Supplementary Material

Appendix S1. R packages used in the analyses.

Library	Version	Citation
General wrangling		
tidyverse	2.0.0	Wickham H, Averick M, Bryan J, Chang W, McGowan LD, François R, Grolemund G, Hayes A, Henry L, Hester J, Kuhn M, Pedersen TL, Miller E, Bache SM, Müller K, Ooms J, Robinson D, Seidel DP, Spinu V, Takahashi K, Vaughan D, Wilke C, Woo K, Yutani H (2019). Welcome to the tidyverse. Journal of Open Source Software, 4, 1686. <https: 10.21105="" doi.org="" joss.01686="">.</https:>
widyr	0.1.5	Robinson D, Silge J (2022). widyr: Widen, Process, then Re-Tidy Data. R package version 0.1.5, https://CRAN.R-project.org/package=widyr .
Scopus API		
rscopus	0.7.1	Muschelli J (2023). rscopus: Scopus Database 'API' Interface. https://dev.elsevier.com/sc_apis.html, https://github.com/muschellij2/rscopus.
Networks		
circlize	0.4.15	Gu Z, Gu L, Eils R, Schlesner M, Brors B (2014). circlize Implements and enhances circular visualization in R. Bioinformatics 30(19): 2811–2812.
igraph	1.5.1	Csardi, G., & Nepusz, T. (2006). The igraph software package for complex network research. InterJournal, Complex Systems, 1695.
network	1.18.1	Butts C (2008). network: a package for managing relational data in R. Journal of Statistical Software, 24(2). https://doi.org/10.18637/jss.v024.i02 >.
RСу3	2.20.2	Gustavsen, J. A., Pai, S., Isserlin, R., Demchak, B., & Pico, A. R. (2019). RCy3: Network biology using Cytoscape from within R. F1000Research, 8, 1774. https://doi.org/10.12688/f1000research.20887.3

ggraph	2.1.0	Pedersen T (2022)ggraph: An Implementation of Grammar of Graphics for Graphs and Networks R package version 2.1.0, .
tnet	3.0.16	Opsahl, T., 2009. Structure and evolution of weighted networks. University of London (Queen Mary College), London, UK, pp. 104-122. Available at http://toreopsahl.com/publications/thesis/; http://toreopsahl.com/tnet/
Text handling & modelling		
tidytext	0.4.1	Silge, J., & Robinson, D. (2016). Tidytext: Text mining and analysis using tidy data principles in R. The Journal of Open Source Software, 1(3), 37. https://doi.org/10/gfwcjt
stm	1.3.6.1	Roberts, M. E., Stewart, B. M., & Tingley, D. (2019). stm: An R package for structural topic models. Journal of Statistical Software, 91(2). < <u>https://doi.org/10.18637/jss.v091.i02</u> >
SnowballC	0.7.1	Bouchet-Valat M (2023). SnowballC: Snowball Stemmers Based on the C 'libstemmer' UTF-8 Library. R package version 0.7.1, ">https://CRAN.R-project.org/package=SnowballC> .
tm	0.7.11	Feinerer I, Hornik K, Meyer D (2008). Text mining infrastructure in R. Journal of Statistical Software 25 (5), 1-54. https://doi.org/10.18637/jss.v025.i05 >.
franc	1.14	Csardi G, Wormer T, Ceglowski M, Rideout JR, Johnson AKS (2021). franc: Detect the Language of Text. R package version 1.1.4, https://CRAN.R-project.org/package=franc >.
word2vec	0.3.4	Wijffels J (2021)word2vec: Distributed Representations of Words. R package version 0.3.4, <https: cran.r-project.org="" package="word2vec">.</https:>

textstem	0.1.4	Rinker, T. W. (2018). textstem: Tools for stemming and lemmatizing text version 0.1.4. Buffalo, New York. http://github.com/trinker/textstem
wordcloud		Fellows I (2018)wordcloud: word clouds R package version 2.6, <https: cran.r-<br="">project.org/package=wordcloud>.</https:>
textclean	0.9.3	Rinker, T. W. (2018). textclean: Text Cleaning Tools version 0.9.3. Buffalo, New York. https://github.com/trinker/textclean
Other		
densityClust	0.3.2	Pedersen T, Hughes S, Qiu X (2022). densityClust: clustering by fast search and find of density peaks. R package version 0.3.2, https://CRAN.R-project.org/package=densityClust .

