



Local-scale variation in trace elemental fingerprints of the estuarine bivalve *Austrovenus stutchburyi* within and between estuaries

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ABSTRACT: Trace elemental fingerprinting has been used to estimate population connectivity of marine species by using variations in the chemistry of shell produced in different water masses to determine the natal location of settlers. This method has been successfully used to differentiate between estuaries and coasts at spatial scales of 10s to 100s of km. Here, we investigated the suitability of trace elemental fingerprinting to distinguish subpopulation connectivity at management scales suitable for local fishery or conservation management. The trace element chemistry of *Austrovenus stutchburyi* shells was used to distinguish intra- and inter-estuary subpopulations at local (~1 to 10 km) scales, with ~99% success. Differences between 8 locations within a large harbour and 4 smaller neighbouring estuaries were detected. Sr:Ca, Zn:Ca and Mn:Ca elemental ratios were the primary drivers of inter-site trace elemental fingerprint differences. Intra-shell variation in elemental fingerprints was used to examine temporal stability and demonstrated that, although intra-shell differences were present, site-specific differences in elemental fingerprints were maintained. The results of this study suggest trace elemental fingerprinting can be a viable method to estimate population connectivity over small inter- and intra-estuary spatial scales.

KEY WORDS: LA-ICP-MS · Microchemistry · Larval dispersal · DFA

INTRODUCTION

Population connectivity occupies a central role in population dynamics (Cowen & Sponaugle 2009, Kool et al. 2013) and is relevant to marine management activities, including controlling the spread of invasive species, tracking species range shifts due to changing temperatures, marine reserve design and restoration efforts (Almany et al. 2009, Delaney et al. 2012, Kim et al. 2013, Sorte et al. 2013). The life history of many marine species consists of a pelagic larval phase which is capable of being transported distances rang-

ing from a few metres to 1000s of km (Grantham et al. 2003, Shanks 2009). When mature individuals exhibit no or limited mobility, this larval phase is critical to population connectivity (Largier 2003).

Quantifying the dispersal of marine larvae, however, is extremely difficult. Although artificial tagging techniques have met with some degree of success (Jones et al. 2005, Moran & Marko 2005), propagules are subject to significant mortality, advection and diffusion processes over large scales, which makes the recapture of tagged individuals unlikely (Levin 2006, Gawarkiewicz et al. 2007). While examination of the

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genetic structure of populations can be used to resolve broad-scale connectivity patterns, the temporal scale over which information can be obtained limits the applicability of these methods to local-scale management (Hedgecock et al. 2007, Ross et al. 2012).

Geochemical fingerprinting methods have considerable promise for overcoming the problems previously associated with tracking marine larvae. These techniques take advantage of natural variations in multi-elemental signals in carbonate structures (trace elemental fingerprints) accreted in geographically and chemically distinct water masses (e.g. Becker et al. 2005, Carson 2010, Sorte et al. 2013, Cathey et al. 2014). If larval calcified structures are metabolically inert, are formed in the natal water mass and are retained after settlement, their chemical signatures can be used to infer the origin(s) of individuals (Walther & Thorrold 2006, Becker et al. 2007). The major advantage this methodology has over other techniques is that all members of a population potentially carry markers of larval origins. Bivalves are ideal candidates for the use of trace elemental fingerprinting, as the larval shell (prodissoconch) forms within 24 h of the veliger larva hatching, and juveniles subsequently retain their prodissoconch after settlement (Moueza et al. 2006).

Prior to the application of trace elemental fingerprinting techniques to track the dispersal of marine larvae, it is essential that the spatial resolution at which sites can be distinguished from each other is established (Campana et al. 2000, Gillanders et al. 2001, Becker et al. 2005). To date, individuals collected from sites located at distances ranging from ~1 to >200 km have been able to be differentiated based on their trace elemental fingerprints (Dunphy et al. 2011, Cathey et al. 2014). Most studies have focused on establishing variation in trace elemental fingerprints over open coastlines or between closely located estuaries (e.g. Cathey et al. 2012, Sorte et al. 2013). In estuarine species, both biophysical models and empirical studies suggest that a large portion of larvae may be retained within the natal estuary (Lundquist et al. 2009, Peteiro & Shanks 2015). Given that many species exhibit a patchy distribution within estuaries (Thrush et al. 2003), intra-estuary dispersal over small scales (km) may be more relevant to the management of populations than inter-estuary dispersal.

Elements responsible for successfully discriminating individuals collected from different sites vary based on the species investigated, the area over

which samples were collected and the time of sampling (Cathey et al. 2012, Carson et al. 2013, Sorte et al. 2013). Nevertheless, there are some elements which are commonly used to differentiate between individuals including Mg, Cr, Mn, Co, Cu, Zn, Sr, Cs, Ba, Pb, and U (Carson et al. 2007). Identifying which elements contribute to variation in elemental fingerprints for a species of interest can allow for an optimal elemental suite to be selected for analysis, which enhances analytical throughput, therefore reducing costs (Dunphy et al. 2011).

The use of multiple complementary techniques may provide a more holistic understanding of larval dispersal than can be achieved using one method alone (Abaunza et al. 2008, Tanner et al. 2016). In this study, we investigated elemental fingerprinting in a region where biophysical larval tracking models have already been developed to predict larval dispersal and population connectivity of estuarine bivalves (Lundquist et al. 2009, Lundquist & Broekhuizen 2012). In addition to biophysical models existing in the area studied, the genetic structure of bivalve populations throughout New Zealand has been previously examined (Ross et al. 2012). The large scales of genetic population differentiation (>100 km), however, have limited applicability for local management of shellfish populations.

