

Patterns of regional endemism among New Zealand invertebrates

Briar Taylor-Smith, Mary Morgan-Richardson and Steven A. Trewick

Ecology, SAE, Massey University, Palmerston North, New Zealand

ABSTRACT

Biodiversity is unevenly distributed worldwide in terms of both species diversity and species endemism. Although centres of endemism are a conservation priority, both patterns and drivers of endemism are poorly understood in New Zealand. Here we explore whether invertebrate species distribution records in New Zealand represent the complete geographic range of species. We use distribution records of 2,322 invertebrate species to survey variation in range size and regional-endemism among 28 New Zealand regions, and explore the correlates of diversity and regional-endemism. Our data suggest patterns of regional-endemism in New Zealand invertebrates are not artefacts of sampling effort and the majority of species are not widespread. We found that endemism-score (which is a measure that corrects for species diversity) correlates positively with the relative size of the region three million years ago. Five variables (and their interactions) contributed to the relative level of invertebrate species endemism within a region (in a generalised linear model). Level of endemism tends to be lower in regions with greater geographic connectivity. This suggests that high levels of regional-endemism are not simply the product of the accumulation of species over time, but depends on the ability of a region to retain local species.

ARTICLE HISTORY

Received 10 April 2019
Accepted 14 October 2019

HANDLING EDITOR
Adrian Paterson

KEYWORDS

Diversity; endemism; extinction; geology; invertebrates of New Zealand; range size

Introduction

Diversity is unevenly distributed around the globe (Wallace 1879) with more variation in life existing at the tropics than in temperate regions (latitudinal biodiversity gradient; Dowle et al. 2013). Nevertheless, temperate New Zealand has a high proportion of species that are found here and nowhere else, thus it has high species endemism and is considered a biodiversity hotspot (Myers et al. 2000). Island Biogeography theory (MacArthur and Wilson 1967) might help explain the biodiversity in New Zealand, founded as it is on the principle that the number of species on an island is determined by an equilibrium between the rate of arrival of new species and the extinction of existing species. Distance from sources of colonists and island area are considered the main determinants. As an archipelago that is relatively large and distant from other landmasses (Neill and Trewick 2008)

and low extinction rate for New Zealand. As species can only be viewed as endemic if they occur nowhere else both extinction of source populations and in situ speciation have a role in development of New Zealand's highly endemic biota.

The same factors may influence species diversity at a larger scale and can be revealed by recording regional variation in diversity within a land mass such as New Zealand (e.g. Heenan et al 2017). Here we consider in particular regional endemism with a species regarded as a regional-endemic if it is restricted to a priori area within the landscape. Thus, the term regional-endemism is scale dependent. Our present focus on New Zealand invertebrates utilises published data where regions are already defined (see methods). The number of species recorded within a region and the number of these species that are restricted to it can be compared to similar sized regions to better understand the distribution.

- (3) Climate and habitat type influence endemism and species diversity in many ways. Slope, for example, may impact speciation rates because topographic heterogeneity results in habitat variation (Veech and Crist 2017) and more complex communities (Wollenberg et al. 2009), that may yield differential selection and high rates of speciation relative to extinction (species pump model, Moritz et al. 2000; Rahbek and Graves 2001). A complex environment with many biotic interactions might result in more species in the same geographic space compared to a simple community with few competitive interactions (species packing; MacArthur 1969). In Africa, endemism is concentrated in mountainous regions with low seasonality (Fjeldså and Jetz et al. 2004).
- (4) Climatic/habitat stability is likely to suppress rates of extinction (Wallace 1876; Dowle et al. 2013). In some places the distribution of plants and animals appears to have been influenced by glaciation and aridification during Pleistocene glacial periods. For example, a gradient of European endemism suggests refugia in warmer climates were influential in preserving local diversity (Essl et al. 2013).

