn confirms the 33 general finding that the turnover of the IPL pool, and thus the release of CL-

isoGDGTs, is faster than the degradation of these CLs. Regardless, the increase in
isoGDGT concentrations in all incubation setups suggests that availability of water is
likely more important that the type of water that is added to the soil.

#### 4 3.4 GDGT distribution changes

5 To evaluate if and how soil GDGT signatures are modified during the incubations, we 6 calculated brGDGT and isoGDGT-based indices commonly used in paleoclimate and 7 carbon cycle studies (i.e. the MBT', CBT, BIT, TEX<sub>86</sub>: Supplementary Table).

8 Next to the absence of changes in brGDGT concentration, also the MBT' and CBT 9 indices for the ocean and river water experiments exhibit minimal variation over time, remaining within a range of 0.04 for the MBT' index, and 0.02 for the CBT (Fig. 6). 10 11 Surprisingly, the largest changes are observed in the control experiment using distilled 12 water. The MBT' index for the IPL-derived brGDGT fraction in the control setup increases from 0.23±0.00 to 0.29±0.04 ( $r^2 = 0.57$ , p = 0.023; Fig. 6), and is primarily 13 caused by an increase in the concentration of brGDGT-Ia (from 23±0.3 to 29±3.9%). 14 15 A closer look at the relative distributions of each of the brGDGT types at t=0 and 16 t=152 days reveals a subtle shift in the contribution of IPL-derived brGDGT-III 17 (decreasing from 23±1.3 to 19±0.1%) to brGDGT-II (increasing from 54±0.4 to 18 58±0.4%) in river water (data not shown). Since both brGDGTs-II and III in the 19 denominator of the MBT' index (Eq. 2), and the fraction of IPL-derived brGDGT-I 20 remains stable, this change is not reflected by the index values. Given that the 21 absolute amounts of IPL-derived brGDGTs do not significantly change with time, it is 22 hard to determine the exact processes that cause the distributional changes, as both in 23 situ production and preferential degradation of specific brGDGTs could influence the 24 total concentration. Since brGDGTs with cyclopentane moieties were only present 25 below detection limit in most of the IPL-derived fractions, it was not possible to 26 reliably calculate CBT index values for any of these fractions.

The BIT index is clearly influenced by the substantial increase in crenarchaeol during the experiment, and starts to decrease between t=30 and t=91 days in all water types (Fig. 6). Due to the large percentage of crenarchaeol that is present as IPL compared to that of the brGDGTs, the BIT index of the IPL-derived fraction is much (0.48) lower than that of the CL fraction (0.88) at t=0. The lowest BIT index value for the IPL-derived fraction is 0.35 at t=152 days in ocean water (Fig. 6), and stems from the

1 increase in crenarchaeol (and isoGDGTs in general) in this experiment. The trend of a 2 lower BIT value for the IPL versus the CL fraction has also been found for river SPM 3 from the Yenisei (De Jonge et al., 2014) and Amazon (Zell et al., 2013b), where the 4 difference was explained by a lower degradation rate of soil-derived crenarchaeol 5 IPLs compared to brGDGTs and/or a contribution of *in situ* produced crenarchaeol 6 IPLs in the river (Zell et al., 2013b). Our incubation results suggest that the 7 degradation of IPLs from soil-derived crenarchaeol is faster than that of brGDGTs, 8 and that the difference in BIT index between the IPL-derived and CL fractions is thus 9 mainly caused by the higher production rate of crenarchaeol. This is in agreement 10 with the recent finding that the BIT index of river SPM primarily tracks the seasonal 11 aquatic production of crenarchaeol rather than that of soil input (Yang et al., 2013; 12 Zell et al., 2013a). Our results furthermore indicate that care should be taken with 13 using the BIT index to constrain the input of fluvially transported soil OC into a 14 marine system. However, the stability of the soil brGDGT pool supports earlier 15 observations that tracing the absolute amount (as opposed to the relative abundance) 16 of brGDGTs in rivers during land-sea transport may provide a more reliable tracer of 17 soil OC at our study site (c.f. Fietz et al., 2011; Smith et al., 2012; Zell et al., 2013b). 18 Moreover, the spread in GDGT response factors between laboratories resulting in a 19 range of BIT values for the same sample provides another, independent argument to 20 use absolute concentration measurements (Schouten et al., 2013a).