The primary objective of this study was to examine the spatial scale at which geochemical methods (i.e. trace elemental fingerprinting) can be used to differentiate between individuals of the bivalve *Austrovenus stutchburyi*. Small-scale (1 km to 10s of km) differences both within a single estuary and between closely located estuaries were examined. Intra-shell differences in elemental fingerprints were examined to determine if temporal variation in shell chemistry existed over the period of shell accretion. In addition, the trace element (TE) analytical suite which provided optimal classification success rates was determined to optimise future studies in this area.

MATERIALS AND METHODS

Study species

The New Zealand cockle *Austrovenus stutchburyi* is abundantly distributed within intertidal sand and mudflats in estuaries throughout New Zealand (Morton & Miller 1968). Spawning typically occurs in the warmer summer months, with peak spatfall previously observed in Whangarei Harbour between December and February (Lundquist & Broekhuizen

2012). The dispersal period lasts 2 to 3 wk, after which settlement occurs at a size of approximately 180 μm (Stephenson & Chanley 1979). The shell of infaunal bivalves is entirely composed of aragonite, with calcite not being deposited (Green et al. 2013). In addition to larval dispersal processes, there is evidence of post-settlement dispersal, with juveniles (<5 mm) being found in the water column and in the bedload as well as adults being observed burrowing through the sediment (Cummings et al. 1995, Hewitt et al. 1996, Lundquist et al. 2004).

Study site

Whangarei Harbour (Fig. 1) is located in the north-east of New Zealand ($35^{\circ}48' \text{S}$, $174^{\circ}26' \text{E}$), covers an area of approximately 102 km^2 at high tide and is approximately 24 km in length. The harbour consists of 54 km^2 of intertidal flats and includes a range of habitats such as mangrove forests, seagrass meadows, shellfish beds, and intertidal mud and sandflats (Morrison 2003). The harbour receives runoff from a catchment area which is approximately 300 km^2 in area (Griffiths 2012). The land use in the catchment is highly varied and includes urban inputs from Whangarei urban areas (10%), indigenous forest (20%), plantation forestry (10%) and high-productivity agricultural grassland (50%) (Griffiths 2012). The majority of freshwater input is from the Hātea River, located in the upper reaches of the harbour, which has a mean flow of $10 \text{ m}^3 \text{ s}^{-1}$ (Inglis et al. 2006).

Whangarei Harbour supports a significant commercial, recreational and traditional fishery of *A. stutchburyi*, with commercial efforts restricted to Snake Bank (Williams et al. 2006). There has been interest in the restoration of historically important *A. stutchburyi* populations in recent years (Cummings et al. 2007, Lundquist et al. 2009, Hewitt & Cummings 2013). These efforts have focused on the Takahiwai and Parua Bay sites (Fig. 1). There are a number of smaller estuaries located to the north and south of Whangarei Harbour which also support *A. stutchburyi* populations. Hydrodynamic models predict that larvae are highly connected within the harbour as well as being exported from the harbour to the adjacent coastline, promoting inter-estuary exchange (Lundquist et al. 2009, Lundquist & Broekhuizen 2012).

