High congruence of isotope sewage signals in multiple marine taxa

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Abstract

Assessments of sewage pollution routinely employ stable nitrogen isotope analysis ($\delta^{15}\text{N}$) in biota, but multiple taxa are rarely used. This single species focus leads to underreporting of whether derived spatial N patterns are consistent. Here we test the question of ‘reproducibility’, incorporating ‘taxonomic replication’ in the measurement of $\delta^{15}\text{N}$ gradients in algae, seagrasses, crabs and fish with distance from a sewage outfall on the Adelaide coast (southern Australia). Isotopic sewage signals were equally strong in all taxa and declined at the same rate. This congruence amongst taxa has not been reported previously. It implies that sewage-N propagates to fish via a tight spatial coupling between production and consumption processes, resulting from limited animal movement that closely preserves the spatial pollution imprint. In situations such as this where consumers mirror pollution signals of primary producers, analyses of higher trophic levels will capture a broader ambit of ecological effects.

Keywords: sewage; nitrogen isotopes; monitoring; spatial gradients; coastal waters;

1. Introduction

Nitrogen is pivotal in shaping the structure and function of natural systems at multiple levels of ecological organisation (Conley et al., 2009). Humankind has massively altered the cycling and mobility of nitrogen in the biosphere, adding unprecedented amounts of bioavailable N to ecosystems, most of it since the second half of the last century (Canfield et al., 2010; Vitousek et al., 1997). These anthropogenic inputs of N can have serious and widespread environmental consequences, altering productivity, diversity, trophic connections, species composition, organism health, and habitat quality (Cloern, 2001; Deegan et al., 2012).

Excess nitrogen from human sources is a component of a suite of stressors causing impacts that can be severe and widespread in coastal waters (Lotze et al., 2006; Reopanichkul et al., 2009). Fertilization of nearshore marine waters with human-derived nutrients is mainly the result of changing land-use, most notably the widespread and escalating urbanisation of coastal areas (Nixon and Buckley, 2002). Ecological effects of nutrient inputs therefore often centre on coastal cities (Diaz and Rosenberg, 2008; Oczkowski et al., 2009).

Nitrogen contained in human metabolic waste products is a major component of nutrient inputs to coastal waters (McClelland et al., 1997; Schlacher et al., 2005), and monitoring the effects of sewage entering marine ecosystems has become an important activity (Costanzo et al., 2001). One of the most widely applied techniques to identify the presence of sewage-N is the analysis of stable nitrogen isotopes in the tissues of biota putatively exposed to inorganic sewage-N or assimilating it from their diet (Cabana and Rasmussen, 1996; Pitt et al., 2009). The technique is based on predictable differences in the abundance of the heavy (^{15}N) and light (^{14}N) isotopes between sewage and other N sources such as fertilizers; nitrogen in treated wastewater is generally enriched in the heavier isotope ^{15}N , and monitoring of sewage inputs exploits this higher concentration of ^{15}N (Cole et al., 2004; Heaton, 1986).

Studies aimed at detecting nitrogen discharged in sewage based on stable nitrogen isotope ratios ($\delta^{15}\text{N}$) in organisms have drawn upon a wide variety of taxa, including algae (Dailer et al., 2010; Fernandes et al., 2012), rooted macrophytes (Cole et al., 2004; McClelland et al., 1997), sessile invertebrates (Carmichael et al., 2008; Carmichael et al., 2012; Fertig et al., 2010; Fertig et al., 2009; Risk et al., 2009), motile invertebrates (Bucci et al., 2007), and fish (Hoffman et al., 2012; Schlacher et al., 2005). Most report that the chosen taxon is 'suitable' as an ecological indicator of sewage-N, and that spatial patterns in relation to sewage sources can be mapped, or at least interpreted in a spatial context of known inputs (Costanzo et al., 2001; Northington and Hershey, 2006).