Regional-endemism in New Zealand

Endemism appears to be unevenly graded across the New Zealand landscape (Wardle 1963; Rogers and Walker 2005; Millar et al. 2017) with three approximately coincidental regions of relatively high endemisms among vascular plants and invertebrates (Figure 1; Wardle 1963; Trewick et al. 2011). It is also notable that some taxa have disjunct distributions in regions of high endemism and absence between (Trewick and Wallis and Ereckal 2001; Wallis and Ereckal 1976; 38xima Tfnuneve

coverage of New Zealand natural history collection records might create arti

taxon dataset. We investigate potential drivers of New Zealand invertebrate regional-endemism by seeking correlations with environmental variables to infer the combination of environmental variables most associated with regional-endemism patterns. If the dominant influence on relative levels of regional-endemism in New Zealand is:

- (1) the equilibrium between origin (arrival and speciation) and extinction we expect a positive correlation between land area and regional endemism (measured by both the current and past size of the region).
- (2) the ability of species to expand or shift their ranges we expect to see a correlation between land connectivity and regional endemism.
- (3) species-packing and ecological speciation we expect to see a positive correlation between regional diversity of topography and climate and regional endemism.

Regional bias in extinction rates linked to environmental stability (e.g. climatic change and the extent of glaciers) is likely to be involved in determining relative levels of regional-endemism (4: habitat stability/longevity), but the size and heterogeneity of the 28 recognised regions considered in our analyses preclude a test for this.

Material and methods

Invertebrate diversity

Table 1. Invertebrates used to study regional-endemism in New Zealand. FNZ = Fauna of New Zealand series published by Landcare Research (*or other reference).

Class	Order	FNZ edition (common name)	Number of taxa
Insecta	Coleoptera	2: Osoriinae (Staphylinidae) (rove beetles)	37
		3: Anthribidae (fungus weevils)	35
		6: Hydraenidae (aquatic minute moss beetles)	32

evident when site sampling was abundant, we used two densely-sampled datasets for Simuliidae black flies (Insecta: Diptera; Craig et al. 2012) and Lycosidae wolf spiders (Arachnida: Araneae; Vin 2002).

Comparison of endemism with geography/environment

excluding low-lying land and marginal-marine deposits (Figure 2) (Trewick and Bland 2012). As a proxy for the extent of connectivity/isolation of each region we used the number of neighbouring regions currently sharing a boundary. For each region, measures of slope and elevation were included as proxies for habitat heterogeneity. Mean annual temperature and mean annual solar radiation were included because these variables might conceivably influence underlying mutation rates and speciation rate (Dowle et al. 2013). Mean monthly water balance ratio (MMWBR; the average of the monthly ratios of rainfall to potential evaporation) was used as a measure of water availability since aridity may have effects on patterns of regional-endemism, as observed for Madagascan fauna (Wilmé et al. 2006). This information was derived from the Ministry for the Environment data service website. The mean and variance of each region for each variable was calculated in ArcMap from ArcGIS 10.1 (ESRI, CA) using geographic regions (LENZ layer 'NZ Area Codes for recording specimen localities'). As a measure of habitat availability, we included prehuman forest cover, calculated from McGlone (1999) (and an estimate of current forest cover in each region (as a measure of the recent impact of human arrival). We used the relative human population size of each region and the presence of a relevant research institution in the region (a binary variable) as a proxy for sampling bias, because there might be a tendency for more thorough description of local versus more distant fauna. The relative population size of each region was calculated based on 2006 census data using the populated places polygon (by Peter Scott, from Koordinates.com).

