Instream C sources for primary consumers in two temperate, oligotrophic rivers: possible evidence of bryophytes as a food source

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Abstract. Aquatic food webs use C derived from 2 sources: autochthonous (derived from inside the water body) and allochthonous (derived from outside the water body). Various autochthonous sources are available to consumers (e.g., algae, mosses, macrophytes), and stable isotope analysis can be used to differentiate among C sources and to determine which sources sustain aquatic communities. The goal of our study was to determine which C sources contribute to the autochthonous pathways in 2 oligotrophic rivers, the Upper Salmon and Big Salmon, in New Brunswick (Canada). Samples were taken from headwaters to river mouth (above head of tide) at 17 study sites distributed across subbasins and stream orders. Vegetation samples included macrophytes, filamentous algae, biofilm, and aquatic bryophytes. Macroinvertebrate taxa in the scraper (grazer) functional feeding group were collected to represent primary consumers that feed primarily on autochthonous food sources. Macrophyte and cyanobacteria δ¹³C values were correlated with scraper δ¹³C values (r = 0.55 and 0.47, respectively), but δ¹³C of most scrapers was more depleted than δ¹³C of these food sources (>40% and >90%, respectively). Trophic fractionation is primarily an enrichment process, so these results indicated that macrophytes and cyanobacteria were not important food sources for primary consumers. Filamentous algae and biofilm δ¹³C values were poorly correlated with scraper δ¹³C values (r = 0.36 and 0.40, respectively), and >60% of the values were too depleted for these sources to be important food sources for scrapers. Ninety-eight percent of scraper δ¹³C values were enriched relative to bryophyte δ¹³C values, and bryophyte δ¹³C values were correlated with scraper δ¹³C values (r = 0.53), particularly when samples from slow-flowing habitats were removed from the analysis (r = 0.76). Spatial distributions of 2 important bryophyte taxa differed. Fontinalis sp. was abundant in headwater streams, and Drepanocladus sp. was abundant in low-order streams. Scrapers in low-order streams seemed to depend more on Fontinalis sp. than scrapers in high-order streams depended on Drepanocladus sp. Our results suggest that in low-productivity, nutrient-limited rivers, reduced availability of more preferred food sources (e.g., epilithic algae) might cause primary consumers to switch to marginal food sources, such as bryophytes. Further research is required to confirm this hypothesis and to eliminate confounding factors.

Key words: autochthonous, food webs, stable isotopes, carbon, grazer, macroinvertebrate, bryophyte, aquatic moss, New Brunswick.
phic and heterotrophic bacteria, and fungi. Not all autochthonous sources derive their resources from the aquatic environment; e.g., emergent macrophytes can convert atmospheric CO$_2$ into plant tissue during photosynthesis (Madsen and Sand-Jensen 1991).

The relative importance of allochthonous and autochthonous C sources to aquatic consumers can be difficult to determine (Chapman 1966). Traditionally, consumer diets were studied with methods, such as gut content analysis (Petelle et al. 1979, Rounick and Winterbourne 1986), that provided the researcher a point-in-time reference of ingested material. Within the last 30 y, a more integrative approach, stable isotope analysis (SIA), has become increasingly popular (Fry 2006). SIA reflects only materials incorporated into tissues, so it provides a better indication of long-term nutrient assimilation than does gut content analysis (Rounick and Winterbourn 1986). Inferences about a consumer’s diet are made by comparing its isotopic ratios ($\delta^{13}C$) with those of other species in its food web (Fry and Sherr 1984, Peterson and Fry 1987). This comparison is possible because trophic transfer causes predictable changes, or fractionations, in isotopic signatures of consumers relative to their diet (DeNiro and Epstein 1978, 1981).