Investigations that include multiple species from different functional groups or from different trophic levels are less common (but see for example Pitt et al., 2009). Arguably, this lack of taxonomic and functional 'replication' raises two questions in relation to the assessment of isotopic sewage-N distributions: 1.) Are spatial patterns of the sewage signal reproducible using different taxa? ('repeatability'), and 2.) Are effect sizes (i.e. the magnitude of the isotopic sewage signal) comparable among taxa? ('sensitivity'). Here we examine both these questions using a two-staged approach: i) we first test for differences in the strength of the $\delta^{15}\text{N}$ gradient – comparing changes of $\delta^{15}\text{N}$ in multiple taxa with distance from a defined point source of sewage inputs in a shallow marine gulf of southern

Australia, and ii) we then examine the generality of our findings by comparisons with spatial $\delta^{15}\text{N}$ gradients in other settings where multiple taxa have been used to measure sewage-N gradients.

2. Methods

Gulf St Vincent (GSV) is a large, shallow marine embayment in southern Australia, bordering the city of Adelaide (Fig. 1). The general circulation of the gulf is clockwise, the western shores are bordered by mangroves, and the seafloor consists of a mosaic of bare sand and seagrass meadows and reefs (Jones et al., 2008). The GSV is located in an arid climatic zone where the annual evaporation exceeds rainfall, resulting in limited freshwater inputs from small local creeks and rivers (Shepherd and Sprigg, 1976). This makes discharges of sewage effluent on the eastern shores the main source of nutrients to receiving marine waters. Of the 1,431 tonnes of N entering the gulf waters from the city of Adelaide, 827 tonnes (58%) come from waste-water treatment plants (Fernandes et al., 2012). We designed our study to determine spatial gradient of sewage-N in relation to the largest point source, the Bolivar wastewater treatment plant: it is the main contributor of sewage-N to the gulf, discharging 509 tonnes of N per annum, or 62% of all treated N discharged in sewage effluents (Fernandes et al., 2012).

Fig. 1 Location of the study area in Gulf St Vincent of South Australia near the city of Adelaide and map of the sampling stations in relation to the major sewage outfall point on the eastern shores from the Bolivar treatment plant (top row), and distribution of $\delta^{15}\text{N}$ values in the tissues of marine biota sampled to determine the spatial gradients of sewage-N signals in the nearshore marine waters of the gulf (middle and bottom row). In the seagrass map triangles denote *Zostera* samples and circles denote *Posidonia*.

We collected multiple taxa to determine the spatial gradients of the sewage-N signal in relation to the defined sewage outfall of the Bolivar treatment plant (Fig. 1). Biota analysed for tissue $\delta^{15}\text{N}$ included primary producers (the green alga *Ulva lactuca* and two genera of seagrass, *Zostera* and *Posidonia*), mobile benthic invertebrates (the blue swimmer crab *Portunus armatus*), and pelagic fish (small-

mouthed hardyhead, *Atherinosoma microstoma*). Individuals were collected from 33 sites wherever they were present during the austral summer (Jan – Mar).

Muscle tissue was excised from crabs and fish, seagrass blades were cleaned of epiphytes. Samples were analysed on a 20-20 Europa Scientific mass spectrometer at the Stable Isotope Facility of Griffith University. Isotope values are expressed in the δ (‰) notation, $\delta^{15}\text{N}$, relative to the international standard of atmospheric nitrogen: $\delta^{15}\text{N} (\text{‰}) = ((^{15}\text{N}/^{14}\text{N}_{\text{sample}} \div ^{15}\text{N}/^{14}\text{N}_{\text{standard}}) - 1) \times 1000$. Precision of duplicate samples was 0.2‰.

3. Results

The isotope sewage signal in organisms from the Gulf St Vincent showed a strong spatial gradient that was consistent among multiple taxa (Fig. 1). The abundance of the heavier isotope ^{15}N declined significantly with distance from the outfall in the tissues of all taxa examined, consistent with a gradient of decreasing sewage-N contributions further from the point source (Figs. 1 & 2).

Fig. 2 Decline in tissue $\delta^{15}\text{N}$ at increasing distance from the sewage outfall in a) autotrophs (the green alga, *Ulva lactuca*, and two genera of seagrass (*Posidonia*, *Zostera*), and b) consumers (the portunid crab *Portunus armatus*, and hardyheads, *Atherinosoma microstoma*, a small pelagic fish).

