# 1 Geographically widespread <sup>13</sup>C-depletion of grazing

2 caddis larvae: a 'third way' of fuelling stream food webs?

3 Running Title: Widespread <sup>13</sup>C-depletion in grazing caddis 4 5 Aurora Sampson<sup>1\*</sup>, Nicola Ings<sup>1</sup>, Felicity Shelley<sup>1</sup>, Sarah Tuffin<sup>1</sup>, Jonathan Grey<sup>1, 2</sup>, Mark 6 Trimmer<sup>1</sup>, Guy Woodward <sup>1, 3</sup>, Alan Hildrew<sup>1, 4</sup> 7 8 <sup>1</sup>School of Biological and Chemical Sciences, Queen Mary University of London, London, 9 E1 4NS, United Kingdom 10 <sup>2</sup>Lancaster Environment Centre, Lancaster University, Lancaster, LA1 4YO, United 11 Kingdom 12 <sup>3</sup>Department of Life Sciences, Imperial College London, Silwood Park Campus, Buckhurst 13 Road, Ascot, Berkshire, SL5 7PY, United Kingdom <sup>4</sup> Freshwater Biological Association, Far Sawrey, Ambleside, Cumbria LA22 OLP 14 15 \*Corresponding author: aurorasampson1@gmail.com 16 17 18

#### Summary

- 1) Stream ecosystems are supported by both 'green' (i.e, based on grazing) and 'brown' (i.e. detritus) food webs, whereas methane-derived carbon is not considered generally to be important; here we add circumstantial evidence for this potential 'third way'.
- 2) Grazing cased-caddis (Trichoptera) larvae in the family Glossosomatidae can be very abundant in springs and headwaters and frequently have much lower stable carbon isotope ratios (i.e. they are depleted in the heavier <sup>13</sup>C stable isotope) than the biofilm ('epilithon') on the upper surfaces of the stones on which they live, and which is their presumed diet. Evidence for similar isotopic depletion in other lotic invertebrates is currently limited, however; even for glossosomatids it has been observed so far only in some streams draining the southern English cretaceous chalk and in a few headwaters in northern California. If this phenomenon proves to be more widespread, among streams or taxa, it could imply a more general underpinning of stream food webs by 'isotopically light' carbon derived from methane and accessed via consumers feeding on methanotrophic bacteria.
- 3) Here we sampled 58 stream sites to examine whether caddis larvae are also <sup>13</sup>C-depleted in streams draining other geologies. We focused mainly on carboniferous limestone and sandstone, as well as on further chalk streams representative of most of the British chalk aquifer: together, these new sites covered an area of almost 90,000 km², around three times greater than that surveyed previously.
- 4) At all 58 sites methane gas was supersaturated relative to the atmospheric equilibrium, and at 49 of them larvae were conspicuously <sup>13</sup>C-depleted (from -17.5 to -3.6%) relative to the bulk epilithon (components of which we know can oxidise methane). Although still most pronounced on chalk, this phenomenon was geographically and geologically much more widespread than shown previously and

suggests methane-derived carbon could indeed play a prominent role in stream food webs (i.e. the 'third way').

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*Keywords*: methane, carbon, armoured grazers, geology, stable isotopes, streams

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#### INTRODUCTION

much as half of all terrestrial net ecosystem production (Cole et al., 2007; Cole, 2013). Many fresh waters are sources of carbon dioxide and methane to the atmosphere (Striegl et al. 2012; Prairie & del Giorgio 2013), and Bastviken et al. (2011) suggested that, in terms of greenhouse gas potential, their methane emissions may offset the entire terrestrial continental sink of greenhouse gases. Running waters in particular are now recognised as major potential sources of methane, while the review by Stanley et al. (2015) found that river water was almost ubiquitously supersaturated with a mean concentration of 1.35 µM (about two orders of magnitude greater than atmospheric equilibrium). The total annual global emissions of 26.8 Tg of methane from running waters is similar to that released from wildfires, termites, methane hydrates and permafrost combined, and represents about half the total freshwater contribution and 20% of that from wetlands (Kirschke et al., 2013). This lends a new weight to the importance of the global carbon metabolism of rivers and streams. In fresh waters, research has focussed on the generation of methane as a potent greenhouse gas (Bastviken et al., 2004; Walter et al., 2006; Prairie & del Giorgio, 2013; Bridgham et al., 2013). However, not all the methane produced and transported by fresh waters is emitted to the atmosphere. Some is oxidised by methanotrophic bacteria (MOB) and converted to microbial

Inland waters play a key role in the global carbon cycle, transporting, burying or respiring as

biomass and carbon dioxide (CO<sub>2</sub>) (Chen & Murrell, 2010; Trimmer et al., 2012). Both these products can potentially pass through food webs: as microbial biomass consumed directly as food and in the form of CO<sub>2</sub> as an inorganic carbon source for photosynthetic organisms (Grey, 2016). However, the ecological significance of these pathways that divert dissolved methane from the atmosphere remains largely unknown. Secondary production and the trophic basis of food webs in rivers and streams have been widely studied, and the conventional understanding has been that they are based on organic carbon fixed by photosynthesis, either produced within the system (autochthonous) and/or imported from the land (allochthonous) (Hynes, 1970; Vannote et al., 1980; Thorp et al., 1998). The relative importance in various environmental settings of these respective "green" and "brown" pathways is still widely debated (Hein et al., 2003; Carpenter et al., 2005; Doucett et al., 2007; Risse-Buhl et al., 2012; Benke & Wallace, 2014; Carroll, Thorp & Roach, 2016; Guo et al., 2016, 2018; Brett et al., 2017; Neres-Lima et al., 2017). The direct consumption of methanotrophic bacteria, however, represents a potential 'third way', in which methanederived carbon could support river food webs (Trimmer et al., 2012; Shelley, Grey & Trimmer, 2014), as it can in lakes (Taipale et al., 2012). This pathway is of great interest, not only because methane oxidation moderates atmospheric emissions of a potent greenhouse gas, but also because methane-derived carbon is not widely acknowledged as being important in river food webs. The consumption of methanotrophic bacteria by animals can be detected by the marked depletion of <sup>13</sup>C in the tissues of consumers, and is widespread in lakes (Grey & Deines, 2005; Deines et al., 2007; Jones et al., 2008; Taipale et al. 2012). Similarly, low  $\delta^{13}$ C values in lotic animals could signal that the same phenomenon occurs in streams. Some of the first indications of this came from Japan, where Kohzu et al. (2004) found <sup>13</sup>C-depleted values in a variety of

macroinvertebrates in headwater streams, although this was mostly limited to animals found

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near anoxic zones rich in detritus, or in those that lived mostly in the hyporheic zone. More recently, Del Vecchia *et al.* (2016) also suggested that methane-derived carbon supported the biomass of five different species of stonefly (Plecoptera) in the hyporheic zone of four floodplain aquifers in the USA, again based on their low stable carbon isotope values.