$\delta^{13}C$ ratios for primary producers vary because of source C availability and differing methods of photosynthetic processing (Rounick and Winterbourn 1986). Typically, values for terrestrial C$_3$ plants fall within a narrow range around $-28\%$ (Rau 1980, Rounick et al. 1982, France 1995a, Finlay 2001), whereas values for aquatic vegetation can vary from $-50$ to $-10\%$ (Rounick and Winterbourn 1986, France 1995a). Although $\delta^{13}C$ of primary producers can vary significantly from that of their C source, $\delta^{13}C$ increases $<1\%$ with each consumer in the food chain (Post 2002). Therefore, an organism retains the $\sim \delta^{13}C$ value of its primary C source (DeNiro and Epstein 1978, Doucett et al. 1996). This fact, in conjunction with distinct differences in $\delta^{13}C$ values of potential food sources, makes $\delta^{13}C$ values useful for tracing C flow through an ecosystem (Fry and Sherr 1984).

Mixing models can be used to quantify the relative contribution of various food sources to consumers. However, they require isotopically distinct food sources at the local (site) scale (Phillips 2001). An alternative to the use of mixing models within sites is to compare isotope ratios of consumers and food sources across sites with a correlation or regression approach (Bunn et al. 2003). In our study, SIA was used to determine which of many aquatic primary producers contributed most to the autochthonous foods in food webs in the Upper Salmon and Big Salmon River systems, New Brunswick (NB). $\delta^{13}C$ values of aquatic vegetation types were compared with $\delta^{13}C$ values of primary consumers in the scraper (grazer) functional feeding group (FFG). Scrapers primarily consume autochthonous food sources. Allochthonous C sources can be important in food webs of temperate river systems, particularly in headwaters (Vannote et al. 1980), but were not included because our focus was the role of aquatic primary producers in the autochthonous pathway (i.e., grazing invertebrates). Epilithic algae are thought to be the most important autochthonous food source in running waters (Merritt and Cummins 1984, Allan 1995, Finlay 2001), and we predicted that biofilm would be the primary autochthonous food in local food webs. Sampling was done over a large spatial scale to determine if relationships between primary producers and consumers varied with stream order.

**Methods**

**Study locations**

The 2 rivers studied were the Big Salmon River (BSR; Fig. 1A) and the Upper Salmon River (USR; Fig. 1B). Study sites were selected along the lengths of the rivers from headwaters to river mouth to reflect the stream orders present in the 2 systems. The USR runs through Fundy National Park and empties into the Bay of Fundy near the village of Alma, NB, and drains 183 km$^2$. The lower 49% (89 km$^2$) of the basin lies within park boundaries and is classified as a 4th order stream in the North American Datum (NAD83; scale = 1:50,000; NGS 1983). Its annual hydrograph shows a snowmelt-driven high-water peak between mid-March and mid-May, and a smaller peak caused by autumn rains from October to December. The minimum and maximum recorded discharges be-
tween 1967 and 1979 were 0.221 m$^3$/s and 154 m$^3$/s, respectively. During the same period, mean annual discharge was 7.05 m$^3$/s (Environment Canada 2007).

The BSR is ∼20 km west of Fundy National Park and drains 332 km$^2$. The BSR is a 5th-order stream at its mouth (NAD83; scale = 1:50,000). Its annual hydrograph is similar to that of the USR. Mean annual discharge between 1970 and 1979 was 11.78 m$^3$/s. Minimum and maximum discharges were 0.62 m$^3$/s and 140 m$^3$/s, respectively (Environment Canada 2007).

Most land in the 2 basins is forested with only a few small residential developments (<2 people/km$^2$; Cunjak and Newbury 2005). Both catchments are in the Fundy Coast section of the Acadian Forest Region (Rowe 1972). Forest vegetation is mixed hardwood and softwood, and the characteristic tree species are red spruce (Picea rubens), balsam fir (Abies balsamea), sugar maple (Acer saccharum), and yellow birch (Betula alleghaniensis) (Woodley 1985).

Conductivity, a relative measure of enrichment and primary productivity (Biggs 1996), is low in the USR and BSR (range: 22.5–62.7 μS/cm), and N:P ratios range from 20:1 to 60:1. Total N typically is <0.4 mg/L, a value that suggests P limitation in these systems. pH ranges from 6.8 to 7.6 (SMM-H, unpublished data). Based on water chemistry and nutrient levels, these 2 systems can be described as low-productivity, P-limited, neutral-pH rivers.