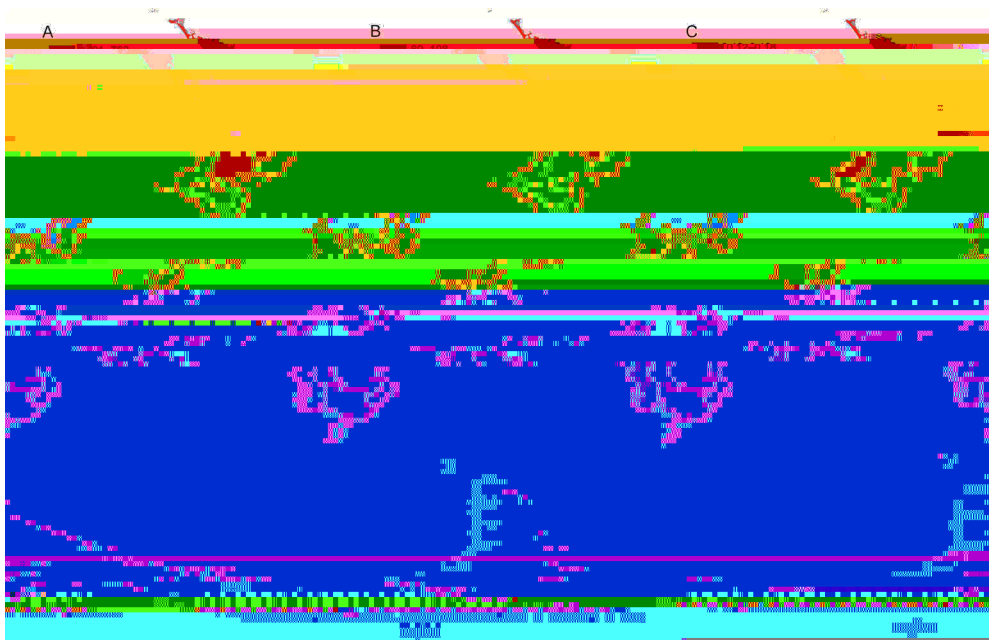


Figure 3. Endemism and diversity levels of invertebrate species are not homogeneous among regions of New Zealand. Four colours represent discretized (binned) numbers per region out of a sample of 2322 taxa. A) Total invertebrate diversity B) Number of endemic invertebrates C) Endemism score using the average level of endemism for each taxon. Bar plots illustrate ranked regional total diversity D) and ranked regional total number of endemic taxa (

regional-endemic taxa, but only 8.8% of the total land area of mainland New Zealand. In contrast, Hawke Bay (HB), Gisborne (GB), Wanganui (WI), Rangitikei (RI), Wairarapa (WA), Kaikoura (KA), north Canterbury (NC) and south Canterbury (SC) accounted for 23% of total land area but only 5% of all endemic taxa. No spatial autocorrelation was detected within the endemism and diversity indices ($P > 0.05$).

Sampling effort

To explore whether the data were dominated by sampling effort we excluded taxa recorded from only one site and that had poor overall sampling of the group. This reduced the number of regional-endemic taxa from 734 to 152 species. The strong relationship between the number of endemic taxa in each region and the number of well-sampled endemic taxa ($R^2 = 0.87$; $P < 0.001$; Figure 4), suggests that overall sampling is representative of actual levels of endemism within regions.

We explored further the influence of variation in sampling effort using patterns of endemism in two intensely sampled families; Simuliidae and Lycosidae spiders (Table 2). Within each region, every collection site provided independent presence/absence records for each of the 11 Simuliidae and 18 Lycosidae species endemic to North Island and/or South Island New Zealand. In both families, most species were recorded from few regions while few species were widespread, in keeping with the predominant pattern of range restriction seen in the larger invertebrate data set (Figure 5).



Figure 4.(A)

Although intensely sampled, neither the Simuliidae nor Lycosidae comprised species diversity to test for significant heterogeneity among regions.

Endemism and environment

The relationship between endemism and the environment was tested using a set of environmental variables (Table S1). There were no simple correlations between endemism or diversity and the environmental variables, however, a significant positive correlation (after Bonferroni correction) was observed between endemism-score (which is a measure that corrects for species diversity) and land area 3 million years ago (34; $P=0.001$; Figure D). Variation in levels of regional-endemism among the regions was best explained by multiple variables in the general linear model (Table 3). Total species diversity ($t = 6.20$; $P=0.00001$), land area 3 Ma ($t=7.45$; $P<0.00001$) and the presence of a research institution ($t = 7.32$; $P<0.00001$) each improved the model, having a significant positive impact on regional-endemism. Climatic variables (represented in this model by the ratio of rainfall to evaporation and seasonal sunshine variation) also contributed to regional-endemicity, and each correlated with other environmental variables not in the model (Table 3 and S2). Interestingly, variance in solar radiation and water availability

(MMWBR) interacted to have a significant negative effect ($t = -3.49; P = 0.003$). The effect of total species diversity and number of neighbouring regions interacted with one another to negatively affect regional-endemism ($t = -3.11; P = 0.007$). Two other variables included in the model, mean slope and current forest cover, did not have a significant effect ($t < 1.25; P > 0.05$).