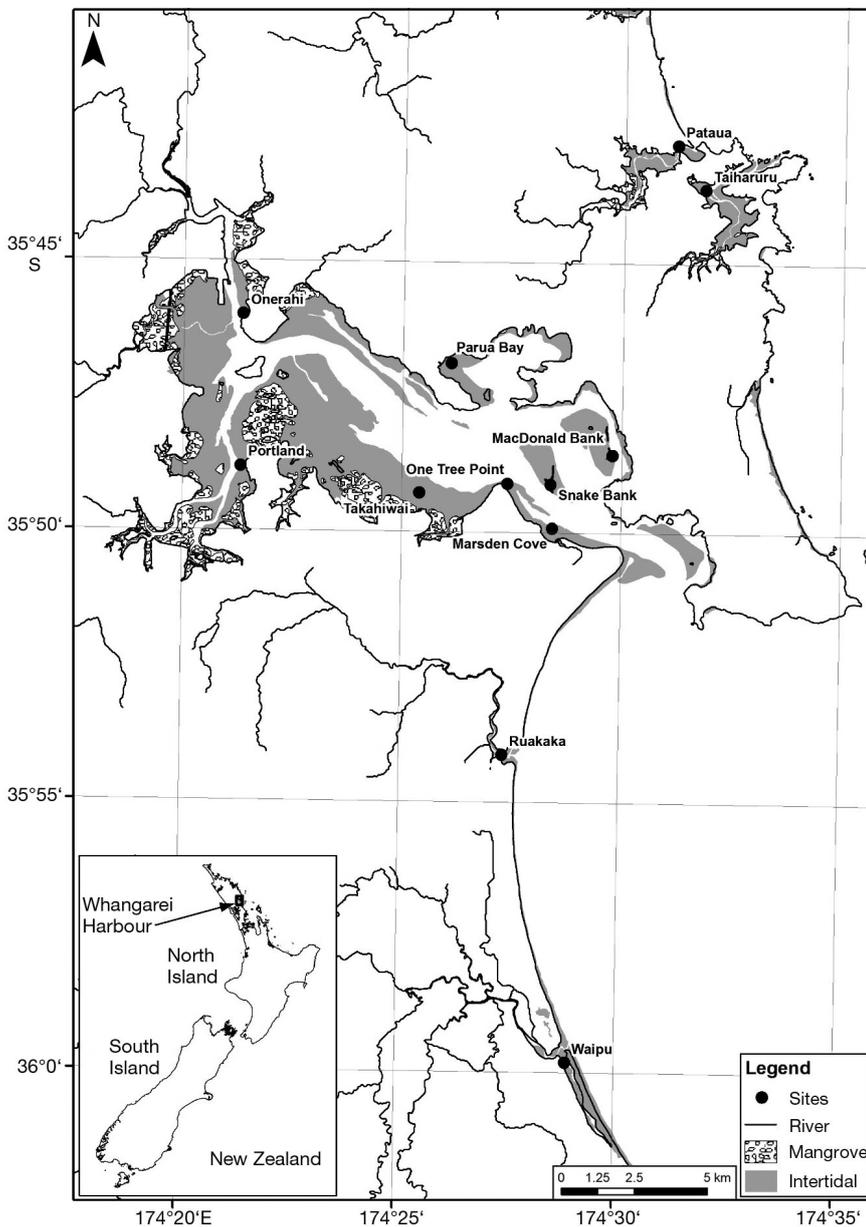


Fig. 1. Study area showing the sites sampled for *Austrovenus stutchburyi*. Shaded areas indicate intertidal regions within the estuaries

Sample collection

Animals were collected in the first week of March 2014 from 8 sites

located within Whangarei Harbour and from 2 estuaries located to the north and 2 located to the south along the same stretch of coast (Fig. 1). All *A. stutchburyi* were collected 2 h either side of low tide by excavating the upper 5 cm of the sediment surface with a plastic trowel and sieving this sediment through 2 stacked plastic sieves with a mesh size of 10 and 2 mm. The contents of the 10 mm sieve were discarded, and all *A. stutchburyi* were picked out of the 2 mm sieve and placed into a zip-lock bag containing seawater from the site. Juvenile bivalves were immediately placed on ice, frozen at the soonest possible opportunity and stored frozen at -20°C until analysis.

Sample preparation

Prior to TE analysis, individuals were removed from the freezer and defrosted. The shells were sorted, and 45 to 50 individuals with a shell height of approximately 5 mm were selected for microchemical analysis from each site. Linear growth rates of 10 mm yr^{-1} typically characterise the first 2 yr, suggesting that individuals selected were <6 mo in age (Adkins et al. 2014). Using stainless steel forceps, valves were split open, the flesh was removed and the shells were physically cleaned to remove any large adhering particles. Valves were placed into 50 ml centrifuge tubes filled with deionised water (which was maintained at a resistivity of $<1\text{ M}\Omega\text{ cm}$) and sonicated for 2 min to remove any remaining organic matter (Strasser et al. 2008). The water was drained, and the shells were allowed to air dry. Finally, the valves were mounted onto microscope slides by placing a bed of adhesive putty on the microscope slides and placing the left valve of each individual onto this base.

Laser ablation inductively coupled plasma mass spectrometry

Elements within juvenile shells were analysed by laser ablation inductively coupled plasma mass spectrometry (LA-ICP-MS). All analyses were conducted

Table 1. Laser ablation inductively coupled plasma mass spectrometry (LA-ICP-MS) operating parameters

Operating parameter	Operating conditions
Tuning	
Tuning standard	NIST 610
Ablation mode	Rastering ($2\text{ }\mu\text{m s}^{-1}$) beneath a $100\text{ }\mu\text{m}$ spot
Monitored isotopes	^7Li , ^{59}Co , ^{88}Sr , ^{140}Ce , ^{208}Pb , ^{238}U
Carrier gas (Ar)	0.85 l min^{-1}
Ablation gas (He)	80–90 %
RF power	1500 W
RF matching	1.72 volts
Repetition rate	5 Hz
Laser power	65 %
Standards and calibration	
Calibration standard	NIST 610
Ablation mode	$100\text{ }\mu\text{m}$ spot
Repetition rate	5 Hz
Laser power	65 %
Background acquisition	30 s
Sample data acquisition	45 s
Washout time	30 s
Measured isotopes	^7Li , ^{11}B , ^{25}Mg , ^{27}Al , ^{43}Ca , ^{47}Ti , ^{51}V , ^{55}Mn , ^{57}Fe , ^{59}Co , ^{60}Ni , ^{63}Cu , ^{66}Zn , ^{88}Sr , ^{89}Y , ^{138}Ba , ^{139}La , ^{140}Ce , ^{208}Pb , ^{238}U
<i>Austrovenus stutchburyi</i> method	
Repetition rate	5 Hz
Ablation mode	$100\text{ }\mu\text{m}$ spot
Laser power	65 %
Background acquisition	30 s
Sample data acquisition	45 s
Washout time	30 s
Measured isotopes	^7Li , ^{11}B , ^{25}Mg , ^{27}Al , ^{43}Ca , ^{47}Ti , ^{51}V , ^{55}Mn , ^{57}Fe , ^{59}Co , ^{60}Ni , ^{63}Cu , ^{66}Zn , ^{88}Sr , ^{89}Y , ^{138}Ba , ^{139}La , ^{140}Ce , ^{208}Pb , ^{238}U

with a New Wave deep ultraviolet (193 nm) excimer laser ablation system (Electro Scientific Industries) coupled to an Agilent 7700 ICP-MS (Agilent Technologies). Two spots were ablated on each shell, one located approximately $50\text{ }\mu\text{m}$ from the shell's most recently formed edge (the ventral margin) and one located approximately halfway between the ventral margin and the umbo. The approximate location of analyses is shown in Fig. 2. Shells were ablated with a laser spot size of $100\text{ }\mu\text{m}$, a repetition rate of 5 Hz and laser power of 65 %. The dwell time was 45 s. Two analyses of the NIST 610 glass standards were performed prior to and at the completion of each analytical session as well after every 25 laser spots performed to correct for elemental fractionation and machine drift. Gas flows and voltages were optimised to maximise sensitivity and signal stability while maintaining an oxide reduction ratio of less than 2 %. This was achieved by monitoring the thorium:thorium oxide ratios. Full LA-ICP-MS operating parameters are listed in Table 1.