**Collections**

Samples were collected in July and November 2004 to represent times with high quantities of either autochthonous (summer) or allochthonous (autumn) C sources available. At each study site, duplicate collections were made in both fast- and slow-flow areas (typically, distinct riffle and pool habitats) to ensure collection of all possible food sources. Autochthonous sources, collected from the stream bottom, consisted of aquatic macrophytes, aquatic bryophytes (mosses and liverworts), filamentous algae, and biofilm (scraped from 4–6 randomly selected rocks to remove the matrix of algae, diatoms, bacteria, and fungi). Samples of different species of aquatic vegetation were collected from each study site and rinsed to remove any attached organisms and fine debris. Vegetation samples were not acidified with 10% HCl because no carbonates were detected (Bunn et al. 1995), and low alkalinity and hardness values were recorded throughout the river systems (SMM-H, unpublished data). The amount of material collected for each autochthonous source was biased by the natural occurrence of sources in various habitats. For example, macrophytes occurred predominantly in slow-moving water; therefore, >80% of these specimens were taken from pools, whereas biofilm was collected only from riffles because sediment might have contaminated samples from pools.

Aquatic macroinvertebrates were collected with a kick net (D-net; mesh size = 250 μm). Four samples were collected from each site, starting at the most downstream habitat to ensure that subsequent sampling locations were not disturbed. Samples were sorted to separate invertebrates from their food sources. Invertebrates were stored at cool temperatures for 24 h to allow gut contents to clear (DeNiro and Epstein 1978). Mollusk shells were removed to avoid contamination from nondietary carbonates (Bunn et al. 1995). Invertebrates were frozen prior to identification and SIA preparation.

**Analysis**

Macrophytes, bryophytes (mosses and liverworts), and algae were identified to genus (Prescott 1980, Sears 1998). All vegetation was diligently hand-picked to remove any invertebrates, detritus, and algae growing on, or entwined within, the various samples to ensure that a pure sample was analyzed. SIA of epiphytic algae removed from macrophyte and bryophyte samples was not possible because of insufficient amounts of algae.

Two groups of bryophytes, mosses and liverworts, were collected and analyzed for stable isotopes. The liverwort species (Chiloscyphus sp.) was isotopically distinct from both moss species (Fontinalis sp. and Drepanoclados sp.). Chiloscyphus sp. had an average δ$^{13}$C of −33%, and >80% of the δ$^{13}$C values were between −28% and −35%. No grazers had δ$^{13}$C values within this range, a result that suggested Chiloscyphus sp. was not important in the food web. Therefore, this species was not included in subsequent analysis or interpretation.

Attempts were made to separate detritus from algae in the biofilm matrix with density fractionation in colloidal silica (Hamilton et al. 2005). Preliminary results were inconclusive, and additional analysis was required to refine the method and assess the efficacy of the separations. Additional experimentation was not possible because of budget and time constraints, so biofilm samples were sorted by hand under a dissecting microscope to remove detritus.

Invertebrates were identified to family with keys in Merritt and Cummins (1984) and were prepared for SIA separately. Whenever possible, individual invertebrates were analyzed. When body size was too small for analysis, several individuals were pooled for
SIA. FFG classification (Merritt and Cummins 1984) was used when comparing invertebrate consumers to potential food sources. Scrapers (grazers) feed primarily on primary producers and were used to identify important autochthonous sources. Invertebrate taxa classified as obligate scrapers in our study were Blephariceridae (Diptera), Gastropoda (Mollusca), Glossosomatidae (Trichoptera), and Psephenidae (Coleoptera). Heptageniidae (Ephemeroptera) were classified as facultative scrapers (Merritt and Cummins 1984). Preliminary SIA of heptageniid mayflies yielded δ^{13}C values comparable with those of scrapers, so heptageniids were used with obligate scrapers to assess important autochthonous food sources.