The potential importance of methane-derived carbon as a basal resource for consumers in the surface waters of a southern English chalk stream draining a largely agricultural catchment was first suggested by Sanders *et al.* (2007), who showed that organic sediment patches under aquatic macrophytes exhibited intense methanogenesis in summer. Dissolved methane reached a mean summer peak of 1.5 µmol L<sup>-1</sup>, with pore-water concentrations two orders of magnitude higher. Sanders *et al.* (2007) showed that much of this methane was apparently oxidised *in situ*, and Shelley *et al.* (2015) found that this process is strongly substrate (i.e. methane) limited and could match the rate of methanogenesis. Further, methanotrophs produce (particulate) carbon equivalent to up to about 13% of that fixed by photosynthesis in some chalk streams (Shelley et al., 2017), with methanotrophic bacteria assimilating methane with an efficiency close to 50% (Trimmer *et al.*, 2015). A substantial amount of methane-derived microbial biomass is therefore potentially available to primary consumers.

Evidence that this potential food source was being exploited by primary consumers was first obtained by Trimmer *et al.* (2009), who found that the mean larval  $\delta^{13}$ C values of two common grazing cased-caddis (Trichoptera), *Agapetus fuscipes* (Glossosomatidae) and *Silo nigricornis* (Goeridae), in one English chalk stream (the River Lambourn) were anomalously low. At -41.2 and -40.4 ‰, respectively, these values were up to 11.1 ‰ lower than any of the various feasible autochthonous and allochthonous (all ultimately produced photosynthetically) food resources sampled. Whilst not the first recorded instance of low glossosomatid larval  $\delta^{13}$ C values (McNeely, Clinton & Erbe, 2006), methane had not been considered as a potential driver of  $^{13}$ C-depletion in caddis larvae.

All the evidence so far comes from a number of southern English chalk streams, which are groundwater-fed, free-flowing systems with very high alkalinity and primary production (Rovelli et al., 2017). Chalk is a form of very soft limestone formed by the shells of coccolithophores laid down in shallow Cretaceous seas and forming highly fractured and extremely permeable thick beds. It is globally a rather unusual geological formation, largely restricted to southern England, northern France and a few other areas of NW Europe (Darling and Gooddy, 2006). However, methane in river waters is abundant almost everywhere (Stanley et al. 2015) and, in UK aquifers, high concentrations in groundwater are by no means confined to the southern chalk (Darling and Gooddy 2006, Bell et al. 2017). This implies that methane could be far more widespread and important as a basal resource than suspected previously. Moreover, grazing glossosomatid caddis species, and potentially other specialist grazers, are often extremely abundant and widespread in suitable headwaters (mainly on various permeable strata, but not only in chalk streams) (Douglas, 1958; Elliott, 1982; Arens, 1990; Becker, 2011). Therefore, we aimed to measure the  $\delta^{13}$ C of glossosomatid larvae across a much greater number and variety of sites than we had done previously. On this evidence, we expected (1) that methane would be ubiquitous (at supersaturated concentrations) in streams on all geologies: in this study these were primarily chalk, carboniferous limestone and sandstone but included opportunistically a few sites draining other formations (permeable and non-permeable) where we knew the larvae occurred. If this proved to be the case, we also expected (2) that low glossosomatid  $\delta^{13}$ C values, relative to epilithon and other putative food resources, would be widespread, suggesting a general role for methane in stream food webs. Finally (3), because methane oxidation by methanotrophs is limited by the supply of the dissolved gas, both epilithic and caddis larval  $\delta^{13}$ C should correlate with

stream methane concentration rather than with the bedrock geology of the streams per se.

#### **METHODS**

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#### Site selection.

Fifty-eight streams were chosen on the basis of geology and the occurrence of glossosomatid caddis larvae (either *Agapetus fuscipes* or *Glossosoma conformis*). Fifty-five of the streams were drawn from various permeable geological formations: chalk (n=31), limestone (9), sandstone (14) and from one site on the south Wales coalfield (mixed sedimentary rock formations of coal, mudstone, siltstone and sandstone) (Table 1; Fig.1). The three remaining sites were on London clay (2) while the last was an isolated spring amongst the volcanic geology of the NW Lake District (mafic lava). Of the 31 sites on chalk, 14 lay outside the area of central southern England sampled in previous work (Trimmer *et al.*, 2009; Shelley *et al.*, 2014), and were either in the north of the Thames basin (sites 1, 8, 9, 41, 44, 52, 53 and 58 in Fig. 1; Table S1), to the south and east of London (sites 2, 3, 42 and 55), or towards the northern extent of the chalk in England (sites 37 and 38). Samples were collected in the spring or summer of 2010, 2011 or 2014 (see Table 1).

#### **Ambient stream water methane concentration**

Samples (analytical replicates) of surface water (n=5) were taken from the mid-channel and mid-depth at each site. Each sample was drawn up with a 60 mL gas-tight syringe (after Pretty, Hildrew & Trimmer 2006) and discharged through a 150 mm length of polytetrafluoroethylene tubing into a gas-tight vial (12.5 mL Exetainer, Labco, High Wycombe, UK), allowed to overflow (~three times) to minimise outgassing to the atmosphere. Samples were immediately fixed with 100 μL ZnCl<sub>2</sub> (50% w/v) bactericide and capped. In the laboratory, a 2 mL headspace of analytical grade helium (BOC) was introduced into each vial using a two-way valve and 5 mL gastight syringe. Samples were placed on rollers (Denley Spiramix, Thermo Denley, Colchester, UK) to allow the gas inside the vials to equilibrate between the water and the headspace. After this, a sub-sample of the headspace (100 μL) was injected into a gas

chromatograph fitted with a flame ionising detector (GC-FID; Agilent Technologies, UK Ltd., South Queensferry, UK (for details see Sanders *et al.* 2007). The methane concentration in the headspace of the vial was calculated from the area of the peak, calibrated against the areas of known standards. The total amount of methane gas in each vial (headspace + water) was calculated using solubility coefficients (Yamamoto, Alcauskas & Crozier, 1976). The mean of the methane concentration in these five samples yielded an overall average value for each site which was then used in subsequent analyses.

### Collection of caddis larvae and putative food resources for stable isotope analysis.

We sought to collect sufficient cased-caddis larvae for at least three measurements (analytical replicates) of  $\delta^{13}$ C (each ~ 15 individuals and totalling ~ 0.5 mg dry mass) to contribute to an overall mean  $\delta^{13}$ C value for caddis for each stream that could be used for further analysis (i.e. treating each stream as a replicate). Caddis were picked directly from rocks and gravels, placed in 50 mL tubes (Falcon, Fisher Scientific, Loughborough, UK) and frozen immediately, therefore including in the sample some gut contents (a potential source of error). In streams where larvae were scarce, fewer analytical replicates could be collected (Table 1).

Glossosomatidae are known to graze on epilithon (both algal and detrital components (Becker, 1990) but, in order to understand how larvae acquire low  $\delta^{13}$ C values, we also collected other

feasible stream food sources (e.g. macrophytes, mosses, filamentous algae and allochthonous leaf litter) for  $\delta^{13}$ C analysis. Epilithon was removed with a toothbrush from the substratum and leaf litter, macrophytes, mosses and filamentous algae were collected by hand. Three samples (analytical replicates, each ~ 0.8 mg dry mass) of each food resource, where present, were collected per site to gain mean  $\delta^{13}$ C site values for each food resource. Fewer than three such samples could be collected for some sites, where food resources were scarce. All samples were then stored in 50 mL tubes (as above) and, along with the caddis larvae, frozen in the field in

a portable freezer (17 L, Engel, Australia). In the laboratory, caddis larvae were thawed out and carefully removed from their cases using forceps under a dissecting microscope, cleaned of any debris and rinsed. Epilithon, terrestrial detritus and macrophytes were acidified (4 % HCl, 24h) to remove any inorganic carbon (Hedges & Stern, 1984). All samples of caddis and putative food sources were then oven dried to constant mass at 60 °C and ground to a fine

196 powder using an agate pestle and mortar.

