Carbon movement and assimilation by invertebrates in estuarine habitats at a scale of metres

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ABSTRACT: Theories of large-scale (kilometres) movement of carbon within and from estuaries are often not supported by empirical data, and this provided the basis for a smaller-scale (i.e. < 100 m) analysis of carbon movement and assimilation between adjacent habitats. We tested 3 models that potentially explained the movement and assimilation of carbon by resident animals in estuarine habitats at different spatial scales: coarse (>30 m), intermediate (2 to 30 m) and fine (<2 m). The carbon stable isotope signatures of 2 crab and 2 gastropod species were measured at different positions in saltmarsh and mangrove habitats (centre, intermediate and edge in the saltmarsh, intermediate and centre positions in adjacent mangroves) at 5 sites. The δ^{13} C signatures of crabs collected from the saltmarsh (-15.6 ± 0.2%) were significantly more enriched than those of crabs from the mangrove habitat $(-22.1 \pm 0.3 \%)$, but did not differ between positions within each habitat. The δ^{13} C signatures of crabs in the saltmarsh were similar to those of the dominant macrophyte, the salt couch grass Sporobolus virginicus ($-14.9 \pm 0.1\%$). The δ^{13} C signatures of crabs in the mangrove habitat were enriched relative to those of the mangroves $(-27.6 \pm 0.2\%)$, but were similar to those of the microphytobenthos in that habitat $(-24.6 \pm 0.7\%)$. The crabs thus fitted the fine-scale model of assimilation of carbon produced in their immediate vicinity, although the signatures for crabs in the mangrove habitat were also consistent with a food source comprising a mixture of mangroves and a more enriched source, possibly the salt couch grass S. virginicus. Gastropods were found only in the saltmarsh habitat. Their δ^{13} C signatures did not differ among central and intermediate positions (-15.3 \pm 0.2%) but were lower at edge positions (-17.0 \pm 0.1%). The δ^{13} C signatures of gastropods indicated assimilation of carbon from sources 2 to 15 m away, at the lower end of the intermediate scale. The extent of carbon movement and assimilation varies among estuaries, and our results show that in some situations it occurs at scales much smaller than previously realised.

KEY WORDS: Trophic ecology · Stable isotope analysis · Estuaries · Crustacea · Gastropoda

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INTRODUCTION

The source of an animal's food is a central organising theme in ecology (Polis et al. 1997). The movement of potential energy sources across spatial boundaries is influenced by the structural complexity of the land-scape (Holt 2002) and the permeability of habitat boundaries (Marquet et al. 1993), not least in aquatic systems. In aquatic systems the potential connection between seemingly fragmented landscapes is en-

hanced by the role of water as a vector for nutrient, detritus and animal transport (Polis et al. 1997). In marine systems, the study of large-scale movement of carbon produced inshore to offshore environments, termed 'outwelling' (Odum 1979), has provided the foundation for a closer analysis of the scale of carbon movement in marine and estuarine systems.

Early studies of outwelling focused on the role of saltmarshes (predominantly in the USA) in supporting offshore fisheries productivity, with the term subsequently also applied to mangroves. However, empirical evidence for large-scale movement of carbon from intertidal habitats is variable. In some estuaries carbon either does not move very far or moves but is not assimilated by consumers. For example, a study in South Africa demonstrated only minor contributions of carbon from a high marsh to offshore waters (Taylor & Allanson 1995). In Malaysia, mangrove carbon was detected several kilometres offshore but was not assimilated by consumers (Rodelli et al. 1984). In a review of mangrove outwelling, Lee (1995) summarised the role of mangroves to include net exporters, net importers, and sinks that retain large proportions of the litter production for *in situ* consumption.

The lack of evidence of large-scale movement and assimilation of carbon from inshore to offshore in some estuaries (e.g. Dittel et al. 2000) led to studies of the movement of carbon between inshore habitats. Marguillier et al. (1997) examined the trophic links between seagrass and mangroves separated by ~1 km, and found evidence to suggest the movement and assimilation of mangrove carbon to nearby seagrass consumers. The shift in focus from carbon transfer between inshore and offshore to exchange between inshore habitats also led to a reduction in the scale at which carbon movement and assimilation was examined, from several kilometres to ≤1 km. A recent study examining assimilation of carbon by consumers at a creek site and a river site separated by hundreds of metres found that the movement and assimilation of carbon might be more appropriately examined at even smaller scales than had previously been considered (Hsieh et al. 2002).