All organic tissue samples were dried at 60°C for 48 h, ground into a fine homogenate with a mortar and pestle, and submitted for C and N SIA at the Stable Isotopes in Nature Laboratory, University of New Brunswick, Fredericton, NB. All samples were combusted and analyzed in a Finnigan Mat Delta Plus continuous-flow, isotope-ratio mass spectrometer (ISOMASS Scientific Inc., Calgary, Alberta) connected to a Thermoquest NC2500 elemental analyzer (ISOMASS Scientific Inc.).

Stable isotope ratios are expressed as δ values and are measured as parts per thousand (‰) differences relative to international standards:

\[ \delta X = \left( \frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \right) \times 1000 \]

where X is 13C, and R is the corresponding ratio 13C/12C. The standard used for 13C was PeeDee Belemnite carbonate (Craig 1957), which is arbitrarily set at 0‰. A greater δ value indicates enrichment of the heavier isotope component, whereas a smaller δ value indicates depletion of the heavier isotope.

Analytical error was calculated per Jardine and Cunjak (2005). The commercially available organic analytical standard (Acetanilide; Elemental Microanalysis Ltd., Devon, UK) had a mean (±1 SD) δ^{13}C of −33.62 ± 0.25‰, whereas the internal standard, bovine liver standard, had a δ^{13}C value of −18.70 ± 0.19‰. Within an analytical run, sample repeats never had 1 SD >0.77‰, and 70% of sample repeats had 1 SD <0.20‰. Invertebrate δ^{13}C was lipid-corrected using the lipid normalization model presented in Logan et al. (2008) that uses C:N as a proxy for lipid content.

Three measures were used to help identify important primary producers: 1) the 1:1 line relating δ^{13}C values of primary producers vs consumers, 2) the coefficient of determination \( (r^2) \), and 3) the correlation coefficient \( r \) (Bunn et al. 2003). δ^{13}C increases, on average, by 0.4‰ with each trophic transfer (DeNiro and Epstein 1978, Vander Zanden and Rasmussen 2001, McCutchan et al. 2003, Post 2002). Therefore, primary producers that are directly consumed should have data points that fall slightly above the 1:1 line in a consumer vs producer plot. We used \( r^2 \) and \( r \) as measures of the strength and direction of associations between primary producers and consumers. Mean δ^{13}C of primary producers were plotted against mean values of various primary consumers from respective habitats, sites, and seasons (SigmaPlot 10.0; Systat Software Inc., San Jose, California). Regression and correlation analyses were done for each vegetation type (NCSS 2004; NCSS, Kaysville, Utah). Variation in primary consumer δ^{13}C values should track the spatial and temporal variation in the vegetation δ^{13}C values if the vegetation is an important C source (across all sites and times; Bunn et al. 2003).

**Results**

Five primary producer taxa were identified as potential autochthonous food sources for consumers in the study rivers: biofilm, macrophytes, filamentous algae (red and green), and aquatic bryophytes. δ^{13}C of biofilm was poorly correlated with δ^{13}C of scrapers \( (r = 0.40, r^2 = 0.16) \), and >60% of the data points fell below the 1:1 line (i.e., most scrapers were more depleted in 13C than were the biofilm samples; Fig. 2A). δ^{13}C of macrophytes was correlated with δ^{13}C of scrapers \( (r = 0.55, r^2 = 0.30) \), and >40% of the data points fell below the 1:1 line (Fig. 2B). δ^{13}C of cyanobacteria was correlated with δ^{13}C of scrapers \( (r = 0.47, r^2 = 0.22) \), and >90% of the data points fell below the 1:1 line (Fig. 2C). δ^{13}C of filamentous algae was poorly correlated with δ^{13}C of scrapers \( (r = 0.36, r^2 = 0.13) \), and >60% of the data points fell below the 1:1 line (Fig. 2D). δ^{13}C of mosses was correlated with δ^{13}C of scrapers \( (r = 0.53, r^2 = 0.28) \), and 98% of the data points fell on or above the 1:1 line (Fig. 3A). Thus, most scrapers were slightly enriched in 13C compared with this potential food source.