Appendix S2. Custom stopwords

For the STMs we used the Snowball stopwords: "new", "zealand", "zealand's", "south", "north", "australia", "australian", "copyright", "sp", "sp.", "nov.", "found", "increase", "suggest", "low", "ii", "iii", "iv", "NA", "time", "female", "male", "study", "studied", "studies", "result", "size", "species", "rate", "difference", "effect", "effects", "differences", "females", "males", "significant", "significantly", "results", "test", "increase", "sample", "tests", "tested", "samples", "sampled", "increases", "increased", "similar", "occur", "site", "suggest", "include"

Appendix 3. Interactive network figures

Interactive versions of some of the network figures (Figs. 4,5, and 7) are available as follows. Note that these interactive versions include more information than those in the printed version although the underlying data are the same; in other words, they are not intended to be exact replicas. The interactive versions were made using Cytoscape and the rcytoscape packages (Shannon et al. 2003).

- Figure 4 affiliations networks are available at <<u>https://spatialecol.com/presentations/cyto/affiliations/web_session/index.html#/</u>>, set the visual style on the tab to **AffiliationsCirc**
- Figure 5 co-author networks are available at <<u>https://spatialecol.com/presentations/cyto/coauthors/web_session/index.html#/</u>>, set the visual style on the tab to **Coauthors**
- Figure 7 keyword co-occurrence networks are available at <<u>https://spatialecol.com/presentations/cyto/keywords/web_session/index.html#/</u>>, set the visual style on the tab to **Keywords**



Organisational group — CRIs — Govt — University

Appendix S4. Number of records associated with universities, CRIs, and central government in NZ. Upper columns show the absolute count (each author on a paper gets a score of 1) while the lower show the fractional count (each author on a paper gets a score of 1/*n*, where *n* is the number of co-authors). Vertical lines are the formation of CRIs in 1992 and the start of the PBRF in 2004



Appendix S5. Cumulative plots of (A) number of unique journals published in over time, (B) number of unique author affiliations, (C) number of unique countries, and (D) number of unique authors.



Appendix S6. Summary of the topics from the *k* = 24 structural topic models with year, NZ journal, and their covariates for (A) publications with NZ-affiliated authors (B) publications on a NZ topic, and (C) all publications (A and B combined). Colours denote topic prevalence across the period 1980–2020, labels are the top ranked terms based on probability (b) for each topic.