Discussion

The patterns

The same pattern of regional variation was apparent in our dataset of endemic invert-

sampled families of New Zealand invertebrates, Lycosidae spiders and Simuliidae black flies, provided information on absences and were not biased by sampling. Within these families, species distributions were heterogeneous with few widespread and many restricted species, as seen in the larger invertebrate dataset (Figure 5). We are confident that the distributions of invertebrate species, as recorded in the Fauna of New Zealand series, do contribute meaningful data for analysing patterns of New Zealand regional-endemism.

As observed in plants (Wardle 1963; McGlone et al 2001; Heenan et al 2017), levels of invertebrate endemism are not homogeneous among New Zealand regions. Rather, some regions have high endemism while others had few or no endemic invertebrate taxa in our sample (Figure 3).

invertebrate species diversity was positively correlated with endemism in New Zealand (Figure 4). However, species diversity interacted with number of connecting (neighbour-

et al. 2018. Although we have evidence that climate and habitat may influence relative levels of invertebrate species endemism the correlation of variables and heterogeneity of regions makes isolation of the relevant drivers difficult.

Climatic and habitat stability

The distribution of plants and animals appears to have been influenced in some places by glaciation and aridification during Pleistocene glacial periods. For example, a gradient in levels of European endemism suggests refugia in warmer climates were essential in preserving local diversity (Essl et al. 2013). During New Zealand's LGM (Williams et al. 2015

Dn

of isolates T331.491 (EsTJ .6 (warmtionr0.9(an)]gnt)s)-339.5 (and) could 0.2 (in)-po

Conclusions

Small-ranged species are a conservation priority, therefore, understanding current patterns of regional-endemism in New Zealand is important for protection of biodiversity. We investigated regional-endemism using a rich invertebrate dataset and found that although variation in sampling effort does influence the data, most species are range restricted and endemism levels are not evenly distributed across the country. High invertebrate species diversity and high endemism can both be conserved if Northland and Nelson are provided with resources to protect their natural habitats. No single variable appears to drive variation in regional-endemism. Those variables that were important appear to be related to differences in geologically recent land availability, and connectivity between regions. High levels of endemism in some regions are not simply a product of the accumulation of species over time, rather regional-endemism depends on the ability of a region to retain local species. Regions with fewer neighbours have lower migration rates, whereas those with many neighbours are less able to retain endemic taxa. The absence of land in southern North Island until recent geological time shows that tectonic activity in New Zealand continues to have a significant influence on current endemism levels.

Acknowledgements

Matt Irwin, Lesley Essen and Jean Sanderson helped organise data. The research was aided by a Massey University Research Fund grant to M.M.R. and scholarships to B.T.S. from Massey University, the Brian Mason Scientific & Technical Trust, and the NZ Entomological Society. We appreciate the support of the Phoenix group (www.evolve.massey.ac.nz). The manuscript was improved by constructive criticism from Matt McGlone and two anonymous reviewers.

Disclosure statement

No potential conflict of interest was reported by the authors.

Funding

This work was supported by Massey University under Grant MURF 2014 & Doctoral scholarship to BTS; Brian Mason Scientific & Technical Trust and New Zealand Entomological Society.

Data availability statement

Data associated with the paper from Fauna of New Zealand <https://www.landcareresearch.co.nz/publications/books/fauna-of-nz-series>