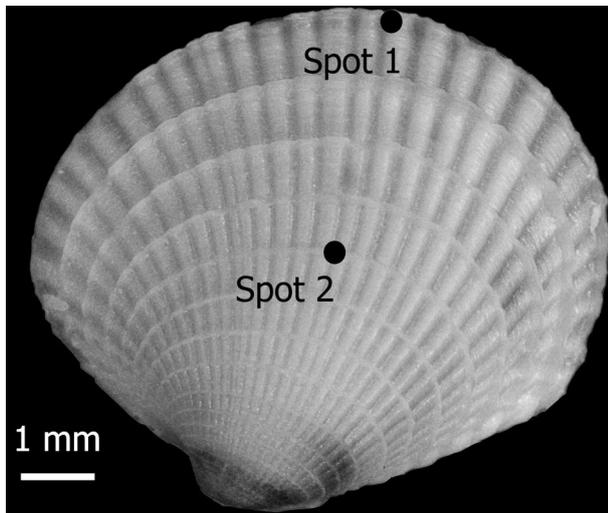


Fig. 2. Locations at which LA-ICP-MS spot analyses were performed on juvenile *Austrovenus stutchburyi*. Black spots mark analytical locations on the shell

The internal precisions of the TE:Ca ratios (% 2 SD relative) of the LA-ICP-MS were calculated from the NIST 610 standard analyses performed in each analytical session. Internal precisions for TE:Ca analyses were Sr:Ca $\pm 7\%$, Zn:Ca $\pm 6\%$, Mn:Ca $\pm 5\%$, Fe:Ca $\pm 18\%$, B:Ca $\pm 6\%$, Ba:Ca $\pm 4\%$, V:Ca $\pm 5\%$, Mg:Ca $\pm 5\%$, U:Ca $\pm 6\%$, La:Ca $\pm 6\%$, Al:Ca $\pm 5\%$, Co:Ca $\pm 5\%$, Ce:Ca $\pm 6\%$, Pb:Ca $\pm 6\%$, Y:Ca $\pm 7\%$, Cu:Ca $\pm 14\%$, Li:Ca $\pm 6\%$, Ti:Ca $\pm 5\%$, Ni:Ca $\pm 6\%$. Detection limits were calculated as the ratio of 3 times the standard deviation of the background to the NIST 610 analyses. The detection limits ($\mu\text{mol}:\text{mol}$) were Sr:Ca 2.1, Zn:Ca 1.6, Mn:Ca 1.3, Fe:Ca 1.3, B:Ca 0.2, Ba:Ca 3.3, V:Ca 1.2, Mg:Ca 0.6, U:Ca 5.8, La:Ca 3.4, Al:Ca 0.7, Co:Ca 1.4, Ce:Ca 3.4, Pb:Ca 5.1, Y:Ca 2.1, Cu:Ca 1.6, Li:Ca 0.2, Ti:Ca 1.2, Ni:Ca 1.4.

Ablation procedure

No chemical pre-cleaning was performed; rather, a pre-ablation technique was performed to remove contaminants from the shell surface. This procedure has previously been used when examining biogenic carbonates (Creech et al. 2010, Bolton et al. 2011, Marr et al. 2011). In these studies, inspection of the ablation profile was used to establish the removal of the contaminated surface veneer through laser ablation and to establish that the structure of interest was being ablated. In the present study, particular attention was paid to the Mg:Ca ratio, as organic material has been shown to be enriched in Mg relative to the

carbonate crystal matrix (Schöne et al. 2010). A typical ^{25}Mg ablation profile is shown in Fig. 3. Organic matter was deemed to have been removed when the ^{25}Mg ablation counts stabilised after an initial rapid decrease. The drop in the ^{25}Mg ablation counts always occurred within the first 10 s of analysis; therefore, this period was excluded from calculations of elemental concentrations. ^{43}Ca ablation counts per second were also monitored throughout analysis to ensure that shell material was being analysed and that ablation of adhesive putty and glass slides did not accidentally occur. The final 10 s of the laser ablation signal was also excluded to prevent complications associated with the drop in laser power at the end of the dwell time.

Due to the exclusion of the first and last 10 s of ablation data, only the middle 25 s of data was included in calculations of elemental concentrations. Data from the middle 25 s of the laser dwell period were then background corrected by subtracting the background average counts prior to analysis from the ablation average counts. The background-corrected counts per second were converted to TE:Ca ratios and normalised using the most recently published NIST 610 values (Jochum et al. 2005) to correct for instrumental drift and elemental fractionation. Finally, TE:Ca ratios were converted to $\mu\text{mol}:\text{mol}$ ratios, which were used in all statistical analyses.

Statistical analyses

To determine variation in elemental fingerprints, a discriminant function analysis (DFA) was performed in JMP 12.0 software (SAS Institute). Intra-shell variation in elemental fingerprints was examined using

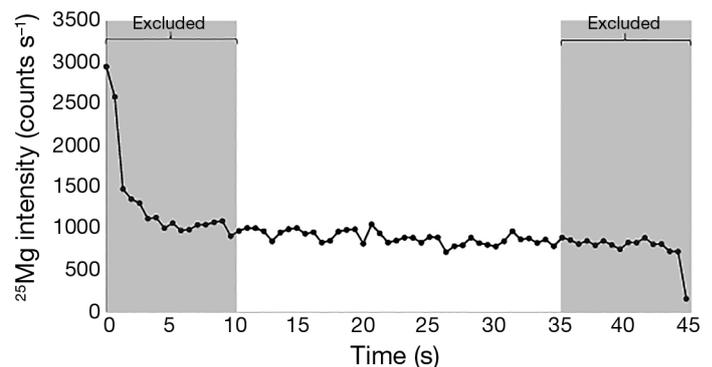


Fig. 3. Representative Mg ablation profile during a 45 s analysis showing the excluded pre- and post-ablation periods

the TE:Ca ratios of each individual laser spot (2 spots within each shell). Correlation matrices were generated for each site/spot using MANOVA, and due to differing covariance estimates among sites, a quadratic DFA (Q-DFA) was performed. As non-normality was detected, data were rank transformed (Bello 1993) prior to the Q-DFA. The mean canonical scores for each spot were calculated to show how elemental fingerprints varied within each shell collected from each of the sampling sites.