Bryophytes in these rivers generally occur in riffles (i.e., fast-flow areas) rather than in pools. The pool samples were collected from transitional habitats (i.e., runs, tails of pools). When pool samples were removed from the analysis, the strength of the correlation between δ^{13}C of mosses and δ^{13}C of scrapers increased \( (r = 0.76, r^2 = 0.57; \text{Fig. 3B}) \). When pool samples were removed from the analyses for the other vegetation types, the strength of the correlations decreased or increased slightly (biofilm: not collected from pools; cyanobacteria: \( r = 0.15, r^2 = 0.02 \); algae: \( r = 0.39, r^2 = 0.15 \), with the exception of macrophytes.
However, the sample size for macrophytes was very small ($n = 7$), and all data points fell below the 1:1 line.

The relationship between invertebrate and moss $\delta^{13}C$ signatures did not differ between obligate scrapers and heptageniids (no apparent separation of invertebrate groups in Fig. 4A). This result indicates that the ranges of $\delta^{13}C$ values of these invertebrate groups are similar throughout the 2 rivers. However, spatial distributions of bryophyte taxa varied. Both moss species, *Fontinalis* sp. and *Drepanocladus* sp., occurred in all stream orders, but *Fontinalis* sp. was more abundant in low- than in high-order streams and *Drepanocladus* sp. was more abundant in high- than in low-order streams. $\delta^{13}C$ values of the 2 moss species differed, reflecting their spatial distributions (*Fontinalis* sp.: $\delta^{13}C = -38$ to $-33\%o$; *Drepanocladus* sp.: $\delta^{13}C = -30$ to $-25\%o$; Fig. 4B). The correlation and 1:1 line analyses (Fig. 4B) indicated that invertebrate $\delta^{13}C$ in low-order streams was better correlated with $\delta^{13}C$ of *Fontinalis* sp. ($r = 0.73$) than invertebrate $\delta^{13}C$ in high-order streams was correlated with $\delta^{13}C$ of *Drepanocladus* sp. ($r = 0.63$).

**Discussion**

**Biofilm**

We found little evidence to support our prediction that biofilm was the most important primary producer in the autochthonous pathway in these rivers. $\delta^{13}C$ results indicate that most scrapers were more depleted in $^{13}C$ than were biofilm samples, a result suggesting that scrapers were not feeding primarily on biofilm. Moreover, very little biofilm grows on the stream beds of these rivers because of low productivity and nutrient limitation, especially in headwaters where closed canopies also reduce insolation. Grazing is an unlikely explanation for the scarcity of biofilm. Invertebrate abundance was not measured, but grazers constituted a very small proportion of individuals collected during our study (SMM-H, personal observation).

Biofilm assemblages consist of attached algae and detritus and includes microfloral and microfaunal elements (Wetzel 1975). Diatoms are the critical nutritional component of the biofilm matrix in streams (Allan 1995) and are consumed primarily by scrapers (Merritt and Cummins 1984, Finlay 2001). However,
they often constitute the smallest fraction of biofilm on stream rocks (Madsen 1972, Cummins 1975). Particularly in headwater streams, detritus and other organic matter (e.g., bacteria, fungi) constitute most of the matrix, and isolation of a pure sample of epilithic microalgae is extremely difficult (Finlay 2001). Diatoms, bacteria, and fungi can have different δ^{13}C values, and the signatures of components, such as detritus and heterotrophic bacteria, can mask the true δ^{13}C signature of microalgae in the biofilm matrix. Selective consumption or selective assimilation of ≥1 biofilm component could cause consumer δ^{13}C values to differ from the δ^{13}C value of the entire biofilm assemblage (Rounick et al. 1982). Therefore, despite its scarcity, biofilm cannot be eliminated completely as a possible C source, and it might be consumed in conjunction with another C source in these rivers.