Samples were weighed into ultra-clean tin capsules, then combusted using an elemental analyser coupled to a continuous flow isotope ratio mass spectrometer (CF/IRMS, Thermo-Finnigan, Delta Matt Plus, Bremen, Germany). Isotope calibration was carried out using the international standard for carbon (Ref. 8542, sucrose -10.47 ‰  $\delta^{13}$ C vs. Vienna-PeeDee Belemnite [VPDB], National Institute for Standards and Technology) and values are expressed using the standard delta ( $\delta$ ) notation with units of per mille (‰) (Equation 1).

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$$\delta = [(R_{SAMPLE} / R_{STANDARD} - 1)] *1000$$
 [Equation 1]

Urea was used to determine carbon content and check linearity, and Cyclohexanone-2,4-dinitrophenylhydrazone ( $C_{12}H_{14}N_4O_4$ ), was used as an internal standard to check the precision of the carbon stable isotope values (Trimmer *et al.* 2009).

#### Data analysis.

Statistical analyses were performed with the R statistical software v3.1.1. (R Core Development Team, 2014). To test for differences in methane concentration in the water between the three main geologies (chalk, limestone and sandstone), we performed a one-way ANOVA with a *post hoc* Games-Howell test to assess differences between pairs. Methane concentration was first log<sub>10</sub> transformed to normalise for positive skew.

Caddis larvae and each putative food resource  $\delta^{13}$ C values were pooled for each stream to provide one mean  $\delta^{13}$ C value for caddis and one for each food source per site for analysis (i.e. the stream was the replicate). Equation 2 was used to calculate the isotope difference between glossosomatid larval  $\delta^{13}$ C and bulk epilithon (after Fry 2006), where  $\delta 1$  is the  $\delta^{13}$ C (‰) of the larval sample and  $\delta 2$  is that of the epilithon:

 $\delta 1,2 \text{ (\%)} = [(\delta 1 - \delta 2) / (\delta 2 + 1000)] * 1000$  [Equation 2]

Potential variation in the trophic fractionation of  $\delta^{13}C$  from source to consumer was incorporated assuming a discrimination factor of  $0.4 \pm 1.20$  % (mean  $\pm$  1SD), after Moore & Semmens (2008). Thus, 95% of larvae were expected to fall within -1.95 to + 2.75 % of the epilithic baseline at each site.

To test whether caddis and epilithic signatures were related to bedrock type, two separate one-way ANOVAs with *post hoc* Games-Howell tests were performed, with caddis and epilithon  $\delta^{13}$ Cs (mean values per site) as the dependent variables and bedrock as the independent variable. Linear and polynomial regressions were used to assess whether there was a linear or curvilinear correlation between methane concentration (log transformed) and the  $\delta^{13}$ C of the caddis larvae or of the epilithon of all of the stream sites.

#### **RESULTS**

#### **Ambient methane concentration in stream water**

At every site, methane was supersaturated in stream water relative to the atmosphere (equilibrial value =  $0.002~\mu mol~L^{-1}$  at  $10^{\circ}C$ ), arithmetic mean ranging from 0.007 -  $0.58~\mu mol~L^{-1}$  with a mean of  $0.11~\mu mol~L^{-1}$  across the 58 streams, thus sustaining hypothesis (1). After logarithmic transformation of the data, chalk streams had significantly higher concentrations

(back-transformed mean 0.08  $\mu$ mol L<sup>-1</sup>) and, on average, around twice that of the untransformed values for those streams draining sandstone (0.04  $\mu$ mol L<sup>-1</sup>) (Fig. 2); concentrations in limestone streams were lower again, and half that of sandstone (0.02  $\mu$ mol L<sup>-1</sup>) (Fig. 2).

#### <sup>13</sup>C-depletion in caddis larvae relative to putative food resources

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Two species of glossosomatid caddis (Agapetus fuscipes and Glossosoma conformis) were found in the 58 streams surveyed, although they did not coexist at any site: Agapetus fuscipes was present in 56 streams and Glossosoma conformis in the two (sandstone) streams in South Wales (sites 13 and 20: Table 1). Caddis larval  $\delta^{13}$ C ranged from -44.3 to -27.0 %, and epilithon  $\delta^{13}$ C from -39.5 to -22.8 % (Table 1). Larvae were depleted in  $^{13}$ C relative to the epilithon at the same site by at least -1.95 ‰ (i.e. values were below the lower 95 % CL) at 49 of the sites, and in some cases by > -10 \% (Fig. 3) (sustaining hypothesis 2). This depletion varied among geologies but was not restricted to chalk. Larval  $\delta^{13}$ C values fell within the expected limits of potential fractionation relative to the epilithon at a further eight out of 58 sites: five draining sandstone, two on chalk and one on the coal measures (Fig. 3). One limestone stream (site 18) contained caddis larvae that were enriched (by 5.6 %) relative to the epilithon. Among the other feasible (even if unlikely) food resources available at each site, particulate detritus of terrestrial origin and various macrophytes were the most common (see Supporting Information Table S2). The  $\delta^{13}$ C of terrestrial detritus ranged from -31.0 to -27.2 ‰, and that of mosses and macrophytes from -44.5 to -24.2 ‰. In most cases, larvae had very depleted carbon isotope signatures relative to these alternatives (Fig. 4) and there was no consistent relationship between larval signature and any of the resources considered. Overall, no feasible mixture of these alternative food sources could explain the isotopic signature of the larvae at most of the sites.

#### The effect of geology

Caddis larvae exhibited statistically different  $\delta^{13}$ C values between the three main geological types of chalk, limestone and sandstone ( $F_{2,51}$ =13.79, P<0.001), with significantly more depletion on chalk than sandstone (P=0.003) but no other pairwise geological differences. Thus,  $^{13}$ C-depletion of caddis larvae was especially pronounced on chalk but also widespread elsewhere. There was no statistically significant difference in the  $\delta^{13}$ C of bulk epilithon among the three main geologies. This partially supports hypothesis 3.

## Caddis $\delta^{13}$ C and methane concentration

There was no overall relationship between methane concentration and caddis  $\delta^{13}C$  for the 58 streams (Fig. 5), nor between methane concentration and epilithon  $\delta^{13}C$ , thus the further stipulation of our third hypothesis, that bulk methane concentration would correlate with the stable carbon isotopic ratios of larvae and their putative food source (epilithon), was not supported.

#### **DISCUSSION**

#### Methane is ubiquitous

Methane exceeded atmospheric equilibrial concentrations at all 58 sites, supporting suggestions that supersaturated methane may be common in running waters worldwide (De Angelis & Lilley, 1987; Jones & Mulholland, 1998; Stanley *et al.*, 2015). Methane is therefore, a pervasive potential basal resource in lotic food webs, particularly since it is so readily and efficiently transformed into microbial biomass (Trimmer *et al.*, 2015; Shelley *et al.*, 2015). The methane in these streams is likely to have come from a variety of sources, including groundwater aquifers linked to underlying ancient oil and gas reserves (Cramer *et al.*, 1999;

Bell *et al.*, 2017) as well as more recent 'autochthonous' methane produced by *in-situ* anaerobic methanogenesis under macrophyte beds (Sanders *et al.*, 2007; Wilcock & Sorrell, 2008; Old *et al.*, 2014). This suggests there may be considerable amounts of previously ignored forms of both new and old carbon cycling through riverine food webs, with potentially important implications for river ecosystems.