Stable isotope analysis of carbon is an effective method of tracing energy transfer from autotrophs to consumers (e.g. Deegan & Garritt 1997, Loneragan et al. 1997, Chong et al. 2001). This method is able to distinguish among carbon sources, where autotrophs have distinct ratios of ¹³C/¹²C, and consumers assume the ratio of their food source (Peterson & Fry 1987). The saltmarsh and mangrove habitats used in this study are ideally suited to this approach, as the ¹³C/¹²C ratios of the dominant autotrophs in each habitat are clearly separated. Stable isotope analysis of carbon also has advantages over other techniques used to study food-web processes as the δ^{13} C values of consumers experience only minor enrichment across trophic levels (typically <1‰; McCutchan et al. 2003). This means that despite multiple trophic shifts (for example via a microbial loop or through the consumption of small invertebrates), stable isotope analysis of carbon can trace the original (autotrophic) source of carbon and therefore does not rely on the direct consumption of plant material by the study organism.

In this study, we used stable isotope analysis to test 3 potential models of carbon movement and its assimilation by invertebrates in adjacent saltmarsh and mangrove habitats, where the carbon signatures of autotrophs from each habitat could be distinguished (Fig. 1). The coarse-scale model (Model a) comprises large-scale movement (>30 m) of carbon and therefore no pattern in carbon isotope signatures of consumers across the habitat boundary. The intermediate-scale model (Model b) comprises more limited carbon movement and assimilation across the habitat boundary (2 to 30 m): the prediction of this model is that δ^{13} C signatures of animals will be influenced by their position within a habitat—the closer to an adjacent habitat, the greater the contribution of carbon from the adjacent habitat. The fine-scale model (Model c) predicts that there is no movement and assimilation of carbon across the habitat boundary: in this scenario, an animal obtains its carbon from autotrophs in its immediate vicinity. The coarse model represents the current understanding of estuarine food webs; the intermediate and fine-scale models represent patterns and extents of movement and assimilation not addressed in previous studies. The labels 'coarse', 'intermediate' and 'fine' were chosen to avoid confusion with other terms such as 'local', which is used in the literature to refer to kilometre-scale movement of carbon.

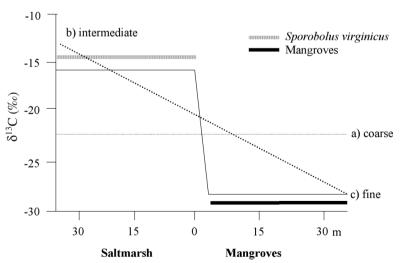


Fig. 1. Patterns of animal isotope signatures predicted by 3 possible models explaining carbon movement and assimilation between adjacent estuarine habitats at different spatial scales: coarse (>30 m: Model a), intermediate (2 to 30 m: Model b), and fine (<2 m: Model c)

MATERIALS AND METHODS

We sampled 5 saltmarshes adjacent to mangroves and dominated by the salt couch grass Sporobolus virginicus, in March 2003 in southern Moreton Bay, Queensland, Australia (Fig. 2). The dominant mangrove species was Avicennia marina, but occasional examples of Ceriops tagal, Rhizophora stylosa and Bruquiera gymnorhiza also occurred. Saltmarshes form part of a complex mosaic of estuarine habitats common in this region, and were defined visually by the presence of discrete habitat boundaries marked by an abrupt change in vegetation type (i.e. mangroves and water). The areas of the saltmarsh habitat were between 1 and 2 ha, and adjacent mangroves were at least 10 ha in extent. The marshes selected were lowlying and were inundated on at least 1 high tide each day for 2 out of 4 wk during the sampling period.

We chose 2 crab species, Australoplax tridentata (Ocypodidae) and Parasesarma erythrodactyla (Grapsidae), and 2 gastropod species, Salinator solida (Pulmonata: Aphibolidae) and Ophicardelus quovi (Pulmonata: Ellobiidae) for analysis. The crabs were chosen because they are found in both saltmarshes and mangroves, and whilst there is little documentation of the movement of A. tridentata and P. erythrodactyla, they are considered to be faithful to their burrows (M. A. Guest pers. obs.) and are therefore assumed to have restricted home ranges. The gastropods were chosen because anecdotal evidence also suggests they are of limited mobility. The crabs are thought to consume detritus, which consists of plant material, microbes and small invertebrates; the snails are either grazers and/or detritivores.