To examine inter-site differences, the mean of the 2 spots performed on each shell was used to obtain a single measurement of TE:Ca ratios for each *A. stutchburyi* shell analysed. The mean of the 2 spots (Fig. 2) was used to reduce the effects of small temporal fluctuations in shell chemistry. Correlation matrices were generated for each site using MANOVA, and a Q-DFA was performed due to differing covariances. Data were also rank transformed due to non-normality. The Q-DFA was performed stepwise by first running the analysis on all elemental ratios and recording the overall classification success. Next, the elemental ratio with the least discriminatory influence was removed, and the Q-DFA was run again with the difference in classification success noted. This was repeated for all elemental ratios to identify the point at which a major decrease in the classification success was observed and thus which combination of elements could provide successful discrimination of sites. A classification matrix was generated to gauge the success of the Q-DFA in assigning shells back to their site of collection based on elemental ratios, and mean canonical scores were calculated.

When performing DFAs, Meyers et al. (2013) have suggested that, to be statistically robust, the sample size (n) for the smallest group (i.e. site) should at least exceed the number of predictor variables (i.e. elemental ratios) used. In this case, the number of predictor variables was 19 overall, 10 of which contributed to classification success, and the smallest sample size was 38 (from the Waipu site).

Univariate tests were performed on the 4 elements which were responsible for the largest portion of classification success in the Q-DFA examining inter-site differences. A 1-way ANOVA was performed on the Sr:Ca, Mn:Ca, Zn:Ca and Fe:Ca ratios. The data were examined, and as they did not fit the assumptions of normality, data were log transformed. Outliers which fell more than ± 3 SD from the mean were removed. Tukey's HSD tests were then performed on those elemental ratios which differed significantly between sites to determine pairwise site differences.

RESULTS

Intra-shell variation

The Q-DFA examining the intra-shell differences in elemental fingerprints was able to assign shells to their collection location with 92% accuracy and to the collection location/spot with 82% accuracy. These results, as well as the plot of the canonical scores (Fig. 4), indicate that although intra-shell differences did exist in the elemental fingerprints of *Austrovenus stutchburyi*, these differences did not mask inter-site differences.

Spatial differences in trace elemental fingerprints

The Q-DFA performed using the mean of the 2 spots analysed on each shell was able to correctly assign juvenile *A. stutchburyi* to their site of collection with 99% accuracy (the classification success rates expected by chance are 8.3%) (Table 2; Fig. 5). This result was achieved with the stepwise inclusion of 10 elemental ratios (Sr:Ca, Zn:Ca, Mn:Ca, Fe:Ca, B:Ca, Ba:Ca, V:Ca, Mg:Ca, U:Ca and La:Ca) into the Q-DFA model (Table 3). The high overall accuracy of the Q-DFA was also reflected in the individual site classification success rates, which ranged from 95 to 100% (Table 2). Sites within the harbour located as close together as 1.2 km (Snake Bank and One Tree Point) were able to be reliably distinguished from each other with 100% accuracy. The maximum dis-

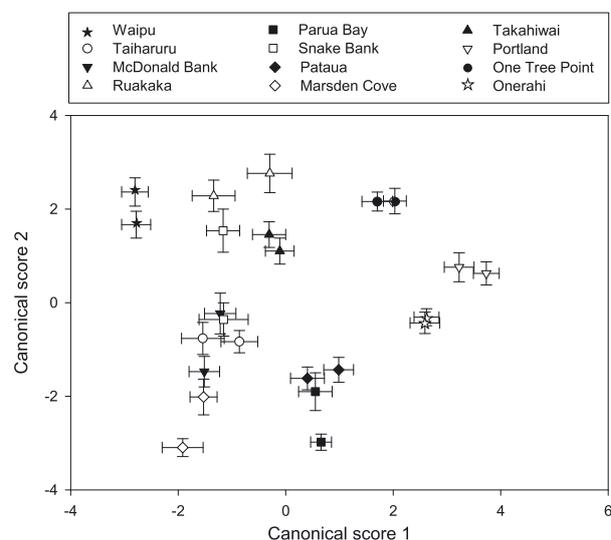


Fig. 4. Mean canonical scores ($\pm 95\%$ CI) from the quadratic discriminant function analysis investigating variation in the elemental fingerprints between the 2 spots performed on each shell

Table 2. Classification matrix from the quadratic discriminant function analysis of trace element:calcium ratios of juvenile *Austrovenus stutchburyi* shells collected from the Whangarei region in March 2014. Sites are arranged from west to east and grouped according to the region from which they were collected. Values in bold indicate the number of individuals for which the actual and predicted site were the same