**Macrophytes**

More than 40% of scrapers had δ^{13}C values more depleted than macrophyte δ^{15}C values. Macrophyte
abundance and diversity were low at most study sites (8 sites had 0 species, 6 sites had ≤3 species, 3 sites had 4–6 species). These results indicate that macrophytes were not an important direct source of C for scrapers in these rivers. Vascular macrophytes serve an important indirect role in aquatic food webs, but their use as a food source has long been an area of debate (Shelford 1918, Frohne 1956, Lodge 1991, Newman 1991, France 1995b). Hynes (1970) stated that it is rare to find the leaves of higher plants (i.e., macrophytes) damaged by invertebrate grazing in running water and, summarizing Whitehead (1935), that “remarkably few invertebrates feed directly on them.” Newman (1991) presented evidence suggesting that macrophytes were important food sources for aquatic communities but concluded that most herbivory was by rare taxa that were specialized oligophagous herbivores that represented families and orders of primarily terrestrial insect groups.

In an effort to resolve this controversy, France (1995b) compiled δ13C results from the literature for benthic animals, macrophytes, and associated epiphytic algae from 4 different environments in North America (seagrass meadows, estuarine salt marshes, rivers, and lakes) and found little evidence for direct assimilation of macrophyte C in any of these environments. Bunn et al. (1999) found that macrophytes were isotopically distinct from most consumers in Australian rivers and probably contributed little to aquatic food webs in these rivers.

**Cyanobacteria**

Cyanobacteria were 13C enriched compared with scrapers, and scraper and cyanobacterial δ13C were poorly correlated. These results suggest cyanobacteria are not a C source in the study rivers. Many cyanobacteria possess morphological defenses that reduce grazing pressure. Long filaments and large colonies, called flakes, can be difficult for herbivores to ingest (Lampert 1987). Freshwater cyanobacteria have toxic defense compounds (Carmichael 1981) that deter grazing by herbivorous zooplankton or reduce herbivore abundance by interfering with physiological processes (Lampert 1987), and toxic compounds produced by marine cyanobacteria deter feeding by fish and invertebrates (Nagle and Paul 1999, Thacker et al. 2001). Three of the 5 genera found in our study contain toxins that deter herbivory: *Tolypothrix* (Thacker et al. 2001), *Lyngbya* (Yin et al. 1997), and *Nostoc* (Carmichael 1997).

**Filamentous algae**

Filamentous algae also were 13C enriched compared with scrapers, and scraper and algal δ13C were poorly correlated. These results suggest filamentous algae are not a C source in the study rivers. Doucet et al. (1996) found no evidence of direct grazing of the filamentous chlorophyte, *Cladophora* sp., by the primary consumer, *Glossosoma* (Trichoptera), in another NB river system (Miramichi). Like cyanobacteria, green and red filamentous algae have physical properties that reduce herbivory. For example, gelatinous green algae pass through the guts of zooplankton unharmed because they have an indigestible sheath (Porter 1977). Bunn et al. (1999) found that filamentous algae were isotopically distinct from most consumers in Australian rivers and probably contributed little to aquatic food webs in these rivers.

**Bryophytes**

Mosses influence the structure of invertebrate communities by providing habitat for colonization, surfaces for the growth of epiphytic algae and accumulation of detritus, refuge from predators or high current velocities, and material for case construction (Glime and Clemons 1972, Suren 1988, Suren and Winterbourn 1992). However, few invertebrates are thought to consume aquatic bryophytes directly (Bunn et al. 1989, Glime 2006). Our results might suggest otherwise. Most fractionation values associated with trophic transfer (Δδ13C) fell within a range of 0 to 5‰ enrichment. The amount of trophic fractionation can vary between individuals and consumer species, and similar ranges of Δδ13C were reported by DeNiro and Epstein (1978) and Post (2002). However, trophic fractionation is primarily an enrichment process (DeNiro and Epstein 1978) rather than a depletion process. Therefore, our δ13C results for mosses were significant when compared with δ13C of the other primary producers tested.