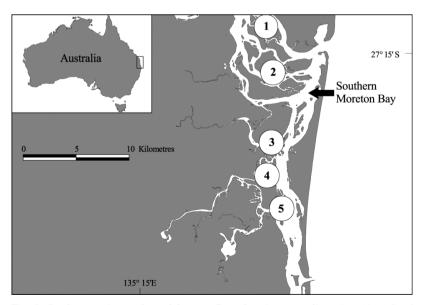


Fig. 2. Study area in southern Moreton Bay showing sampling sites (1 to 5) of saltmarsh adjacent to mangroves

At each of the 5 sites (Fig. 2), 5 positions across the 2 habitat types were chosen: centre and intermediate positions in each habitat, and a position near the saltmarsh/mangrove interface. This last position is labelled 'edge', but was actually 2 m inside the saltmarsh habitat (Fig. 3). Setting the edge position slightly inside the saltmarsh gave us more resolution of carbon movement than having it exactly on the interface between habitats. Positions were separated by about 15 m, so the central positions were about 30 m inside each habitat. We collected 3 samples of both crab species from each position (i.e. 15 samples per site per species), and 3 samples of both gastropod species were collected from central, intermediate and edge positions in the saltmarsh habitat at 3 sites, but no gastropods were collected from mangrove habitats. Each sample consisted of several individuals collected immediately adjacent to a randomly selected point at each position in the habitat. We collected 3 samples of live Sporobolus virginicus from central and edge positions at each site (i.e. 6 samples at each site), and 3 samples of fresh leaves for each mangrove species from intermediate and central positions. Apart from the dominant autotrophs, numerous but inconspicuous algal and cyanobacteria cells are present in the superficial sediment of both habitats; 1 sample of this microphytobenthos (MPB) was collected at the intermediate position in the saltmarsh and in the mangrove habitat at each site, adjacent to where collections of crabs and gastropods were made, by scraping the top 1 cm of sediment from the mud surface.

Gastropod samples were allowed approximately 8 h before freezing for extrusion of gut contents. All other samples were frozen immediately after collection.

Samples were thawed prior to processing. The exoskel-eton and shell were removed from the crabs and gastropods, the gut removed from the crab, and soft flesh selected from both for analysis. MPB samples were washed through 53 µm mesh to remove infauna, and material passing through the mesh was then washed through 5 µm mesh. Material retained on this mesh was added (9 ml) to a centrifuge tube containing 21 ml of colloidal silica (LUDOXTM AM30, density = 1.21) and centrifuged at 10000 rpm for 10 min. A band of diatoms, some organic matter and silica particles formed at the top of the centrifuge tube. This band was removed and again washed through a 5 µm mesh to remove the silica and any remaining microbes. Inspection of

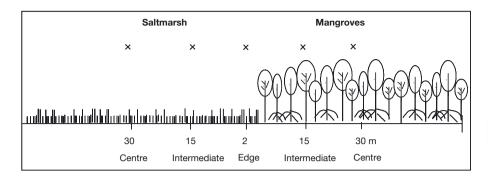


Fig. 3. Sampling transect for each site of saltmarsh adjacent to mangroves. Sampling positions transecting the 2 habitat types are: centre, intermediate, edge (saltmarsh) and intermediate and centre (mangroves). X: Replicates for each position within each habitat (n = 3 for each position at each site)

samples showed that they consisted predominantly of microalgae (mainly diatoms) with occasional contamination by very fine detrital fragments. All samples were dried and ground, placed into tin capsules and their isotopic signatures analysed on an Isoprime mass spectrometer. The ratios of $^{13}\mathrm{C}/^{12}\mathrm{C}$ for all samples were calculated as the relative per mil (%) difference between the sample and the recognised international standard (Pee Dee belemnite limestone carbonate) and expressed as $\delta^{13}\mathrm{C}$ values.