Widespread  $^{13}$ C-depletion in cased caddis Irrespective of its ultimate provenance, remarkably low  $\delta^{13}$ C values in these grazing cased-caddis suggests that the incorporation of methane-derived carbon into stream food webs may be far more geographically and geologically widespread than lowland southern English chalk streams (Trimmer et al. 2009). If these low  $\delta^{13}$ C values are indeed caused by the assimilation of methane-derived carbon (Trimmer et al., 2009; Grey 2016), this has important implications for our wider understanding of the extent to which resources based on methane-derived carbon contribute to stream food webs in general.

The most straightforward explanation for <sup>13</sup>C-depleted caddis is that they are feeding on methanotrophic bacteria that have assimilated isotopically light carbon from methane, a pathway postulated by Trimmer et al. (2009). This suggests that many stream communities could be partially powered by chemosynthesis – the 'third way' of fuelling stream food webs. Recently, Shelley *et al.* (2017) demonstrated that the production of methanotrophs can be equivalent to up to about 13% of the total of net primary production in chalk streams (reaching a maximum in shaded patches and where methane is abundant). This represents a considerable potential basal resource for primary consumers and is in addition to the well-known "brown" (based on terrestrial detritus) and "green" (based on *in situ* photosynthesis) pathways in stream food webs (e.g. Trimmer et al. 2012).

Nevertheless, glossosomatid  $\delta^{13}$ C varied markedly across our 58 sites, suggesting the relative contribution of methanotrophic bacteria to the diet may be influenced by local factors such as larval density, micro-distribution and the availability of other food. The  $\delta^{13}$ C of other feasible, although highly unlikely, food sources were also explored as potential alternatives to epilithic biofilms, but in only a few cases (resources and sites) did they give a better match to the isotopic ratio of the larvae. At a few sites, moss (six sites), macrophyte tissues (two), or even leaf litter (three) could have contributed to the larval carbon isotopic ratio, but this was patchy and inconsistent (Fig. 5; Table S2): there is no compelling case to suggest the larvae are feeding widely on resources other than epilithon, which reflects what is known of their feeding ecology (Douglas, 1958; Arens, 1990; Becker, 1990, 2011; Nijboer, 2004). Are there other possible explanations for the depleted stable carbon isotopic ratios of these epilithic grazers? Chemolithoautotrophy driven by the oxidation of ammonium and reduced sulphur could both contribute <sup>13</sup>C-deplete carbon to the biofilm (Ruby, Jannasch & Deuser, 1987; Sakata et al., 2008). While we recognise the significant potential for ammonium oxidation in both chalk-gravels and sand sediments (Lansdown et al., 2016), its likely depletion in <sup>13</sup>C carbon could not account for that measured here in the caddis and we have no evidence to suggest significant sulphur cycling either (see Supporting Information; Fig. S1, Table S3 and supplementary Discussion). Further, in streams where dissolved inorganic carbon is very abundant, algae can discriminate against <sup>13</sup>C and take up the lighter <sup>12</sup>C isotope (Finlay, Power & Cabana, 1999; Finlay, 2001). As a consequence, algal cells favoured by grazing caddis may be more <sup>13</sup>C-depleted than the bulk biofilm within which they are embedded (and whose  $\delta^{13}$ C we measured). In addition, methanotrophs produce CO<sub>2</sub> as a product of methane oxidation. This CO<sub>2</sub> is likely to be <sup>13</sup>Cdepleted (although very scarce compared to the bulk DIC) and could 'label' constituent organisms, such as algae, disproportionately to other non-photosynthetic components. Where

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methanotrophic bacteria and algal cells are abundant, and in intimate proximity within the biofilm, carbon exchange may be more readily facilitated, so these small-scale transfers could be key to shaping the entry of methane-derived carbon into the wider food web.

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This last explanation for low  $\delta^{13}$ C values in caddis larvae, which can contribute a large fraction of overall secondary production in streams, demands that some algae or cell components are very highly <sup>13</sup>C-depleted relative to the bulk biofilm and are themselves grazed or assimilated highly selectively by the caddis. There is some support for selective feeding within the genus Glossosoma: McNeely et al., (2006) attributed the low glossosomatid  $\delta^{13}$ C values in small unproductive streams in Northern California to selective uptake of algae, although the influence of methane was not assessed in their study. Similarly, Glossosoma intermedium caddis larvae are thought to employ nutrient-specific foraging, grazing on phosphorus-rich periphyton on conspecific cases instead of phosphorus-limited periphyton on stream bed cobbles (Mooney, Strauss & Haro, 2014, 2016). Overall, therefore, the uptake of methane derived carbon in our study could be by direct ingestion of methanotrophs, which we think the most likely. Methane could also be accessed indirectly by grazing epilithic algae that have assimilated the very C<sup>13</sup>depleted inorganic carbon respired by methanotrophs. In either case (i.e. direct or indirect uptake), we would expect the amount of dissolved methane in stream water to correlate with the  $\delta^{13}$ C of the caddis larvae (since methane oxidation rates in rivers rise linearly with methane concentration; Shelley et al., 2014, 2015). This was not found, although methane oxidation rates can vary between different stream gravel biofilm communities (Shelley et al., 2014), and the exact circumstances in which methane-derived carbon is ingested and subsequently assimilated by caddis larvae remain uncertain.

Caddis larval  $\delta^{13}$ C could be independent of stream methane concentration if MOB are consumed only sporadically, perhaps as a "supplementary" resource. Methane-oxidising bacteria could thus constitute a kind of 'ship's biscuit' for these grazers – basic rations when

higher quality food is unavailable. Glossosomatid larvae characteristically form very high-density aggregations and have been shown to graze on the surfaces of neighbouring conspecifics' cases (Cox & Wagner, 1989; Cavanaugh, Haro & Jones, 2004; Mooney et al., 2014, 2016). Note also that biofilms on caddis cases were shown to oxidise methane (Trimmer et al., 2009) and thus must contain MOB. Little is known about the dietary quality of MOB as a food resource for invertebrate grazers. However, in feeding trials carried out by Taipale et al. (2012), the zooplankter *Daphnia magnia* could not grow and reproduce on MOB alone, but did so on various mixtures of bacteria with phytoplankton. On the assumption that depleted stable carbon isotope values in *Agapetus* were attributable solely to the direct assimilation of methanotrophs, Trimmer et al. (2009) used a simple mixing model to calculate that up to 30% of the biomass production of the caddis in a single stream could be accounted for by methanederived carbon. Similar assumptions here suggest this value is not exceptional and indeed, at the sites with the greatest disparity between caddis and bulk epilithon, estimates would evidently be even higher. Such a conclusion is still premature, however, since our evidence is circumstantial.