This relationship between distance from the treatment plant's outfall and organism $\delta^{15}\text{N}$ values did not deviate significantly from a linear model in any species analysed (Runs test: min. $p = 0.236$). Tissue concentrations of ^{15}N declined at similar rates in all species, irrespective of their trophic position, or the mode of nitrogen acquisition in plants. There was no significant ($F_{4,39} = 0.28$, $p = 0.88$) difference in regression slopes among the five taxa analysed, with a common (pooled) slope estimate of -0.74 ; this translates to a decrease of 1 ‰ in $\delta^{15}\text{N}$ for every 1.33 km in distance from the outfall.

This consistency in the strength and direction of the sewage-N gradient among taxa has not been reported previously (Table 1, Fig. 3). On the contrary, taxa usually differ considerably in whether they express a sewage gradient (i.e. non-zero regression slopes), and if they do, they differ in the rate at which the isotopic sewage signal declines away from the source (Table 1, Fig. 3).

4. Discussion

We show that 1.) multiple taxa from different functional groups express spatial patterns of sewage-N at comparative levels (i.e. all species showed significant spatial declines in their ^{15}N tissue concentrations), and 2.) a very similar spatial gradient is reproduced by different taxa (i.e. essentially the same rates of ^{15}N decline). These findings are remarkable because other studies that have employed multiple taxa to assess spatial patterns of sewage-N show inconsistencies between taxa in terms of detecting a gradient, in terms of the strength of the inferred gradient, or both (Table 1, Fig. 3). Also, compared with other published accounts of variations in $\delta^{15}\text{N}$ in relation to sewage outfalls in multiple taxa, we report the strongest gradient of a sewage signature in nearshore marine biota to date (Table 1).

With the exception of Pitt et al. (2009), who also measured significant gradients in the three taxa they examined, species are generally reported to vary considerably in how strongly the spatial sewage pattern is expressed (Table 1). For example, Hansson et al. (1997) report strong gradients for zooplankton and epibenthic mysids, but failed to detect a pattern consistent with a sewage N-source in the smallest and largest size class of herring. Both Conlan et al (2006) and Gartner et al. (2002) found declines in organism $\delta^{15}\text{N}$ consistent with a spatial cline of the sewage signal in only a single species out of 5 and 7 examined (Table 1).

Our result of sewage-N gradients across all taxa is also unexpected because the taxa analysed differ fundamentally in their mode of nitrogen acquisition and subsequent rates of tissue turnover of the assimilated N. Primary producers incorporate inorganic N in dissolved form, whereas animals assimilate N from dietary organic particulates. Uptake of N in algae occurs only via their thalli, whereas seagrasses have a complementary uptake route via their roots (Touchette and Burkholder, 2000). The fact that all species, irrespective of uptake routes or the form of nitrogen used, carried a clear sewage isotope signal in their tissue indicates that sewage-N enters the food via several primary producers to be transferred upwards.

Table 1 Synopsis of spatial gradients in nitrogen isotopic sewage signals measured in organism away from defined point sources of sewage inputs to nearshore estuarine/marine waters. Compilation includes studies that determined spatial patterns of $\delta^{15}\text{N}$ in at least two different taxa with a minimum samples size of $n = 8$. Statistics are for log-linear models of organism $\delta^{15}\text{N}$ in relation to distance from the outfall.