Differences in the isotope signatures of animals at different positions within each habitat and among sites were tested using a 2-way analysis of variance separately for each species. The 2 factors were position (fixed, 5 levels for crabs, 3 for gastropods) and site (random, 5 levels for crabs, 3 for gastropods). Differences among sites were not of specific interest but were included to allow the variation due to positions to be fully partitioned. Data were checked for homogeneity of variances using Cochran's *C*-test, and no transformations were necessary. Where the position factor was significant, differences among positions were tested using a Student-Newman-Keuls (SNK) test.

RESULTS

Autotrophs

The δ^{13} C signatures of *Sporobolus virginicus* varied by <1% among habitat positions (δ^{13} C centre: -14.7 ± 0.1%; edge: -15.0 ± 0.1%). There was also little variation (<2%) in δ^{13} C among mangrove species *Avicennia marina* (-27.4 ± 0.2%), *Ceriops tagal* (-28.2 ± 0.3%), *Rhizophora stylosa* (-26.3 ± 0.2%) and *Bruguiera gymnorhiza* (-28.2 ± 0.5%). The mean δ^{13} C of MPB in the saltmarsh (-23.0 ± 0.5%) was 1.4% higher than in mangroves (-24.4 ± 0.2%). The δ^{13} C of saltmarsh and mangroves did not vary significantly among sites throughout the estuary (overall means shown in Fig. 4).

Consumers

The δ^{13} C signatures of each animal species differed significantly among habitat positions (Fig. 4). Both species of crabs showed the same pattern of δ^{13} C. The δ^{13} C signatures of crabs differed significantly among habitats but not among positions within the same habitat. The $\delta^{13}C$ values were significantly higher for crabs in the saltmarsh (-15.2 \pm 0.2% for Australoplax tri $dentata_{i}$ -16.1 ± 0.2% for Parasesarma erythrodactyla) than for those in the mangroves (-21.1 ± 0.4 and -23.1± 0.2‰, respectively; ANOVA, factor position, p < 0.001 for both species, Fig. 4). The δ^{13} C signatures of A. tridentata (-15.2%) and P. erythrodactyla (-16.2%) in the saltmarsh were close to those of Sporobolus virginicus (-14.9 \pm 0.1%). The δ^{13} C of both species of crabs in the mangrove habitat were higher than those in the mangroves ($-27.6 \pm 0.2\%$). Values for *P. erythro*dactyla were very similar to those of MPB, whereas A. tridentata was slightly enriched relative to MPB.

A significant interaction between position and site was found for both *Australoplax tridentata* and *Parasesarma erythrodactyla* (ANOVA, interaction term, p < 0.001 for both species). The source of this variation is attributable to a single site (Site 5, Fig. 5) that had lower δ^{13} C values (*A. tridentata* –19.5 ± 1.2‰, *P. erythrodactyla* –20.2 ± 1.1‰) than all other sites (*A. tridentata* –17.1 ± 0.4‰, *P. erythrodactyla* –18.5 ± 0.4‰) predominantly due to the lower δ^{13} C values of crabs at the edge position. The δ^{13} C signatures of both crab species collected at the edge position at Site 5 (*A. tridentata* –19.4 ± 0.7‰, *P. erythrodactyla* –20.1 ± 0.6‰) were significantly lower than those from the other saltmarsh positions at that site (*A. tridentata* –14.4 ± 0.1‰, *P. erythrodactyla* –15.5 ± 0.7‰).

The δ^{13} C signatures of the gastropod *Ophicardelus quoyi* from the saltmarsh habitat did not differ significantly between the central (-14.9 ± 0.2%) and intermediate (-15.8 ± 0.2%) positions, but were significantly higher than those of snails from the edge positions (-17.1 ± 0.5%) (Fig. 4). The δ^{13} C of *Salinator solida* showed the same pattern of variation as for

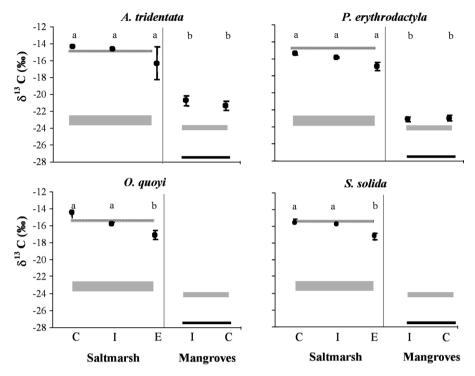


Fig. 4. Carbon stable isotope signatures for autotrophs (mangroves, Sporobolus virginicus and microphytobenthos, MPB), and for each crab (Australoplax tridentata and Parasesarma erythrodactyla) and gastropod (Ophicardelus quoyi and Salinator solida) species across