Overall, our study shows quite clearly that conspicuous <sup>13</sup>C-depletion of glossosomatid caddis larvae and high concentrations of dissolved methane are common. Widespread food-web linkages involving methane-derived carbon, probably via direct grazing of MOB or perhaps via photosynthetic uptake of depleted carbon respired by MOB, could be routine rather than exceptional. Distinguishing unequivocally between these two would require further evidence, such as the presence of assimilated biomarkers of MOB in the tissues of the caddis.

#### Acknowledgments

This work was funded by a Natural Environment Research Council grant (NE/H02235X/1) awarded to JG, MT, GW & AH. We thank Valentine Seymour for assistance in the field and

- 381 Michelle Jackson for helpful comments on an earlier version. The authors declare no conflict
- 382 of interest.

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#### 568 Figures and Tables

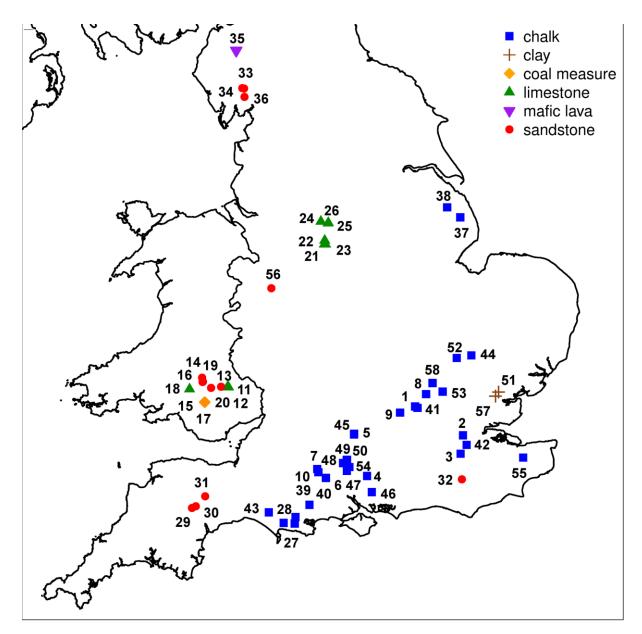
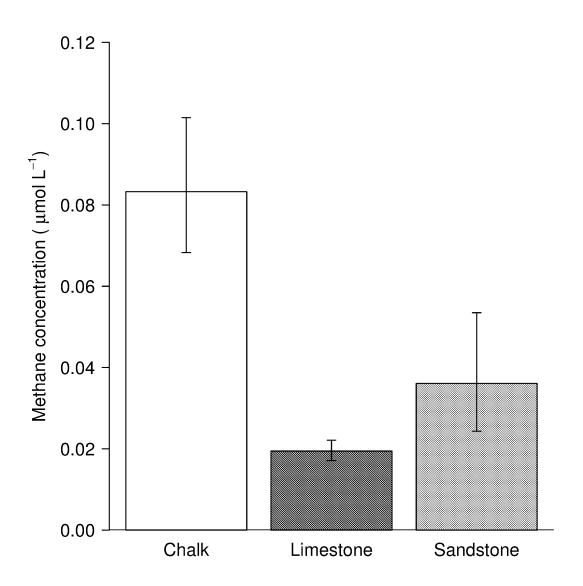
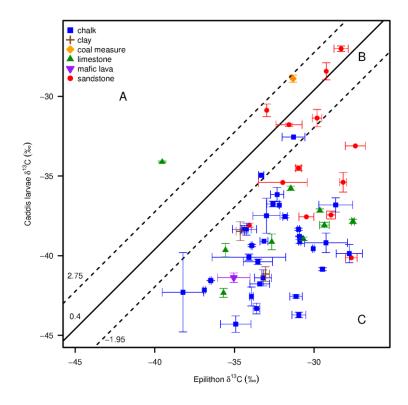


Figure 1. Map of England and Wales showing the 58 sites across six geologies 1. Chess (upper), 2. Cray, 3. Darent (upper), 4. Itchen, 5. Lambourn at Bagnor, 6.Test, 7. Till, 8. Ver, 9. Wye at High Wycombe, 10. Wylye, 11. Nant Onneu, 12. Nant Wenllan, 13. Nant Cleisfer, 14. Craig Cerrig Gleisiad (lower), 15. Craig Cerrig Gleisiad (middle), 16. Craig Cerrig Gleisiad (upper), 17. Dare, 18. Upper Tawe, 19. Traeth Mawr, 20. Owls Grove, 21. Manifold, 22. Milldale, 23. Dove, 24.Wye (upper), 25.Wye (middle) 26. Wye (lower), 27.Tadnoll, 28. Bere, 29. Creedy, 30. Stockleigh, 31. Lowman, 32. Crowborough Warren, 33. Hodgehow Wood, 34.

Belle Grange Beck, 35. Carrock Beck, 36. Ghyll Head, 37. Belleau, 38. Welton le Wold, 39. Allen, 40. Bourne, 41. Chess (lower), 42. Darenth, 43. Frome, 44. Granta, 45. Lambourn at Boxford, 46. Meon, 47. Test (upper), 48. Pillhill, 49. Andover, 50. Anton, 51. Brain, 52. Shep, 53. Mimram (lower), 54. Sheringham, 55. Great Stour, 56. Telford, 57. Ter, 58. Mimram (upper).



**Figure 2.** Mean ( $\pm$  1 SE) methane concentration ( $\mu$ mol L<sup>-1</sup>, after back transformation) for the three main geological types; chalk (n = 31), sandstone (n = 14), limestone (n = 9).



**Figure 3.** Caddis  $\delta^{13}$ C relative to epilithon  $\delta^{13}$ C (points are site means, limits around site points are calculated from analytical replicates). Black and broken lines are the mean expected larval signatures and upper and lower 95% CL, calculated assuming a fractionation of  $0.4 \pm 1.2\%$  (mean  $\pm 1$  SD; Moore & Semmens 2008) and that epilithon is the sole food source. Sites above the upper limit (area A on the plot) indicate larval enrichment in  $^{13}$ C, while those below indicate depletion (area C; sites within the expected limits are in area B). Colours indicate the different geologies.

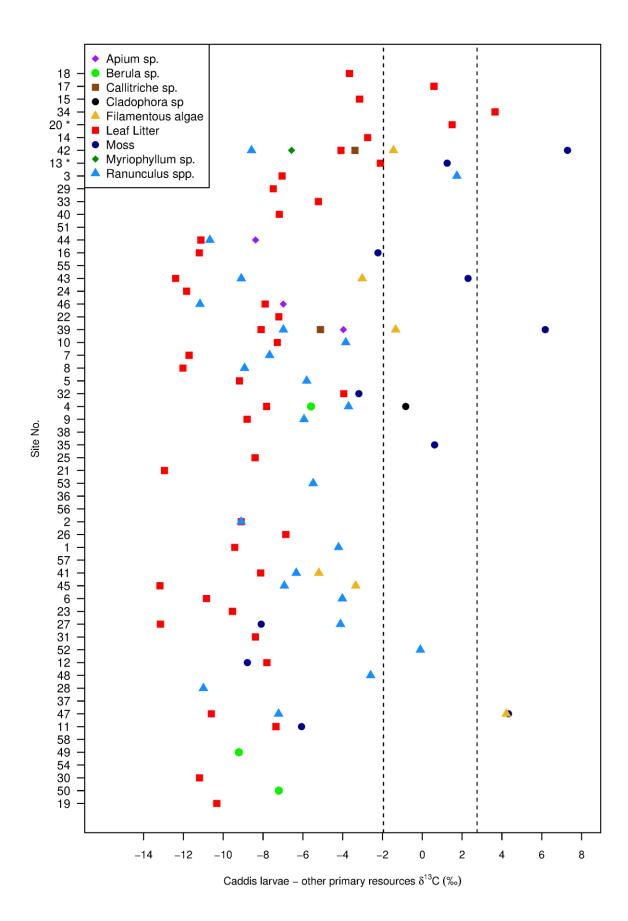
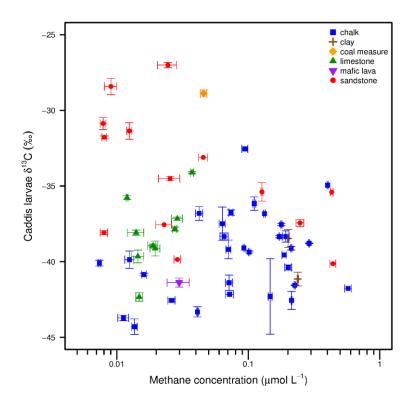