Actual site	Predicted site												Total n	n classified correctly	% classified correctly
	Onerahi	Portland	Takahiwai	Parua	Marsden Cove	One Tree Point	McDonald Bank	Snake Bank	Ruakaka	Waipu	Pataua	Taiharuru			
Onerahi	47	0	0	0	0	0	0	0	0	0	0	0	47	47	100.0
Portland	1	42	0	0	0	0	0	0	0	0	0	0	43	42	97.7
Takahiwai	0	0	44	0	0	0	0	0	0	0	0	0	44	44	100.0
Parua	0	0	0	43	0	0	0	0	1	0	0	0	44	43	97.7
Marsden cove	0	0	0	0	42	0	0	0	0	0	0	0	42	42	100.0
One Tree Point	0	0	0	0	0	30	0	0	0	0	0	0	30	30	100.0
McDonald Bank	0	0	0	0	0	0	46	0	0	0	0	0	46	46	100.0
Snake Bank	0	0	0	0	0	0	0	45	0	0	0	0	45	45	100.0
Ruakaka	0	0	0	1	0	0	0	0	42	0	0	0	43	42	97.7
Waipu	0	0	0	0	0	0	0	0	0	38	0	0	38	38	100.0
Pataua	0	0	0	0	0	0	0	0	0	0	45	0	45	45	100.0
Taiharuru	0	0	0	0	0	0	1	0	1	0	0	39	41	39	95.1
Total	48	42	44	44	42	30	47	45	44	38	45	39	508	503	99.0

tance between sites which were able to be distinguished based on their trace elemental fingerprints was 35 km (Waipu and Portland). Inter-estuary differences in trace elemental fingerprints were also pronounced, and only 3 individuals (0.7% of those collected) were classified to an estuary other than that from which they were collected (21% would be expected by chance). The smallest distance between estuaries in this study was 1 km (Pataua and Taiharuru); despite their proximity, the Q-DFA was 100% successful at distinguishing between them, with no cross-classification errors occurring.

The results of the stepwise variable selection procedure showed that although 10 TE:Ca ratios were able to produce a DFA model which classified settled shells to their site of collection with 99% accuracy, fewer ratios contributed the majority of discriminating power of the model (Table 3). Sr was the primary driver of spatial variation, with 43% of individuals classified correctly using only the Sr:Ca ratio. The B:Ca and Mn:Ca ratios were also important discriminators for the accuracy of the model, contributing to 20 and 15%, respectively, of the model's discriminating power, bringing the total classification success to 80% using only these 3 elements.

Results of the ANOVA indicated that significant differences existed between sites in the 4 elemental ratios responsible for the majority of classification success (Table 4) (Sr:Ca $p < 0.01$, Mn:Ca $p < 0.01$, Zn:Ca $p < 0.01$, Fe:Ca $p < 0.01$). Tukey's HSD post hoc tests indicated that significant pairwise differences existed between most sites in the TE:Ca ratios examined (Table 5). The proximity of sites to each

other did not appear to influence whether pairwise differences in the Sr:Ca, Mn:Ca, Zn:Ca or Fe:Ca ratios were detected. Parua Bay and Snake Bank, for example, are separated by only 1.3 km but differed significantly in all TE:Ca ratios with the exception of Fe:Ca. In contrast, Tahiruru and Onerahi were located the furthest distance apart of any 2 sites in this study but did not differ significantly in their Sr:Ca, Mn:Ca or Fe:Ca ratios.

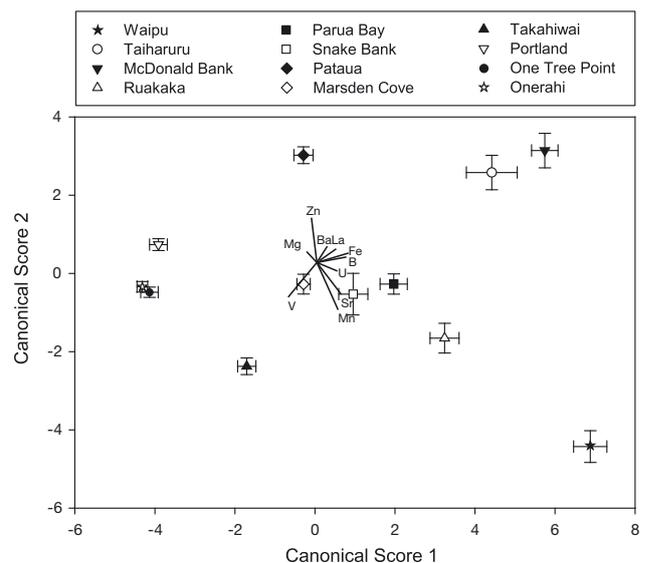


Fig. 5. Mean canonical scores ($\pm 95\%$ CI) from the quadratic discriminant function analysis (Q-DFA) examining the inter-site differences in mean trace elemental fingerprints of both analytical locations within each shell. Overlaid elemental vectors show how each of the elements included in the Q-DFA is driving differentiation between sites

Table 3. Results of the stepwise variable selection procedure of the quadratic discriminant function analysis model comparing elemental ratios of *Austrovenus stutchburyi*. Note: shaded elements indicate those used to produce the classification matrix in Table 2

Elemental ratio	Total % classified correctly	Individual % classified correctly
Sr:Ca	42.91	42.91
Zn:Ca	65.16	22.25
Mn:Ca	80.91	15.75
Fe:Ca	88.19	7.28
B:Ca	91.14	2.95
Ba:Ca	93.7	2.56
V:Ca	95.47	1.77
Mg:Ca	97.05	1.58
U:Ca	98.42	1.37
La:Ca	99.02	0.6
Al:Ca	99.21	0.19
Co:Ca	99.61	0.4
Ce:Ca	99.61	0
Pb:Ca	99.61	0
Y:Ca	99.8	0.19
Cu:Ca	99.8	0
Li:Ca	100	0.2
Ti:Ca	100	0
Ni:Ca	100	0

DISCUSSION

The results of this study demonstrate (1) that trace elemental fingerprinting techniques can be used to differentiate between *Austrovenus stutchburyi* populations over small intra- and inter-estuary scales (as small as 1.2 km) relevant to their management; (2) that although intra-shell variation in elemental fingerprints exists, inter-site differences are maintained regardless of the location of the analysis within settled shell; and (3) that an elemental suite consisting of 10 TE:Ca ratios provides optimal classification success, with the Sr:Ca, Mn:Ca and Zn:Ca ratios together providing over 80% classification success.