ORCID

Mary Morgan-Richard 

- Anderson S. 1994 Area and endemism. *Quarterly Review of Biology*. 69:471–481.
- Birand A, Vose A, Gavrillets S. 2012 Patterns of species ranges, speciation, and extinction. *The American Naturalist*. 179:21.
- Bulgarella M, Trewick S, Minards NA, Jacobson MJ, Morgan-Richards M, Kissling M. 2019 Shifting ranges of two tree weta species (*Hemideina* spp.): competitive exclusion and changing climate. *Journal of Biogeography*. 41(3):525–536.
- Craig DA, Craig REG, Crosby TK. 2012 Simuliidae (Insecta: Diptera). *Fauna of New Zealand*, Vol. 68. Lincoln: Manaaki Whenua Press.
- Crampton JS, Cooper R, Beu AG, Foote M, Marshall BA. 2010 Biotic influences on species duration: interactions between traits in marine molluscs. *Paleobiology*. 38:204–214.
- Crisp MD, Laan SW, Linder HP, Monro A. 2001 Endemism in the Australiana. *Journal of Biogeography*. 28:189–198.
- Crosby TK, Dugdale JS, Watt J. 1976 Recording specimen localities in New Zealand: An arbitrary system of areas and codes. *New Zealand Journal of Zoology*. 3:69.
- Crosby TK, Dugdale JS, Watt J. 1998 Area codes for recording specimen localities in the New Zealand subregion. *New Zealand Journal of Zoology*. 25:135–137.
- Daru BH, Bank M, Davies TJ, Thuiller W. 2014 Spatial incongruence among hotspots and complementary areas of tree diversity in southern Africa. *Diversity and Distributions*. 21:786–799.
- Don W. 2007 *Ants of New Zealand*. Dunedin: Otago University Press.
- Dowle E, Morgan-Richards M, Trewick S. 2013 Molecular evolution and the latitudinal biodiversity gradient. *Heredity*. 110:561–570.
- Essl F, Moser D, Dirnböck T, Dullinger S, Milasowsky N, Winter M, Rabitsch A. 2013 Native, alien, endemic, threatened, and extinct species diversity in European countries. *Biological Conservation*. 164:907–917.
- Fjeldså J, Ehrlich D, Lambin E, Prins L. 1997 Are biodiversity hotspots correlated with current ecoclimatic stability? A pilot study using the NOAA-AVHRR remote sensing data. *Biodiversity & Conservation*. 6:404–422.
- Foote M, Crampton JS, Beu AG, Marshall BA, Cooper RA, Maxwell PA, Matczak J. 2007 Rise and fall of species occupancy in Cenozoic fossil mollusks. *Science*. 318:1331–1334.
- Goldberg J, Knapp M, Emberson RM, Townsend JL, Trewick SA, Colwell RK. 2013 Species radiation of Carabid beetles (*Broscini*: *Mecodema*) in New Zealand. *PLoS ONE*. 9(1):e86185.
- Heads M. 1997 Regional patterns of biodiversity in New Zealand: one degree grid analysis of plant and animal distributions. *Journal of the Royal Society of New Zealand*. 27:337–347.
- Heenan PB, Millar TR, Smitsen RD, McGlone MS, Wilton DJ. 2017 Phylogenetic measures of neo- and palaeo-endemism in the indigenous vascular flora of the New Zealand archipelago. *Australian Systematic Botany*. 30:423–433.
- Jetz W, Rahbek C, Colwell R. 2004 The coincidence of rarity and richness and the potential signature of history in centres of endemism. *Ecology Letters*. 7:1190–1200.
- Kisel Y, Barraclough TG. 2010 Speciation has a spatial scale that depends on levels of gene flow. *The American Naturalist*. 175:343–354.
- MacArthur RH. 1969 Species packing, and what interspecies competition minimizes. *Proceedings of the National Academy of Sciences of the United States of America*. 64:318–321.
- MacArthur RH, Wilson EO. 1967 *The theory of island biogeography*. Princeton, NJ: Princeton University Press.
- Martin RD. 1993 Primate origins Plugging the gaps. *Nature*. 363:223–224.
- McGlone M. 1989 The Polynesian settlement of New Zealand in relation to environmental and biotic changes. *New Zealand Journal of Ecology*. 12:129–135.
- McGlone MS, Duncan R, Heenan PB. 2001 Endemism, species selection and the origin and distribution of the vascular flora of New Zealand. *Journal of Biogeography*. 28:296–306.
- McGlone MS, Newnham RM, Moar N. 2010 The vegetation cover of New Zealand during the last glacial maximum: do pollen records under-represent woody vegetation. *Terra Australis*. 32:49–68.