Figure 4. The  $\delta^{13}$ C of glossosomatid larvae, relative to alternative primary food resources. For each site, each symbol (specifying a putative resource) denotes the isotopic ratio of the caddis relative to that particular resource. Sites have been ordered by the largest negative difference between larvae and epilithon (from least difference at the top to greatest at the bottom). The area between the broken lines (95% CL) indicates the expected larval  $\delta^{13}$ C assuming a mean fractionation value of +0.4  $\pm$  1.2% (mean  $\pm$  1 SD; Moore & Semmens 2008).





**Figure 5.** Stream ambient methane concentration  $\pm$  1 SE (logarithmic scale) and caddis larval  $\delta^{13}$ C  $\pm$  1 SE, suggesting no overall difference in methane concentration among sandstone, limestone and chalk, but greater overall depletion of caddis on chalk than the others (see text for details).

**Table 1.** Mean  $\delta^{13}C \pm 1SE$  for epilithon and caddis larvae. Sites are grouped in to geologies then numerically by site number (facilitating comparison with Fig. 1). † indicates the two sites where *Glossosoma conformis* was found rather than *Agapetus fuscipes*. Where only two samples contribute to a mean, both values are given as a range.

Bedrock geology	Site no. (year	Epilithon δ13C	n =	Agapetus fuscipes δ13C	n =
Chalk	sampled) 1 (2010)	-30.93 to -30.91	2	-38.79 ± 0.02	3
Chalk	2 (2010)	-31.03 to -30.92	2	$-38.75 \pm 0.02$ $-38.35 \pm 0.08$	3
Chalk	3 (2010)	$-33.32 \pm 0.17$	3	-34.95	1
Chalk	4 (2010)	$-33.82 \pm 0.17$ $-31.82 \pm 0.14$	3	$-34.53$ $-37.54 \pm 0.07$	3
Chalk	5 (2010)	$-33.91 \pm 0.04$	3	$-39.37 \pm 0.07$	3
Chalk	6 (2010)	$-30.90 \pm 0.04$	3	$-39.12 \pm 0.07$	3
Chalk	7 (2010)	$-36.49 \pm 0.05$	3	$-41.56 \pm 0.02$	3
Chalk	8 (2010)	$-36.89 \pm 0.08$	3	-41.30 ± 0.02	1
Chalk	9 (2010)	$-33.15 \pm 0.25$	3	-39.09	1
Chalk	10 (2010)		3		1
	, ,	$-32.18 \pm 0.01$	3	-36.83	
Chalk	27 (2011)	$-33.41 \pm 0.62$		-41.77	1
Chalk	28 (2011)	-30.11 to -29.99	2	-39.56	1
Chalk	37 (2011)	$-33.61 \pm 0.21$	3	$-43.32 \pm 0.34$	3
Chalk	38 (2011)	$-34.09 \pm 2.31$	3	$-40.09 \pm 0.20$	3
Chalk	39 (2011)	$-32.99 \pm 0.85$	4	-38.59 to -36.39	2
Chalk	40 (2011)	$-32.32 \pm 0.41$	6	$-36.16 \pm 0.45$	5
Chalk	41 (2011)	$-28.64 \pm 1.06$	6	$-36.81 \pm 0.45$	5
Chalk	42 (2011)	$-31.29 \pm 0.72$	6	-32.55	1
Chalk	43 (2011)	$-38.23 \pm 1.30$	6	$-42.30 \pm 2.49$	3
Chalk	44 (2011)	$-34.37 \pm 0.76$	5	$-38.34 \pm 0.23$	3
Chalk	45 (2011)	$-33.20 \pm 0.54$	5	$-41.39 \pm 0.51$	6
Chalk	46 (2011)	$-32.58 \pm 0.39$	6	$-36.75 \pm 0.17$	3
Chalk	47 (2011)	$-29.25 \pm 1.35$	6	$-39.19 \pm 0.62$	6
Chalk	48 (2014)	$-34.92 \pm 0.96$	3	$-44.30 \pm 0.51$	3
Chalk	49 (2014)	$-31.13 \pm 0.39$	3	$-42.56 \pm 0.10$	3
Chalk	50 (2014)	-31.39 to -30.52	2	$-43.72 \pm 0.18$	3
Chalk	52 (2014)	$-33.93 \pm 0.11$	3	-43.16 to -41.97	2
Chalk	53 (2014)	$-33.53 \pm 0.92$	3	$-40.38 \pm 0.17$	3
Chalk	54 (2014)	$-27.79 \pm 0.82$	3	$-39.87 \pm 0.57$	3
Chalk	55 (2014)	$-34.25 \pm 0.30$	3	$-38.33 \pm 0.37$	3
Chalk	58 (2014)	-29.47	1	$-40.85 \pm 0.09$	3
Clay	51 (2014)	-34.90 to -34.36	2	$-38.46 \pm 0.60$	3
Clay	57 (2014)	$-33.05 \pm 0.24$	3	-41.6 to -40.7	2
Coal	17 (2011)	-31.32	1	$-28.88 \pm 0.23$	4
Limestone	11 (2011)	$-27.56 \pm 0.09$	3	$-37.84 \pm 0.14$	3
Limestone	12 (2011)	$-29.35 \pm 0.30$	3	$-38.08 \pm 0.17$	3
Limestone	18 (2011)	-39.53	1	$-34.11 \pm 0.05$	3
Limestone	21 (2011)	-35.68	1	$-42.34 \pm 0.28$	3
Limestone	22 (2011)	-31.46	1	$-35.77 \pm 0.13$	3
Limestone	23 (2011)	-30.67	1	$-38.94 \pm 0.11$	3
Limestone	24 (2011)	-35.56	1	$-39.65 \pm 0.42$	3
Limestone	25 (2011)	-32.66	1	$-39.13 \pm 0.49$	3
Limestone	26 (2011)	-29.63	1	$-37.16 \pm 0.11$	3
Mafic Lava	35 (2011)	$-35.04 \pm 1.02$	3	$-41.38 \pm 0.30$	4
Sandstone	13 (2011)	$-29.81 \pm 0.30$	3	$-31.36 \pm 0.55 \dagger$	3
Sandstone	14 (2011)	$-31.57 \pm 0.82$	3	$-31.30 \pm 0.33$ $-31.78 \pm 0.09$	3
Sandstone	15 (2011)	$-33.81 \pm 0.11$	1	$-30.87 \pm 0.39$	3
Sandstone	16 (2011)	-34.07	1	$-38.09 \pm 0.13$	3