An increasing number of studies have reported consumption of aquatic bryophytes (Hynes 1970, Gerson 1972, Winterbourn et al. 1986, Suren 1988, Suren and Winterbourn 1991, Glime 2006). For example, crustaceans, gastropods, mites, tardigrades, and many groups of insects consume bryophytes (Gerson 1972, Richardson 1981). In aquaria, lake-dwelling snails consumed the aquatic moss *Fissidens fontanus* but not *Fontinalis* sp. (Lohammer 1954). Fecal analyses indicated that the aquatic isopod *Asellus militaris* regularly consumed *Fontinalis antipyretica* in the wild (Bowden et al. 1999). Suren (1988) compared invertebrate assemblages on real and artificial mosses and found significant decreases in the abundance of certain taxa on artificial mosses, a result indicating that some organisms were dependent on the bryophytes for food. Moss fragments were found in the
guts of 14 of 23 species collected from shaded and unshaded alpine streams in New Zealand, but the frequency of occurrence was low, except for 3 species that had consumed significant amounts of the bryophytes (Suren and Winterbourn 1991).

Bryophytes can contain a number of phenolic compounds, lignin-like compounds, and terpenes that deter herbivory, particularly in leaves, that can influence the part of the plant that is consumed. Harvester ants (Messor sp.; Loria and Herrnstadt 1980), lemmings (Dicrostonyx spp.; Pakarinen and Vitt 1974), and woodland slugs (Arion spp.; Jennings and Barkham 1975) consume terrestrial moss capsules. Terrestrial isopods consumed significantly more leaves than tough stems of 2 Polytrichum species, despite their high phenolic content (Glime 2006).

Aquatic mosses often are regarded as low-quality food. However, Fontinalis antipyretica is not nutritionally inferior to other plants (Bowden et al. 1999, Glime 2006) and should be a prime food source for many aquatic organisms (Glime 1994). Winterbourn et al. (1986) found that the nutritive value of bryophyte tissues did not differ greatly from that of nearby vascular plants, and 5 species of bryophytes tested by Suren and Winterbourn (1991) had at least as much lipid, carbohydrate, and N content as common grass, shrub, and tree foliage, although holocellulose, crude fiber, and ash content were slightly higher. N content often is used as an indicator of food quality and can be assessed with C:N ratios, where a low value indicates good quality as a food source (Jardine et al. 2005).

Bryophytes in our study system have lower C:N ratios (26.5 ± 6.0) than does conditioned leaf litter (32.9 ± 14.1) from the same sites (SMM-H, unpublished data), further indicating that moss is not nutritionally inferior.

Bryophytes might be an important alternate (marginal) food source when a preferred food is unavailable. In terrestrial environments, when food sources are limited or unavailable (e.g., during winter), harvester ants feed on terrestrial moss capsules (Loria and Herrnstadt 1980), and populations of caribou (Rangifer tarandus grantii) and reindeer (Rangifer tarandus tarandus) feed on various mosses (White and Trudell 1980). In the laboratory, the detritivorous caddisfly larvae Zelandopsyche ingens consumed the aquatic moss Fontinalis rigidulus rather than less desirable terrestrial foods, such as yellow beech (Nothofagus spp.) leaves and clean twigs (Winterbourn and Davis 1976). Thus, aquatic macroinvertebrates that do not typically consume aquatic mosses might do so if preferred food sources (e.g., diatoms) are scarce. Abundances of primary producers were not measured in our study rivers, but water chemistry and nutrient data (SMM-H, unpublished data) show that these rivers have low productivity and are P limited. In unproductive systems, bryophytes are better able than algae to sequester nutrients (Bowden et al. 1999). Therefore, we suggest that the availability of a preferred food source might be so limited in our system that primary consumers (i.e., scrapers) use an alternate food source, aquatic bryophytes.

δ13C of Fontinalis sp. was more strongly correlated with scraper δ13C than was δ13C of Drepanocladius sp. Fontinalis spp. are always found submerged (Grout 1903) and cannot survive even a week of air-drying (Watson 1964), whereas Drepanocladius species are not as sensitive to desiccation and can be found at the edges of brooks and growing on emergent rocks in splash zones (Grout 1903). In our system, Fontinalis sp. was more abundant in low-order reaches, where it grew large strands (≥50 cm) and frequently colonized >75% of the substrate surface. Drepanocladius sp. was more abundant in higher-order reaches and grew in shorter tufts (<5 cm). Bryophyte abundance was substantially lower in high- (~5% of substrate surface) than in low-order reaches. Thus, ecological differences might have limited the availability of Drepanocladius sp. to scrapers and reduced its importance to local food webs relative to Fontinalis sp.