Intra-shell variation

The results of the Q-DFA investigating variation in the trace elemental fingerprints within *A. stutchburyi* shells demonstrated that although intra-shell variation in trace elemental fingerprints existed, this was not

sufficient to mask inter-site differences. It is likely that environmental stability contributed to the lack of variation in TE:Ca ratios within each shell. Shell analysed in this study was likely deposited over approximately 3 mo (using linear growth rates of 10 mm yr⁻¹ from Adkins et al. 2014). Shell material was, therefore, likely deposited over the austral summer, when the environmental conditions which may influence elemental incorporation into bivalve shell (i.e. temperature and salinity) were stable (Beer et al. 2011, Poulain et al. 2015). Mean air temperatures from December 2013 to February 2014 ranged from 19.1 to 20.1°C (NIWA 2016). These results imply that shell material deposited at different times over a period of environmental stability can be compared, therefore reducing the need for frequent sampling to maintain up-to-date data on elemental fingerprints between sites (Carson et al. 2013).

These results also suggest that age effects did not influence the incorporation of TEs into settled juvenile shell. Further work, however, should be performed prior to the extension of this finding to the prodissoconch. There are a number of differences between settled shell and prodissoconch shell which may hinder their comparison. Physical changes in the environment as a shell transitions from a pelagic to an infaunal lifestyle may result in different elements being available for incorporation into growing shell. It has previously been demonstrated that differences in the incorporation of elements to bivalve shell vary between larval and recruited shell (Zacherl et al. 2009). It has also been found that the diet of individuals may influence the uptake of elements into bivalve shell (Thebault et al. 2009). Furthermore, it is

Table 4. Mean elemental levels ($\mu\text{mol}:\text{mol} \pm \text{SE}$) of the 4 elements responsible for the majority of the classification success of the quadratic discriminant function analysis. The p-value for all ANOVAs examining inter-site differences in elemental ratios was <0.01

Collection site	Mn:Ca	Sr:Ca	Zn:Ca	Fe:Ca
Waipu	0.6 \pm 0.05	1455.8 \pm 36.5	6.3 \pm 0.4	861.9 \pm 46.3
Taiharuru	5.1 \pm 0.9	2090.2 \pm 28.9	6.5 \pm 0.7	1316.3 \pm 111.1
McDonald Bank	3.0 \pm 0.2	2176.0 \pm 44.5	2.5 \pm 0.2	531.6 \pm 39.4
Ruakaka	27.2 \pm 4.5	1445.2 \pm 34.7	12.4 \pm 2.2	901.0 \pm 59.5
Parua Bay	28.7 \pm 3.7	1851.1 \pm 46.9	19.7 \pm 1.3	488.6 \pm 29.9
Snake Bank	2.3 \pm 0.2	2173.6 \pm 78.6	8.1 \pm 1.1	703.2 \pm 44.3
Pataua	3.3 \pm 0.3	2528.8 \pm 23.9	1.7 \pm 0.1	573.2 \pm 33.8
Marsden Cove	1.5 \pm 0.0	2605.8 \pm 25.8	20.1 \pm 1.5	280.9 \pm 6.4
Takahiwai	6.2 \pm 1.4	1755.5 \pm 25.8	31.2 \pm 2.8	975.7 \pm 86.5
Portland	36.7 \pm 5.2	2048.3 \pm 18.0	8.3 \pm 0.9	1395.2 \pm 154.0
One Tree Point	24.3 \pm 1.5	2030.4 \pm 21.6	7.2 \pm 0.6	1379.3 \pm 103.2
Onerahi	10.4 \pm 0.7	1968.7 \pm 20.4	19.9 \pm 1.1	797.1 \pm 34.8

elemental fingerprints vary between sites located close to each other in a single estuary and were able to distinguish between sites located less than 1 km apart with over 80% accuracy. The advantage of the current study over Ricardo et al. (2015) is that both a higher number of sites were sampled and more individuals from each site were analysed.

Optimal classification success occurred with the inclusion of 10 TE:Ca ratios into the DFA model. This information is valuable, as it limits the number of elements needing to be quantified in future geochemical fingerprinting studies on this species in this area. The 3 elements (Sr, Zn and Mn) which dictated the majority of classification success are frequently used as part of the trace elemental suite in fingerprinting studies (e.g. Broadaway & Hannigan 2012, Sorte et al. 2013, Cathey et al. 2014). In fact, even using only these elemental ratios achieves a classification success rate comparable to many other studies (e.g. Carson et al. 2007, Dunphy et al. 2011, Fodrie et al. 2011).

Drivers of elemental variation

The differences in the elemental composition of *A. stutchburyi* shells in this study were likely due to a number of factors. The incorporation of elements into bivalve shell has been shown to be influenced by biological and environmental factors such as growth rate, temperature, salinity or the concentration of elements in the surrounding environment (Lloyd et al. 2008, Beer et al. 2011, Tabouret et al. 2012, Poulain et al. 2015). As high-resolution temperature and seawater composition data are not available over the study area, it is only possible to speculate on why Sr, Zn and Mn were responsible for the high classification rates observed.