Sandstone	19 (2011)	$-22.78 \pm 0.34$	3	-39.8639.85	2
Sandstone	20 (2011)	-29.23	1	$-28.42 \pm 0.54$ †	3
Sandstone	29 (2011)	$-31.97 \pm 1.54$	3	-35.4	1
Sandstone	30 (2011)	$-27.66 \pm 0.36$	3	-40.13	1
Sandstone	31 (2011)	-29.2228.66	2	$-37.44 \pm 0.25$	3
Sandstone	32 (2011)	$-27.41 \pm 1.09$	3	-33.11	1
Sandstone	33 (2011)	$-30.98 \pm 0.17$	3	$-34.51 \pm 0.14$	3
Sandstone	34 (2011)	$-28.30 \pm 0.44$	3	$-27.01 \pm 0.17$	3
Sandstone	36 (2011)	$-30.48 \pm 0.47$	3	-37.56	1
Sandstone	56 (2014)	$-28.18 \pm 0.22$	3	$-35.39 \pm 0.61$	3

## **Supporting Information**

Table S1. Stream number, site names, latitude and longitude.

No				No			
	Stream site	Latitude	Longitude		Stream site	Latitude	Longitude
1	Chess (upper)	51.678619	-0.531497	30	Stockleigh	50.828669	-3.596573
2	Cray	51.432946	0.137360		Lowman	50.913828	-3.465383
			0.22.20		Crowborough		
3	Darent (upper)	51.276710	0.104090	32	Warren	51.058296	0.122215
4	Itchen	51.086330	-1.203529	33	Hodgehow Wood	54.385572	-2.921985
5	Lambourn at Bagnor	51.439692	-1.386321	34	Belle Grange Beck	54.388649	-2.947094
6	Test	51.130964	-1.485299	35	Carrock Beck	54.709711	-3.031610
7	Till	51.146125	-1.899308	36	Ghyll Head	54.314272	-2.916455
8	Ver	51.783835	-0.378468	37	Belleau	53.289143	0.099984
	Wye at High						
9	Wycombe	51.626327	-0.741859		Welton le Wold	53.374073	-0.083635
10	Wylye	51.118594	-1.883152	39	Allen	50.840853	-2.007536
11	Nant Onneu	51.844590	-3.153215	40	Bourne	51.070008	-1.778369
					Chess at		
12	Nant Wenllan	51.841701	-3.137769		Chorleywood	51.666700	-0.498589
13	Nant Cleisfer	51.846579	-3.243810	42	Darent (lower)	51.350933	0.188828
14	Craig Cerrig Gleisiad Lower	51.889191	-3.496174	43	Frome	50.777333	-2.576144
14	Craig Cerrig Gleisiad	31.009191	-3.4901/4	43	riome	30.777333	-2.3/0144
15	Middle	51.887046	-3.502176	44	Granta	52.113786	0.254714
10	Craig Cerrig Gleisiad	31.007010	3.302170		Lambourn at	32.113700	0.25 1711
16	Upper	51.887423	-3.500837	45	Boxford	51.446572	-1.383592
17	Dare	51.714891	-3.472319	46	Meon	50.949369	-1.136356
18	Upper Tawe	51.826556	-3.683616	47	Test at Wherwell	51.162231	-1.450025
19	Traeth Mawr	51.923453	-3.508511	48	Pillhill	51.196944	-1.538667
20	Owl's Grove	51.836834	-3.382922	49	Andover	51.219711	-1.487791
21	Manifold	53.058900	-1.800589	50	Anton	51.224461	-1.484427
22	Milldale	53.092537	-1.795177	51	Brain	51.801374	0.634004
23	Dove	53.059931	-1.783559	52	Shep	52.091401	0.049623
24	Wye Upper	53.253473	-1.848458	53	Mimram (lower)	51.804682	-0.146922
25	Wye Middle	53.239425	-1.748029	54	Sheringham	51.224461	-1.484427
26	Wye Lower	53.243428	-1.745233		Great Stour	51.243697	0.977561
27	Tadnoll	50.687380	-2.368156	56	Telford	52.685119	-2.541128
28	Bere	50.736946	-2.203922	57	Ter	51.766550	0.591073
29	Creedy	50.814149	-3.649864	58	Mimram (upper)	51.877870	-0.288111

**Table S2.** Mean  $\delta^{13}C \pm 1SE$  for other primary resources sampled from stream sites. The number of samples contributing to each mean value is indicated in brackets.

Site No.	Terrestrial detritus	Moss spp.	Ranunculu s spp.	Berula sp.	Cladophor a sp.	Apium sp.	Callitriche sp.	Myriophyll um sp.	Filamentou s algae
1	-29.65 ± 1.09 (3)	-	-34.73 ± 0.06 (3)	-	-	-	-	-	-
2	-29.52 ± 0.38 (3)	-	-29.52 ± 0.25 (3)	-	-	-	-	-	-
3	-28.10 ± 022 (3)	-	-36.62 ± 0.08 (3)	-	-	-	-	-	-
4	-29.96 ± 0.06 (3)	-	-33.97 ± 0.20 (3)	-32.14 ± 0.03 (3)	-36.74 ± 0.09 (3)	-	-	-	-
5	-30.46 ± 0.73 (3)	-	-33.75 ± 0.18 (3)	-	-	-	-	-	-
6	-28.59 ± 0.13 (3)	-	-35.24 ± 0.20 (3)	-	-	-	-	-	-
7	-30.20 ± 0.26 (3)	-	-34.15 ± 0.15 (3)	-	-	-	-	-	-
8	-30.49 ± 0.42 (3)	-	-33.52 ± 0.18 (3)	-	-	-	-	-	-
9	-30.56 ± 0.22 (3)	-	-33.35 ± 0.37 (3)	-	-	-	-	-	-
10	-29.76 ± 0.95 (3)	-	-33.11 ± 0.01 (3)	-	-	-	-	-	-
11	-30.72 ± 0.07 (3)	-31.97 ± 0.81 (3)	-	-	-	-	-	-	-
12	-30.51 ± 0.03 (3)	-29.55 ± 0.69 (3)	-	-	-	-	-	-	-
13	-29.32 ± 0.02 (3)	-32.57 ± 1.42 (3)	-	-	-	-	-	-	-
14	-29.11 ± 0.06 (3)	-	-	-	-	-	-	-	-
15	-27.80 ± 0.13 (3)	-	-	-	-	-	-	-	-
16	-27.19 (1)	-35.94 (1)	-	-	-	-	-	-	-
17	-29.45 ± 0.19 (6)	-	-	-	-	-	-	-	-
18	-30.57 ± 0.20 (3)	-	-	-	-	-	-	-	-
19	-29.84 ± 0.26 (3)	-	-	-	-	-	-	-	-
20	-29.88 ± 0.48 (3)	-	-	-	-	-	-	-	-
21	-29.77 ± 1.14 (3)	-	-	-	-	-	-	-	-
22	-28.77 ± 0.99 (3)	-	-	-	-	-	-	-	-
23	-29.70 ± 0.52 (3)	-	-	-	-	-	-	-	-
24	-28.14 ± 0.46 (3)	-	-	-	-	-	-	-	-
25	-30.99 ± 0.15 (3)	-	-	-	-	-	-	-	-
26	-30.52 ± 0.38	-	-	-	-	-	-	-	-
27	-29.00(1)	-33.95 (1)	-37.81 ± 0.43 (3)	-	-	-	-	-	-
28	-	-	-28.88 ± 0.79 (3)	-	-	-	-	-	-
29	-28.16 28.09 (2)	-	-	-	-	-	-	-	-
30	-29.67 28.87 (2)	-	-	-	-	-	-	-	-