Mangroves

S. virginicus

MPB

sampling positions. C: centre; I: intermediate; E: edge. Values averaged across the 5 sites (±1 SE; SE too small to see in some cases). Widths of lines for autotrophs represent ±1 SE. Different letters indicate significant differences in ani-mal signatures among positions

O. quoyi, with snails from the central $(-15.2 \pm 0.2\%)$ and intermediate $(-15.3 \pm 0.1\%)$ positions not differing but being more enriched than snails from the edge $(-16.8 \pm 0.4\%)$ (Fig. 4).

DISCUSSION

Scales of carbon movement and assimilation

This study is the first to use stable carbon isotopes to link animal position and carbon source at the fine-scale between adjacent estuarine habitats. Since isotope signatures of all consumers varied with position, our results do not support the coarse-scale model (Fig. 1, Model a) of large-scale carbon movement and assimilation between adjacent saltmarsh and mangrove habitats. Carbon movement and assimilation by the crabs and gastropods in these 2 habitats therefore occurs at scales of $<\!30~\mathrm{m}$.

Our results indicate that the scale of carbon movement and assimilation is either towards the lower end of the intermediate model range (2 to 15 m) or <2 m (the fine-scale model). Support for the intermediate model is provided by the $\delta^{13} C$ of gastropods at all sites and crabs at Site 5. For gastropods, the slight but significant depletion in their $\delta^{13} C$ signatures at the edge position 2 m inside the saltmarsh habitat suggests movement and assimilation of carbon across the saltmarsh–mangrove interface. Likewise, crabs at Site 5

showed the same depletion in $\delta^{13}C$ at the edge position. Animals may also assimilate MPB produced *in situ* and this may explain the lower $\delta^{13}C$ signatures at edge positions. In this scenario, the depleted $\delta^{13}C$ signatures at the edge position for the gastropods and for the crabs at the edge position at Site 5 support more limited carbon exchange (i.e. the fine-scale model) across the saltmarsh–mangrove interface.

The δ^{13} C signatures of crabs in the saltmarsh habitat at sites other than Site 5 matched the δ^{13} C signature of the salt couch grass Sporobolus virginicus, the macrophyte in their immediate vicinity. Although the δ^{13} C signatures of crabs in the mangrove habitat were enriched relative to those of the mangroves, they matched those of the MPB in the mangrove habitat (closely for Parasesarma erythrodactyla, less so for Australoplax tridentata). The δ^{13} C signatures of crabs were therefore consistent with the fine-scale model of assimilation of carbon from producers in their immediate vicinity (Fig. 1, Model c). However, the δ^{13} C signatures of crabs in the mangrove habitat could also have resulted from a mixture of carbon sources from, for example, a combination of mangrove and salt couch grass, with or without MPB. In this scenario, the signatures of mangrove crabs suggest movement of carbon over larger distances, whilst those of saltmarsh crabs strongly indicate fine-scale movement. Investigating the relative importance of MPB to crabs in mangroves is pivotal to further differentiating among our models of carbon movement. Future work might best examine

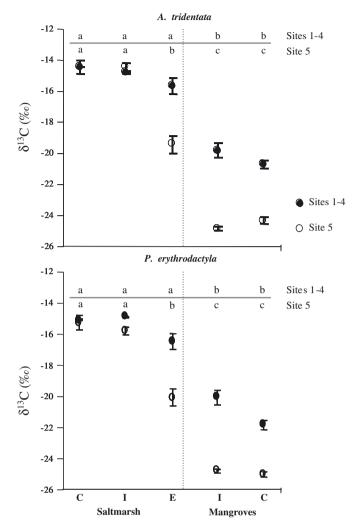


Fig. 5. Australoplax tridentata and Parasesarma erythrodactyla. Mean (±1 SE) isotope signatures at Sites 1 to 4 and Site 5 (separate). Different letters indicate significant differences in isotope signatures among positions. Abbreviations as in Fig. 4

the importance of MPB to mangrove animals using recently developed techniques for manipulating isotopic signatures of algae (e.g. Winning et al. 1999, Carman & Fry 2002).