Whangarei Harbour and its catchment are highly modified with urban, industrial and agricultural inputs present, which may have affected the trace elemental concentrations observed. Coastal geology and geography may also have influenced the incorporation of elements into *A. stutchburyi* shell. Weathering and erosion of rocks are important sources of TEs into marine systems and may influence TE:Ca ratios in shell (Lorens & Bender 1980, Vitousek et al. 1999, Lloyd et al. 2008). From One Tree Point to the south, the coastal plain is primarily composed of Holocene alluvium, while at Pataua, sandstone and argillite rocks dominate the catchment. Whangarei Harbour's catchment is composed of a variety of lithologically distinct geological units, possibly con-

tributing to the strong spatial differences in TE fingerprints observed within the estuary (Brook & Edbrooke 2009).

The Sr:Ca ratio was responsible for the highest degree of classification success in the Q-DFA model investigating inter-site differences. This high discriminatory power of the Sr:Ca ratio suggests that Sr isotope ratios ($^{87}\text{Sr}/^{86}\text{Sr}$) might also be a powerful complementary discrimination tool along with TEs (Hobbs et al. 2005). Drivers of Sr:Ca variation over the study area are difficult to determine due to the number of processes which may control its incorporation into biogenic carbonate such as salinity and ambient Sr concentrations (Lorens & Bender 1980, Zacherl et al. 2003, Heinemann et al. 2011, Poulain et al. 2015). The Sr:Ca ratio in *A. stutchburyi* did not follow a geographic pattern, nor was it related to possible anthropogenic activities at sites. The 2 sites with the highest Sr:Ca ratios, for example, were Pataua, located close to a freshwater input in a small village, and Marsden Cove, located close to a popular recreational marina. While freshwater input at Pataua may have driven the higher Sr:Ca ratios, Waipu and Ruakaka were also located near freshwater inputs but had the lowest Sr:Ca ratios of all the study sites (Poulain et al. 2015).

Zn played a large role in the discriminatory power of the Q-DFA model. Heavy metals such as Zn may enter estuarine systems through runoff from anthropogenic sources. These heavy metals can then bind to sediments and be released, the equilibrium between the 2 being affected by complex factors such as salinity and pH (Hatje et al. 2003, Fukunaga & Anderson 2011). In this study, however, the highest Zn:Ca ratios were found at Parua Bay and Takahiwai; both sites are dominated by agricultural grassland in their catchment area and are not located close to any potential sources of heavy metal contamination. In seawater, bound Zn may be released from sediments more readily than in fresh water, suggesting that salinity may affect its bioavailability (Hatje et al. 2003). This deabsorption of Zn from sediments may explain the low Zn:Ca ratios found in shell collected from Pataua, which is located in an estuary with significant freshwater input.

The Mn:Ca ratio was also responsible for a large portion of the discriminatory power of the Q-DFA. The availability of the Mn^{2+} ion may be influenced by microbial action or through input of high Mn sediments to the marine system (Klinkhammer & McManus 2001, Husmann et al. 2012). Shells collected from Waipu were found to have the lowest Mn:Ca ratios. The Waipu River runs through a large

area of agricultural farmland, from which runoff into the river may promote microbial activity. Microbes form Mn oxides and therefore remove Mn^{2+} ions from the surrounding environment, thereby reducing their availability to organisms, which may explain the low Mn:Ca ratios found at Waipu (Klinkhammer & McManus 2001). The highest Mn:Ca ratios were found at Portland. The sediment in the upper harbour region in which this site is located is primarily mud (<63 μm grain size) (Millar 1980). Sediment with smaller grain sizes is more prone to resuspension than larger grains (Cornwell & Porter 2009). Metals associated with suspended particulate matter may be an important exposure pathway for filter feeders such as *A. stutchburyi* (King et al. 2004). As sediment Mn concentrations may be much higher due to the binding of Mn, this may explain the high Mn:Ca ratios observed in shells from this site (Hatje et al. 2003).

Future studies should further investigate the drivers of inter-site variation so as to provide further insights into the potential temporal and spatial stability of the elemental fingerprints observed. Understanding the conditions which cause variation in elements between sites will aid in the selection of study sites where trace elemental fingerprinting may prove successful in tracking larval dispersal. It should also be noted that mean Zn:Ca and Mn:Ca ratios in some instances were near or below the detection limits of the ICP-MS (Zn:Ca at Pataua, Mn:Ca at Marsden Cove and Mn:Ca at Waipu). Due to the high number of elements used in the multivariate analysis, however, this is unlikely to have influenced the validity of the results obtained.

CONCLUSIONS AND FUTURE DIRECTIONS

Overall, there was a high degree of spatial variation in the elemental fingerprints of *Austrovenus stutchburyi* over the small inter- and intra-estuary scales investigated in this study. This represents an important step in the use of trace elemental fingerprinting to unravel the local-scale connectivity patterns required for the effective management of bivalve species.

Prior to the application of elemental fingerprints to track the dispersal of marine larvae in future studies, the use of a pre-ablation spot which is larger than the analysed spot should be investigated to determine if edge effects influence results (Fowler et al. 2015). A careful comparison of the results obtained from shell cleaned using the pre-ablation technique used here

and other chemical cleaning techniques involving the use of hydrogen peroxide (e.g. Strasser et al. 2008) should also be undertaken to determine the effect of cleaning methodology. Furthermore, multiple sampling events should be incorporated to understand how elemental fingerprints change over seasonal or annual scales. Finally, future studies should examine if the ontogenetic and temporal intra-shell stability observed in settled shell of *A. stutchburyi* also extends to prodissoconch shell material. This will allow the comparison of prodissoconch and settled shell material to be made and allow inference of larval origins of settlers to validate biophysical models of larval dispersal in this area (Lundquist et al. 2009, Lundquist & Broekhuizen 2012).

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