Site No.	Terrestrial detritus	Moss spp.	Ranunculu s spp.	Berula sp.	Cladophor a sp.	Apium sp.	Callitriche sp.	Myriophyll um sp.	Filamentou s algae
31	-29.31 (1)	-	-	-	-	-	-	-	-
32	-29.28 ± 0.71 (3)	-30.12 29.92 (2)	-	-	-	-	-	-	-
33	-29.44 (1)	-	-	-	-	-	-	-	-
34	31.29 29.81 (2)	-	-	-	-	-	-	-	-
35	-	-41.97 (1)	-	-	-	-	-	-	-
36	-	-	-	-	-	-	-	-	-
37	-	-	-	-	-	-	-	-	-
38	-	-	-	-	-	-	-	-	-
39	-29.67 29.59 (2)	-43.40 ± 0.80 (3)	-30.73 ± 0.75 (6)	-	-	-33.66 ± 0.97 (4)	-32.74 32.34 (2)	-	-36.20 (1)
40	-29.19 ± 0.32 (4)	-	-	-	-	-	-	-	-
41	-28.93 ± 0.25 (6)	-	-30.68 ± 0.30 (5)	-	-	-	-	-	-31.78 ± 0.61 (3)
42	-28.58 ± 0.19 (6)	-39.56 ± 1.62 (3)	-24.18 ± 0.44 (6)	-	-	-	-29.27 ± 0.53 (6)	-26.16 ± 0.51 (3)	-31.14 ± 1.36 (6)
43	-30.40 30.17 (2)	$-44.50 \pm 0.80 (3)$	-33.51 ± 2.90 (3)	-	-	-	-	-	-40.10 38.70 (2)
44	-27.52 ± 0.28 (6)	-	-27.97 ± 0.55 (3)	-	-	-30.21 (1)	-	-	-
45	-28.59 ± 0.29 (6)	-	-34.70 ± 1.29 (6)	-	-	-	-	-	-38.17 ± 0.46 (4)
46	-29.74 28.44 (2)	-	-25.87 ± 0.35 (3)	-	-	-29.98 ± 0.78 (3)	-	-	-
47	-28.90 ± 0.29 (4)	-43.33 ± 0.76 (3)	-32.20 ± 0.72 (6)	-	-	-	-	-	-43.23 (1)
48	-	-	-41.81 ± 0.33 (3)	-	-	-	-	-	-
49	-	-	-	-33.66 ± 0.35 (3)	-	-	-	-	-
50	-	-	-	-36.78 ± 2.76 (3)	-	-	-	-	-
51	-	-	-	-	-	-	-	-	-
52	-	-	-42.47 (1)	-	-	-	-	-	-
53	-	-	-35.10 ± 0.20 (3)	-	-	-	-	-	-
54	-	-	-	-	-	-	-	-	-
55	-	-		-	-	-	-	-	-
56	-	-	-	-	-	-	-	-	-
57	-	-	-	-	-	-	-	-	-
58	-	-	-	-	-	-	-	-	-

**Figure S1 and Table S3.** Stable isotope carbon values for dissolved inorganic carbon ( $\delta^{13}$ C-DIC vs VPDB ‰) in river water, epilithon, *Agapetus* (overwhelmingly *Agapetus fuscipes* but including two samples of the confamililial *Glossosoma conformis*) and 'hypothetical' biofilms based on average carbon fractionation, relative to the DIC, during autotrophy coupled to ammonia oxidation [20‰ (Sakata *et al.*, 2008)) or sulphur oxidation [25‰, on average (Ruby, Jannasch & Deuser, 1987). Data for  $\delta^{13}$ C-DIC were collected from only 28 of the 58 sites used in the main study and were analysed as described in Trimmer *et al.*(2015).

**Table S3.** All Pairwise Multiple Comparison Procedures (Tukey Test) for the  $\delta^{13}$ C values in above:

638	Comparison	Diff of Ranks	$\mathbf{q}$	$\boldsymbol{P}$	<i>P</i> <0.050
639	DIC vs Sulphur at -25	2628.000	12.245	< 0.001	Yes
640	DIC vs Agapetus	2426.000	11.304	< 0.001	Yes
641	DIC vs Ammonia at -20	1498.000	6.980	< 0.001	Yes
642	DIC vs Epilithon	1288.000	6.001	< 0.001	Yes
643	Epilithon vs Sulphur at -25	1340.000	6.244	< 0.001	Yes
644	Epilithon vs Agapetus	1138.000	5.302	0.002	Yes
645	Epilithon vs Ammonia at -20	210.000	0.978	0.958	No
646	Ammonia at -20 vs Sulphur at -2	25 1130.000	5.265	0.002	Yes
647	Ammonia at -20 vs Agapetus	928.000	4.324	0.019	Yes
648	Agapetus vs Sulphur at -25	202.000	0.941	0.964	No
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## Supplementary discussion in relation to the data presented in Figure S1 and Table S3.

We know that there is a high ammonia oxidation potential in both chalk-gravel and sand riverbed sediments (Lansdown *et al.*, 2016) that will be adding chemolithoautotrophic carbon to the biofilm and that could have a  $\delta^{13}$ C value (Sakata *et al.*, 2008) practically the same as our bulk-epilithon (*see* rank Anova in Table S3). The glosssomatids are still deplete relative to both (P=0.002 and P=0.019, respectively), however. If the biofilm was predominantly composed of carbon fixed by sulphur oxidising bacteria, fractionating on average by -25‰, relative to the DIC (Ruby, Jannasch & Deuser, 1987), then that could account for the typical

δ<sup>13</sup>C values in our samples of caddis. If this was the case, then we would expect the products of sulphur cycling to be a significant component of the biofilm and to see evidence, for example, of sulphate reduction as a source of reduced sulphur that could then be reoxidised through chemolithoautotrophy to fix <sup>13</sup>C deplete carbon. However, our previous work in the chalk shows sulphate to be conservative in the riverbed (Pretty, Hildrew & Trimmer, 2006). As sulphur is one of the recognised macro-elements (H, O, C, N, P, S – by average rank abundance) in living biomass (Sterner & Elser, 2003), we would expect some reduced sulphur to result during decomposition but, in these predominantly oxic riverbeds, we would not expect it to accumulate sufficiently for it to dominate the biofilm as, say, in a mat of *Beggiatoa* spp. or *Thioploca* spp. where such chemolithoautotrophy is significant (Thamdrup & Canfield, 1996; Hogslund *et al.*, 2009).

#### **Supplementary References**

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