Our results suggest that aquatic mosses might be a primary C source for grazing aquatic invertebrates in the BSR and USR. The possibility that a potential macroscopic autochthonous food source was not sampled is unlikely given the spatial and temporal extent of sampling. However, microscopic food sources, such as epiphytes or bacteria growing on the bryophytes, might have δ13C signatures similar to those of bryophytes, and could provide an alternate explanation for the correspondence between scraper and bryophyte isotope ratios observed. Large amounts of periphyton (Johnson 1978, Suren 1988) and detritus (Johnson 1978) commonly are associated with bryophytes. Dissolved organic matter and plant nutrients released from moss cells (Wetzel 1983) might enhance the growth of the surrounding periphyton. Bryophytes use only dissolved CO2 for photosynthesis, whereas algae can use HCO3- when CO2 is scarce (Bowden et al. 1999, Finlay 2004). However, algae use dissolved CO2 preferentially, and CO2 should be available in high concentrations in low-productivity systems. Thus, epiphytes could have δ13C values similar to those of bryophytes in our system (Finlay 2004). Bacterial C is derived from dissolved organic C and particulate detritus and should have a δ13C signature that reflects these sources (Hall 1995). Therefore, bacteria growing on the surface of bryophytes could have δ13C values similar to those of the bryophytes.
Hall (1995) found that bacterial C was an important C source to the food web in a headwater stream and that the scraper Stenonema (Heptageniidae) fed predominantly on bacterial C. Thus, we are unable to completely exclude epiphytes and bacteria as possible food sources.

FFGs other than scrapers also can rely on autochthonous food sources to varying degrees. Collector-gatherers consume particulate organic matter, which is typically a combination of autochthonous and allochthonous C sources and their associated microbes, from the sediment (Vannote et al. 1980). In our study, allochthonous C remained relatively unchanged (SMM-H, unpublished data). Therefore, spatial and temporal changes in collector-gatherer δ13C should reflect changes in autochthonous C (Bunn et al. 2003). Similarly, collector-gatherer δ13C was poorly correlated with macrophyte (r = 0.29, r² = 0.08, 80% of values below the 1:1 line), cyanobacteria (r = 0.55, r² = 0.30, 88% below 1:1), filamentous algae (r = 0.47, r² = 0.22, 54% below 1:1), and biofilm (r = 0.51, r² = 0.26, 52% below 1:1) δ13C. However, bryophyte δ13C was well correlated with collector-gatherer δ13C (r = 0.76, r² = 0.58), and 97% of values were above the 1:1 line, indicating that collector-gatherer δ13C was enriched relative to bryophyte δ13C. Collector-gatherers might have consumed all vegetation types, but the strong correlation with bryophytes strengthens the idea that aquatic mosses are supporting the autochthonous portion of food webs in these unproductive river systems.

In conclusion, many potential aquatic food sources can contribute to food webs throughout a river system. Our results indicate that macrophytes, filamentous algae, and cyanobacteria were not important food sources to invertebrates in the USR and BSR. Biofilm also seemed unimportant in local food webs but could not be excluded as a potential food source. Scrapers appeared to use aquatic bryophytes as their primary autochthonous food source, although the possibility that grazers were consuming isotopically similar epiphytes and bacteria growing on bryophyte surfaces could not be eliminated. However, this possibility does not exclude use of moss δ13C as an autochthonous C source in subsequent mixing models for the USR and BSR rivers. The low productivity and nutrient-limited nature of these river systems might severely reduce the availability of preferred food sources and require that primary consumers switch to an alternate food source. Confirming that aquatic bryophytes are this alternate food source will require additional studies, such as tracer additions or gut content analysis. However, our results suggest that aquatic mosses are important food sources in low-productivity streams.

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