Source	Location	Habitat	Load	No. Species / Taxa	Spatial scale	Max. $\delta^{15}\text{N}$ contrast	Slopes (range)	No. slopes >0	R ² (range)	ANCOVA (equal slopes?)
Hansson et al. (1997)	Northern Baltic	coastal sea	na	7: invertebrates 2x (zooplankton, mysids (<i>Mysis mixta</i> & <i>M. relicta</i>); fishes 5x (smelt, <i>Osmerus eperlans</i> , sprat, <i>Sprattus sprattus</i> , & herring, <i>Clupea harengus</i> , 3 size classes).	~ 29 km	8.9	1.01 - 19.90	5 (72%)	0.03 - 0.78	$F_{6,69} = 3.15$, $P = 0.009$
Tucker et al. (1999)	Boston Harbor	estuarine / embayment; semi-exposed	~ 11,700 t N y ⁻¹	2: macroalgae (<i>Ulva</i> & <i>Enteromorpha</i>); blue mussels (<i>Mytilus edulis</i>)	~ 17 km	8.3	1.75 - 6.93	1 (50%)	0.23 - 0.60	$F_{1,12} = 3.74$, $P = 0.077$
Gartner et al. (2002)	Western Australia	open coast, nearshore reef; exposed to semi-exposed	140 ML d ⁻¹	5: macroalgae 3x (<i>Ulva australis</i> , <i>Vidalia sp.</i> , <i>Ecklonia radiata</i>); invertebrates 2x (sponge, <i>Clathria sp.</i> , ascidian, <i>Pyura australis</i>)	~ 19 km	6.8	1.4 - 4.82	1 (20%)	0.05 - 0.52	$F_{4,48} = 1.29$, $P = 0.29$
Conlan et al. (2006)	Antarctica	subtidal, exposed	0.6 - 2.1 ML d ⁻¹	7: invertebrates 6x (burrowing clam, ascidian, soft coral, sea star, sea urchin, nemertean worm); fish (<i>Trematomus bernacchii</i>)	< 1 km	7.0	0.15 - 18.02	1 (14%)	0.01 - 0.43	$F_{6,98} = 2.57$, $P = 0.02$
Pitt et al. (2009)	SE Australia, Brisbane Estuary	estuary, sheltered	303 t N y ⁻¹	3: algae (filamentous algae on mangrove pneumatophores); mangrove leaves (<i>Avicennia marina</i>); shore crabs (<i>Australoplax tridentata</i> & <i>Parasesarma eythrogramma</i>)	~ 41 km	26.3	4.30 - 8.80	3 (100%)	0.41 - 0.69	$F_{2,168} = 9.78$, $P < 0.001$
This study	Gulf St Vincent, South Australia	embayment, semi-exposed	Mean flow = 93 ML d ⁻¹ ; Load = 487 t N y ⁻¹	5: macroalgae (<i>Ulva lactuca</i>); seagrasses 2x (<i>Zostera spp.</i> , <i>Posidonia spp.</i>); blue swimmer crab, <i>Portunus armatus</i> ; fish (Hardyheads, <i>Atherinosoma microstoma</i>)	~ 23 km	20.0	11.50 - 17.72	5 (100%)	0.46 - 0.86	$F_{4,39} = 0.31$, $P = 0.87$

Nitrogen is turned over quickly in algal tissue, in the local situation reaching equilibrium with DIN within 8 days (Fernandes et al., 2012). Tissue turnover in seagrasses and animals is considerably slower and more variable, typically ranging from weeks to months (Gaston and Suthers, 2004; Hesslein et al., 1993; Lepoint et al., 2002). Because tissue turnover rates determine the time interval over which biota integrate the sewage-N signals they have been exposed to, the consistency of the isotope patterns documented in this study suggest that the wastewater imprint in the receiving waters fronting the Adelaide metropolitan coast is a persistent feature. In other settings, smaller and short-lived nitrogen and sewage plumes have been traced in coastal waters (Connolly et al., 2009; Gaston et al., 2006; Schlacher et al., 2008), including ^{15}N abundance in beach clams exposed to pulsed outflows from a nearby estuary (Schlacher and Connolly, 2009).

We found that isotopic signals of sewage-N in primary producers were transferred to mobile consumers, including fish (Figs. 2 & 3). This upwards movement of an isotopic pattern through the food web to fishes is not unusual (e.g. Hobson et al., 2012). Mechanistically it depends on two fundamental conditions to be met: i) the diet of fish and other consumers includes plants that have incorporated the sewage-N (a 'dietary match'), and ii) consumers move little between feeding locations ('site fidelity'). In a trophic context, producers and consumers are spatially coupled in such situations where inorganic N (from sewage or other sources) enters plants as an inorganic substrate to be transferred to consumers with little spatial dispersion of the latter. Such consistent consumer-diet relationships (sensu Vanderklift and Wernberg, 2010) appear to operate in the nearshore waters of Gulf St Vincent, resulting in the closely matched spatial patterns between mobile consumers and sessile algae and plants.

Coastal fishes can have pronounced site fidelity (Skinner et al., 2012). More generally, when animals preserve spatial differences in isotope values over prolonged periods, stable isotope analysis becomes a tool to determine animal movement (Hobson, 1999). In our situation, the close match in isotope gradients between sessile plants and mobile consumers strongly suggests that movement of crabs and fish is limited; sewage-N is transferred locally in the food web with conservative dispersion away from the source (Fig. 2).