- Millar TR, Heenan PB, Wilton AD, Smissen RD, Breitwieser A. 2017. Spatial distribution of species, genus and phylogenetic endemism in the vascular flora of New Zealand, and implications for conservation. *Australian Systematic Botany*. 30:147.
- Moran PAP. 1950. Notes on continuous stochastic phenomena. *Biometrika*. 37:17.
- Morgan-Richards M, Bulgarella M, Sivyler L, Dowle EJ, Hale M, McKean NE, Trewick SA. 2015. Explaining large mitochondrial sequence differences within a population sample. *Royal Society Open Science*. 4(11):170730.
- Moritz C, Patton JL, Schneider CJ, Smith AB. 2000. Diversification of rainforest faunas: an integrated molecular approach. *Annual Review of Ecology and Systematics*. 31:533.
- Myers N, Mittermeier RA, Mittermeier CG, Da Fonseca GA, Kesteven J. 2000. Biodiversity hotspots for conservation priorities. *Nature*. 403:853.
- Nattier R, Grandcolas P, Elias M, Desutter-Grandcolas L, Jourdan H, Couloux A, Robillard T, O'Grady P. 2012. Secondary sympatry caused by range expansion informs on the dynamics of microendemism in a biodiversity hotspot. *PLoS ONE*. 7(11):e48047.
- Neall VE, Trewick SA. 2008. The age and origin of the Pacific Islands – a geological overview. *Philosophical Transactions of the Royal Society, London*. 363:3293.
- Papadopoulou A, Anastasiou I, Keskin B, Vogler AP. 2009. Comparative phylogeography of tenebrionid beetles in the Aegean archipelago: the effect of dispersal ability and habitat preference. *Molecular Ecology*. 18:2505.
- Rahbek C, Graves GR. 2001. Multiscale assessment of patterns of avian species richness. *Proceedings of the National Academy of Sciences*. 98(8):4539.
- R Development Core Team. 2010. R: a language and environment for statistical computing. Vienna, Austria: R Foundation for Statistical Computing. [URL: www.R-project.org/](http://www.R-project.org/)
- Ribera I, Barraclough T, Vogler AP. 2001. The effect of habitat type on speciation rates and range movements in aquatic beetles: inferences from species-level phylogenies. *Molecular Ecology*. 10:724-735.
- Rogers GM. 1989. The nature of the lower North Island floristic gap. *New Zealand Journal of Botany*. 27:22-241.
- Rogers G, Walker L. 2005. Evolution of the New Zealand vascular flora: regional and provincial patterns of richness, radiation, and endemism. *New Zealand Journal of Botany*. 43:381.
- Shepherd LD, Perrie LR, Brownsey P. 2007. Fire and ice: volcanic and glacial impacts on the phylogeography of the New Zealand forest fern *Asplenium hookerianum*. *Molecular Ecology*. 16:4536-4549.
- Sivyler L, Morgan-Richards M, Koot E, Trewick SA, Keyghobadi N. 2008. Anthropogenic cause of range shifts and gene flow between two grasshopper species revealed by environmental modeling, geometric morphometrics and population genetics. *Insect Conservation and Diversity*. 1(5):415-434.

j K 3 3 6 0 1 5 2 3 2 2 8 1 2 2 0 0 5 2 8 3 0 7 2 5 5 8 0 0 6 2 6 0 - 7 . 2 (7) T J 0 1 0 s c n 1 . 9 2 2 0 T d [(.) - 2 5 9 . 4 (F i r e) -

- Vink CJ. 2002 Lycosidae (Arachnida: Araneae). Fauna of New Zealand, Vol. 44. Lincoln: Manaaki Whenua Press.
- Wallace AR. 1876 The geographical distribution of animals. London: MacMillan.
- Wallis GP, Trewick SA. 2009 New Zealand phylogeography: evolution on a small continent. *Molecular Ecology*. 18:3543-3580.
- Wallis GP, Waters JM, Upton P, Craw D. 2016 Transverse alpine speciation driven by glaciation. *Trends in Ecology and Evolution*. 31:992-1006.
- Ward DF. 2012 More than just records: analysing natural history collections for biodiversity planning. *PLoS One*. 7(11):e50346.
- Wardle P. 1963 Evolution and distribution of the New Zealand flora, as affected by Quaternary climates. *New Zealand Journal of Botany*. 1:1-173.
- Waters JM, Fraser CI, Hewitt GM. 2013 Founder takes all: density-dependent processes structure biodiversity. *Trends in Ecology & Evolution*. 28:77-88.
- Webb TJ, Gaston KJ. 2003 On the heritability of geographic range sizes. *The American Naturalist*. 161:553-566.
- Williams PW, McGlone M, Neil H, Zhao J. 2015 A review of New Zealand palaeoclimate from the