Topic 1		Topic 2		Topic 3		Topic 4		
pmax	temperature	simco	carbon	borak	naturalise	tree line	forest	
bait hook	lichen	trossulus	mussel	iacaena	subsp	silver beech	fire	
hypoliths	photosynthesis	ocean biogeochemical	isotope	urvilliana	genus	natural seedling	vegetation	
desiccation prone	photosynthetic	downscales	stable isotope	linatella	kermadec island	goshawk	seedling	
rosshavet	coral	bcasts	lipid	triton	name	wildlings	beech	
upper shade	irradiance	tariff	fatty	obconica	comb	aranuian	canopy	
sunshade	alga	carbohydratefat	fatty acid	mordella	synonymy	maquis	tree	
temperature	bacterial community	carbon	isotope	island	ngen	forest	litterfall	
water	conductance	food	stable isotope	record	basionyms	tree	mountain beech	
light	photosynthetic rate	diet	isotope analysis	genus	nomenclatural status	vegetation	tree line	
warm	fucoid	mussel	mytilus	taxon	electronic flora	native	silver beech	
growth	spin-dry valley	energy	galloprovincialis	endemic	name database	plant	charcoal	
community	elevate temperature	Isotope	discount rate	plant	tipu	cover	weinmannia racemosa	
climate	galloway	acid	stable carbon	distribution	tipu aotearoa	site	forest structure	
Тор	pic 5	Topic 6		Тор	Topic 7		Topic 8	
pindope	lamb	polyphylla	invasion	hfes	genetic	byena	community	
oxide technique	possum	channel catfish	alien	grassy community	haplotype	multilaver network	richness	
clover stag	liveweight	machair	native	fernandez firecrown	locus	technological progress	ecosystem	
leaf allowance	milk	pest risk	invasive	avsc versus	genetic diversity	metallophytes	species richness	
lean liveweight	bait	archaeophytes	alien species	river capture	allele	mountain lion	ecosystem service	
winter lean	wean	native deer	invasive species	otsclock1b	microsatellite	ecosystem size	trait	
weight line	heifer	propagule size	native range	ots5a5nwfsc	mitochondrial	kowaro	stressor	
control	lamb	native	alien species	population	haplotype	community	functional richness	
possum	liveweight	invasive	native range	genetic	genetic diversity	ecosystem	natural capital	
bait	heifer	invasion	biological invasion	island	allele	trait	social ecological	
lamb	romney	plant	alien plant	diversity	genetic variation	diversity	stressor effect	
weight	slaughter	introduce	invasive plant	gene	genetic structure	scale	assembly process	
treatment	carcass weight	range	introduce range	variation	mtdna	functional	ecosystem size	
miik	Tieece	Introduction	plant invasion	structure	genetic differentiation	environmentai	alternative stable	
Тор	Topic 9		Topic 10		Topic 11		Topic 12	
eradication effort	eradication	node appearance	clover	sphenodonti	parasite	sclerotia	endophyte	
eradication success	antarctic	pitau	rvegrass	metacercariae	host	tuki	radiata	
eradication campaign	ross	sugar yield	white clover	gnathiid	trematode	grain angle	fungus	
astrolabe	rodent	maku	cultivar	shoal size	snail	holopsis	mycorrhizal	
pseudoeconesus	gracilipes	regenerants	sward	gallinarum	antipodarum	cinnamomi	pinus radiata	
gracilipes	expedition	scheuchzeri	tiller	cobble size	parasite species	simile	pinus	
cadophora	rodent eradication	unattacked	herbage	arboreal plant	host species	stem wood	wood	
island	gracilipes	plant	tiller	host	parasite	tree	mycorrhizal	
eradication	rodent eradication	clover	clover trifolium	parasite	trematode	wood	phytophthora	
antarctic	eradication program	root	lotus	snail	flea	radiata	ddon	
rodent	rena	growth	trifolium repens	population	parasite species	fungus	radiata ddon	
ross	grandis	yield	stoion	Intection	nost parasite	plant	wood property	
hait	invasive rodent	loof	sood vield	individual	intermediate host	arowth	mycorrhizal fungus	
Dait	invasive rouent	ical	seed yield	individual	antermediate nost	giowai	mycormizariungus	

Metric a FREX a Lift a Prob (β) a Score

Appendix S7. Labels for 1–12 of the 24 topics identified by the STM. The purple boxes show the terms with the highest probability in the identification of the topics (b). The other metrics are FREX (terms that are frequent and exclusive), the term-lift metric (Taddy 2012), and the score metric (Chang 2015).