Fig. 3 Relationship between $\delta^{15}\text{N}$ in biota and distance from defined sewage outfall points in marine waters. Regression lines are shown for each taxon measured within a study: shades of green denote algae, brown colours rooted macrophytes, blues are for invertebrates and red shades for fishes; taxa included in each study are listed in Table 1 which also contains regression statistics for each panel shown here. Solid lines have slopes >0 (at $P < 0.05$), whereas dashed lines have slopes not significantly different from zero ($P > 0.05$).

The sewage-N signal was equally strong in crustaceans and fish as it was in autotrophs, providing unequivocal evidence that anthropogenic nitrogen is transferred upwards through the food chain in this system. Fundamentally, this transfer of N can have two, opposing, ecological consequences for consumers: positive effects in the form of greater secondary production and a flow-on effect of enhanced biological carbon sequestration in the system supplied with anthropogenic N; negative effects where the health and fitness of organisms becomes impaired (Schlacher et al., 2007), or where

habitats and ecosystems are degraded as a consequence of massive and persistent nutrient loadings (Diaz and Rosenberg, 2008). Elevated tissue ^{15}N concentrations are reliable indicators of the presence of assimilated sewage-N in organisms (Schlacher et al., 2005), but do not necessarily by themselves imply impaired health in all cases. Measurements of assimilated sewage-N represent a surrogate for complex contaminant pathogen mixtures released into the system via sewage. The specific pathological and chemical components within sewage effluent are not necessarily identified, but may be responsible for the impaired health in fish (*sensu* Mondon et al., 2001).

Lower water quality may contribute to the decline of seagrass meadows and coastal fisheries (Lotze et al., 2006; Orth et al., 2006; Waycott et al., 2009), but anthropogenic nutrients that are transferred to coastal consumers (*sensu* Connolly et al., 2009) also stimulate fisheries production (Oczkowski et al., 2009). Fisheries productivity has increased over time in areas under the influence of land-derived nutrient enrichment, and ‘anthropogenically enhanced fisheries’ can depend strongly on nutrient inputs from sewage and fertilizers (Oczkowski et al., 2009).

Indubitably, the growing nutrient enrichment of coastal marine waters is of concern (Cloern, 2001; Wulff et al., 2011), but the system-wide consequences of reducing nutrient inputs from sewage, including putative impacts on fisheries, may extend beyond simple improvements in water quality. For example, in the Gulf St Vincent, seagrasses are a large nitrogen sink, trapping up to 862 tonnes of N per year which represents about one-third of all nitrogen inputs; a sizeable fraction (38%) of that N is delivered in sewage discharges (Fernandes et al., 2009). Seagrasses are linked to fisheries production, chiefly because of their high tissue replacement (e.g. shedding of leaves) which liberates organic matter to become available for animal consumption (Connolly et al., 2005; Winning et al., 1999). It follows that reductions in nutrient loads through improved wastewater treatment processes may, hypothetically, have knock-on effects on fisheries species via lower carbon sequestration by seagrasses and other primary producers. Such higher ‘order effects’ are unknown and, if present, need to be carefully weighed against documented impacts of sewage on marine biota (Schlacher et al., 2007) and the broader environmental consequences resulting from inadvertent anthropogenic fertilization of coastal seas (Diaz and Rosenberg, 2008).

5. Conclusion

Marina taxa exposed to sewage discharged from a treatment plant displayed spatial gradients in the ^{15}N content of their tissues that reflected diminishing sewage-N contributions with distance from an outfall. We incorporated ‘taxonomic replication’ in the pollution assessment, using multiple taxa to gauge these spatial gradients. In contrast to previous accounts where sewage-N was mapped

isotopically in multiple taxa, isotope distributions were remarkably consistent amongst taxa. This congruence amongst taxa indicates several features of anthropogenic nitrogen movement in this system: i) sewage N enters food webs via several primary producers, ii) sewage-N is transferred up the food chain to fish, and iii) a tight spatial coupling between production and consumption processes resulting from limited animal movement that preserves the spatial sewage imprint.

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7. References

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