The interface between the saltmarsh and mangrove habitats was very useful for testing among our models: although the carbon isotope values of salt couch grass and mangroves were very different, the values for each autotroph within any one habitat were similar at all positions. Elsewhere, the carbon isotope signatures of a single autotroph species adjacent to mangroves have themselves been shown to be affected by proximity to mangroves. For example, in a bay in Kenya, the δ^{13} C signatures of seagrass were 9% lower in seagrass adjacent to mangroves than in seagrass several kilometres away, because of the lower isotope value of the

dissolved inorganic carbon generated in the mangroves (Marguillier et al. 1997). The slight depletion of MPB we found in the mangrove relative to the saltmarsh habitat might well result from the same phenomenon. Our test among models for 2 adjacent habitats, focusing on differences among positions rather than on whether mangrove material per se was an important source to animals, also avoided the issue of generalised depletion of carbon values in the whole mangrove biota (France 1998).

The results of this study are in contrast to historical expectations of large-scale carbon movement in aquatic systems but are consistent with a recent study of the movement and assimilation of carbon between adjacent habitats at a smaller scale. Hsieh et al. (2002) analysed the carbon isotopes of ocypodid crabs at sites separated by hundreds of metres. They concluded that ocypodid crabs derive their carbon from the sites in which they reside rather than from sites further away. Ocypodid crabs (*Uca arcuata* and *U. borealis*) collected at a creek site were considered to have derived their carbon from MPB, although a mixed diet of algae, particulate organic matter or the detritus of a terrestrial grass was also a plausible source (Hsieh et al. 2002). However, at river sites 600 m from the creek site, the main source of carbon for these species was the terrestrial grass nearest to the river site. Our own work suggests that even the site-specific work of Hsieh et al. (2002) was at a scale much coarser than the movement and assimilation of carbon by shore crabs, which probably obtain their nutrition from sources in the surrounding few metres.

In this study, the change in $\delta^{13}C$ signatures between the saltmarsh and mangrove habitats was the same for both crab species and among all sites sampled. However, the isotope signatures of crabs at 1 site (Site 5) were lower at the edge and mangrove positions than at these positions in all other sites. Observations while sampling showed that the edge positions at this marsh appeared to drain more slowly than the edge positions on the other marshes. Whilst this difference in hydrology was not sufficient to influence the isotopic signatures of the plants, it may have influenced the feeding behaviour of the crabs at the edge positions.

Crab and gastropod diets

Both crab species collected in the saltmarsh in this study appear to derive their carbon from salt couch grass. Crabs in the mangroves may obtain their carbon predominantly from MPB, although their $\delta^{13}C$ signatures are also consistent with a mixture of carbon sources such as mangroves and salt couch grass, with or without a contribution from MPB. So far, no studies

have specifically examined the diet of these 2 crab species, but observations of feeding behaviour typically ascribe a wide range of feeding habits to ocypodid and grapsid crabs (Lee 1998).

The feeding habits of grapsid crabs include the harvesting of mangrove leaves from the canopy (Warner 1967) and the consumption of saltmarsh plants (e.g. Spartina densiflora, Iribarne et al. 1997) and mangrove leaf litter (Lee 1997). Sesarmid crabs (e.g. Parasesarma erythrodactyla) are typically considered to be herbivores that carry leaf litter into their burrows (Lee 1998), but recent evidence from gut-content analysis suggests a lack of dietary specialisation for some species (e.g. Sesarma leptosoma) that indicates a generalist mode of feeding and dietary assimilation (Dahdouh-Guebas et al. 1999). However, the δ^{13} C signatures of *P. aspe*rum collected from mangrove habitat were slightly more enriched than those of mangroves (Bouillon et al. 2002), a result similar to that for P. erythrodactyla in the current study. The slightly enriched δ^{13} C of *P. asperum* relative to those of mangroves was thought to result from a diet comprising a mixture of mangrove material and sediment scrapings.