21 TEX<sub>86</sub> index values start to change between 30 and 91 days of incubation for the IPL-22 fraction in river water and the CL-fraction in ocean water experiments (Fig. 6). We 23 find no clear explanation for the diverging trends for the different water types or IPL 24 versus CL fractions, but there are several factors that may contribute to the observed 25 changes. For example, shifts in the initial isoGDGT-community due to water contact, 26 different membrane adaptation rates, or preferential release from the soil matrix may 27 all have occurred. Alternatively, the relative distribution of isoGDGTs may differ per 28 type of headgroup in both marine (Schouten et al., 2008; Pitcher et al., 2011; Lengger 29 et al., 2012) and soil Thaumarchaeota (Sinninghe Damsté et al., 2012), so that the 30 deviation in TEX<sub>86</sub> values may also be explained by different turnover rates per 31 headgroup. However, even though the total variation in TEX<sub>86</sub> corresponds with a 32 4°C SST change in the experiment, given the low absolute amounts of terrestrial 33 isoGDGTs that are finally discharged to the ocean in this case, this will have a minor 1 impact on TEX<sub>86</sub> records from the marine environment where their abundances are
2 much higher.

### 3 3.5 Experimental setup evaluation

Despite prior evidence for in situ production of brGDGTs in several large rivers (e.g. 4 5 Kim et al., 2012; Zhang et al., 2012; Zell et al., 2013b; Yang et al., 2013; De Jonge et 6 al., 2014), we have not observed this in our experiment. The absence of an in situ 7 production signal in our data may be a consequence of the soil:water ratio of 1:10 that 8 we used in our experimental setup, which is likely more concentrated than in most 9 natural systems, so that the relatively large amount of soil-brGDGTs may have 10 overprinted any evidence of aquatic brGDGT production during the experiment. An 11 indication that in situ production in the Rakaia River may take place after all is 12 reflected by the offset between brGDGTs in river SPM and that in the soil (Fig. 6). 13 Although this soil-SPM offset is based on a single sample location in the entire river 14 basin and brGDGT distributions in soils may show substantial variation within a small 15 area (e.g. Weijers et al., 2007b; Naeher et al., 2014), it may also support the idea that 16 brGDGTs in soils and river SPM may have different sources. Notably, also the 17 brGDGT distributions in river and ocean SPM show an offset (Fig. 6), despite the 18 ocean water sample location close to the river mouth. Differences in biomarker 19 distributions in SPM from the fresh-saline water transition have previously been 20 attributed to hydrological sorting (e.g. Goñi et al., 1998), or, more specifically in case 21 of brGDGTs, additional marine production (e.g. Zhu et al., 2011; Zell et al., 2014). In 22 our case, this offset may have also been introduced by the use of filters with different 23 pore size to obtain the SPM from the river and ocean water (0.2  $\mu$ m and 0.7  $\mu$ m, 24 respectively). This has resulted in a comparison of 'free living' and 'suspended' lipid 25 fractions, which at least in marine water may have a different composition (Ingalls et 26 al., 2012; Close et al., 2014).

Furthermore, 152 days of incubation appears not to have been long enough to determine the sensitivity of brGDGTs to degradation in aquatic environments based on changes in their distribution or abundance. The unexpected apparent stability of the IPL-derived brGDGTs suggests that the presence of IPL-brGDGTs in aquatic systems does not necessarily indicate that they are produced in situ, and rather support the earlier finding that IPLs containing ether lipids are relatively resistant against degradation, even on a longer time scale than has previously been shown (t=97 days; Logemann et al., 2011). Alternatively, the water-saturated conditions may have inhibited both growth and degradation of brGDGTs to take place. However, without analyzing the exact headgroup composition of the brGDGTs in our experiment it is not possible to distinguish between these processes.

6 For this study, it was assumed that the soil-derived (IPL-)GDGTs would be degraded 7 in aquatic environments, yet our data indicate that the concentration of isoGDGTs 8 actually increased over time in all setups (Fig. 4). This suggests that isoGDGTs are 9 either produced, or that they are (preferentially) released from the soil matrix during 10 the experiment. In order to get a better control on the exact processes taking place, the 11 control setup could have benefitted from the addition of a chemical agent to prevent 12 any microbial activity, and thus to avoid additional GDGT-production during the 13 experiment.

14 To summarize, follow-up experiments could benefit from a more natural soil:water 15 ratio, so that a potential contribution of aquatic GDGTs can be detected by analyzing 16 the soils. The control setup should be treated to prevent any microbial activity. In 17 addition, determining the abundance and distribution of GDGTs present in the 18 incubation water could likely provide further information on water column processes, 19 provided that the amount of water added to the soil is sufficient to obtain a large 20 enough sample upon filtering for GDGT analysis. Finally, to better understand the 21 apparent stability of IPL-brGDGTs in aquatic environment, the type of headgroup 22 should be monitored during the incubation experiment, which should run longer than 23 the 152 days of this study.

24

### 25 4 Conclusions

Laboratory incubation experiments involving admixture of soil and fresh/ocean water indicate that soil-derived brGDGTs appear to be surprisingly stable in aquatic environments in both IPL- and CL configurations. Our observations suggest that soil brGDGT signatures will likely be unaltered during fluvial transport from land to sea, although any influence of aquatic degradation and/or production processes may be obscured due to the higher-than-natural soil:water ratio used in our experimental setup. Indeed, distributional offsets between brGDGTs in river SPM and catchment soil, as well as between brGDGTs in river SPM and ocean SPM indicate that additional sources may contribute to the final distribution in which brGDGTs are delivered to the ocean. This means that the exact sources of brGDGTs in a river system need to be well constrained before these compounds can be used as reliable tracers for land-sea transport of soil organic carbon, as well as for paleoclimate reconstructions.

7 As opposed to brGDGTs, the concentration of isoGDGTs increased during all 8 incubation experiments, altering their initial relative distribution. Although the 9 changes in TEX<sub>86</sub> index values correspond with a maximum change of about 4°C in 10 reconstructed SSTs in the ocean water setup, the low abundance of isoGDGTs in soils 11 relative to in marine settings suggests that their land-sea transport would have 12 minimal impact on TEX<sub>86</sub> recorded in the marine sedimentary record. Nevertheless, 13 after 30 days of incubation, the increase in isoGDGTs was sufficient to affect BIT 14 index values. The observed increase in isoGDGT concentration in the control setup 15 using distilled water indicates that the availability of water is more important for the 16 behavior of isoGDGTs than the properties of the aquatic system in which the soil is 17 introduced. Our findings may have consequences for the interpretation of  $TEX_{86}$ -18 based SST records for coastal marine settings with a BIT index close to the cutoff 19 value of 0.3 (Weijers et al., 2006b), as the actual input of soil-derived GDGTs upon 20 delivery to the ocean may be larger than anticipated.

21

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Figure 1. Molecular structures of the brGDGTs and isoGDGTs monitored during the
 incubation experiment.



- 1 Figure 2. Overview map of the Rakaia River, South Island, New Zealand, indicating
- 2 the sampling locations.



Figure 3. Total organic carbon (TOC) and total nitrogen (TN) concentrations in a
 sandy loam soil from the Raiaka River catchment, New Zealand, during incubation (t
 = 152 days) in river (RI), ocean (OC), and distilled (C) water.





Figure 4. Concentrations of core lipid (CL) and intact polar lipid (IPL)-derived a)
brGDGTs, b) crenarchaeol + cren', c) isoGDGT-1,2,3, and d) isoGDGT-0 in a sandy
loam soil from the Raiaka River catchment, New Zealand, during incubation (t = 152
days) in river (RI), ocean (OC), and distilled (C) water. Average concentrations are
indicated by solid (CLs) and dashed (IPL-derived) lines.



Figure 5. Percentages of the total pool of a) brGDGTs, b) crenarchaeol + cren', c)
 isoGDGTs-1,2,3, and d) isoGDGT-0 present in 'living', or intact polar lipid (IPL) derived form. Average concentrations are indicated by solid lines.



Figure 6. Distribution of core lipid (CL) and intact polar lipid (IPL)-derived GDGTs in a sandy loam soil from the Rakaia River catchment, New Zealand, as reflected by the a) MBT' index, b) CBT index, c) BIT index, and d) TEX<sub>86</sub> index during incubation (t = 152 days) in river (RI), ocean (OC), and distilled (C) water under quasi-aerobic conditions. Horizontal straight lines represent GDGT composition in river (black) and ocean water (grey) suspended particulate material (SPM). Average concentrations are indicated by solid (CLs) and dashed (IPL-derived) lines.



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