Ocypodid crabs have been described as feeding on bacteria (e.g. *Uca vocans* and *U. polita*, Dye & Lasiak 1986) or microalgae (e.g. *U. dussumieri*, *U. forcipata*, *U. rosea* and *U. triangularis*, Rodelli et al. 1984), and a variety of plant matter including particulate organic matter and terrestrial grass (e.g. *U. arcuata* and *U. borealis*, Hsieh et al. 2002). Carbon stable isotope studies of *U. pugnax* in USA saltmarshes found that the signature of crabs closely matches that of the cordgrass *Spartina alterniflora* (Haines & Montague 1979, Currin et al. 1995). Even where ocypodids consume plant material, however, it is mostly as detritus, and they are also likely to ingest microbes and microscopic invertebrates.

There is little information on the diets of Salinator solida and Ophicardelus quoyi, but they are generally considered to either graze on microalgae or behave as detritivores (e.g. S. burmana, Rodelli et al. 1984). Microalgae are generally considered an important food source for macrobenthos in saltmarshes (Sullivan & Moncreiff 1990). In our study, however, the δ^{13} C signatures of gastropods in the saltmarsh habitat indicated that they obtain their carbon predominantly from salt couch grass. Analysis of the diet of 2 depositfeeding gastropods, Assiminea japonica and Angustassiminea castanea, revealed that although these snails feed mainly on phytoplankton and benthic algae in the wild, they were nevertheless able to assimilate organic matter from reed detritus under laboratory conditions (Kurata et al. 2001). This demonstrates, at least for these species, that they are physiologically capable of assimilating carbon from grasses.

Seagrass is also present in southern Moreton Bay,

seaward of the mangroves, and has an enriched δ^{13} C signature (-12%) similar to that of saltmarsh grass (Melville & Connolly 2003). Seagrass carbon could potentially be deposited on the marshes and be utilised by crabs and gastropods directly or indirectly. However, as there was no seagrass in the vicinity of the saltmarsh sites and no seagrass detritus observed at the site, it is unlikely that seagrass contributed much, if at all, to the consumer carbon source. The carbon isotope signatures of phytoplankton (~20%) (Fry 1984, Bouillon et al. 2000, Davenport & Bax 2002) also make it an unlikely source of carbon for consumers residing on the saltmarsh. Truly terrestrial vegetation can show similar isotopic signatures to that of mangroves (e.g. -27 to -30%, Boon & Bunn 1994), but the sites used in this study were several kilometres from terrestrial sources, minimising the likelihood of terrestrial vegetation contributing to the isotopic signatures of the consumers. As animal samples were composites of several individuals, it is also possible that differences in individual feeding behaviour, such as animals feeding entirely on MPB at the edge position, will have been undetected. Future studies using δ^{13} C values of individual animals would clarify individual feeding behaviour.

Whilst the aim of this study was to examine carbon movement and its assimilation by estuarine invertebrates, the movement of the target animals across habitat boundaries could also explain the observed patterns in δ^{13} C (e.g. Hobson 1999). Such movement is considered unlikely for the crabs and gastropods in this study. Little information on the movement of crabs and gastropods is available, but both crab species are considered to be faithful to their burrows, and the gastropods appear to move predominantly vertically within the sediment profile (M. A. Guest pers. obs.). This study used invertebrates of limited mobility that were either detritivores or grazers, to highlight potential carbon movement between adjacent habitats, but the results may have differed had animals of greater mobility and/or different feeding modes been examined. For example, had we examined sedentary filterfeeding bivalves on the marsh, their carbon signature might have reflected that of the water-column phytoplankton produced remotely rather than any in situ production. Further work on animals of differing mobility and feeding modes is therefore needed to achieve a more general conclusion about the scale of carbon movement and assimilation in estuarine systems.

It is possible that carbon may move among habitats at any of the scales presented but not be assimilated by the consumers sampled (Duarte & Cebrian 1996). Stable isotope analysis of consumers tests only the movement *and* assimilation of carbon. As this study set out to determine the trophic importance of discrete habitats over different spatial scales, only the movement

and assimilation of carbon has been examined. Movement of carbon that does not contribute to trophic dynamics has therefore not been considered. In this study, carbon isotope results from 2 types of estuarine invertebrates showed that the movement and assimilation of carbon occurs at scales much smaller than originally conceived in theories about the movement of carbon from inshore to offshore habitats.

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