Topic 13		Topic 14		Topic 15		Topic 16		
crambina	clade	freshwater megafauna	management	world country	prev	lobster	fish	
nabkha	lineage	badge	conservation	ladder	spider	inanga	trout	
diadema	genus	buckwheat	policy	gelotia	salticids	trout salmo	spawn	
herbertus	seta	cage clump	biodiversity	icdd	salticid	jasus edwardsii	lobster	
awaous	vicariance	wood rise	threaten	display posture	portia	salmo trutta	reef	
hamuli	gondwana	cache space	threaten species	dawn chorus	predator	trout	otolith	
afro	monophyletic	theft	urban	insect biomass	courtship	metamorphosis	galaxias	
genus	electron microscope	management	conservation action	prey	salticids	fish	trout	
island	rygmodus	conservation	policy	predator	salticid	larva	lobster	
sequence	axial parenchyma	research	people	spider	portia	habitat	otolith	
analysis	nrdna	impact	decision maker	behaviour	jump spider	adult	reef fish	
taxon	species delimitation	risk	conservation	display	vibratory	larval	kokopu	
southern	nabkha	approach	management	response	salticidae	juvenile	brown trout	
region	tree topology	biodiversity	stakeholder	nest	silk	density	rainbow trout	
То	Topic 17		Topic 18		Topic 19		Topic 20	
sorption	soil	ecosanctuaries	teote	mistletoe	flower	thief	penguin	
irrigator	fertiliser	beech mast	dolphin	pollination service	pollinator	fast sperm	mate	
biochar	pasture	wasp vespula	skink	alepis	pollen	parental effort	chick	
post confluence	urine	common wasp	rattus	enantiostvlv	pollination	food input	sperm	
hawkweed cover	lime	vespula	prey	kamchatica	mistletoe	input ratio	breed	
application depth	olsen	laysan	wasp	sunbird	seed	hare wallaby	fledge	
effluent irrigation	superphosphate	mahout	home range	herkogamy	fruit	host coral	colony	
soil	lime	island	vespula	flower	mistletoe	breed	nestle	
pasture	pasture production	population	common wasp	plant	tetrapetala	mate	extrapair	
concentration	potassium	predator	german wasp	pollen	stigma	population	sperm competition	
water	olsen	nest	bottlenose	pollination	peraxilla	colony	skua	
graze	soil water	bird	bottlenose dolphin	fruit	ovule	individual	ejaculate	
fertiliser	leach loss	habitat	mohua	pollinator	pollination service	condition	foal	
application	lysimeter	density	artificial nest	seed	alepis	forage	harem	
То	pic 21	Topic 22		Topic 23		Topic 24		
vbnc	sediment	vertical object	pheromone	final sweep	simulation	germination	seed	
sediment concentration	stream	pheromone titre	acetate	enfa model	model	brigalow	germination	
terrestrial sediment	lake	paludosa	apple	unweighted regression	spatial	subtorquata	fruit	
cdom	periphyton	dead seed	trap	temporary migration	error	seed bank	germinate	
calamoecia	phytoplankton	asubunit isoform	edna	bien database	accuracy	tetrapathaea	seedling	
froude	benthic	hypanthial	moth	master sample	sdms	bilge	dispersal	
aphanizomenon	macroinvertebrate	zeac	volatile	gcgms	prediction	scatterhoarders	broom	
water	sediment	trap	acetate	model	sdms	seed	germination	
stream	macropnyte	pheromone	polymerase	datum	tree model	fruit	seed dispersal	
sediment	periphyton	sequence	postvittana	estimate	monte	dispersal	broom	
river	macroinvertebrate community	compound	polymerase chain	predict	sensitivity analysis	germination	seed bank	
flow	suspend sediment	extract	tran bait	analysis	monte carlo	piant	frugivore	
community	periphyton biomass	identify	ctenopseustis	scale	neural network	seedling	frugivores	
connunty	periphyton biomuss	lucitury	cichopsousis	State	Heural Hetwork	seconing	indgivores	

Metric a FREX a Lift a Prob (β) a Score

Appendix S8. Labels for 13–24 of the 24 topics identified by the STM. The purple boxes show the probability weight used to identify the topics (b). The other metrics are FREX (terms that are frequent and exclusive), the term-lift metric (Taddy 2012), and the score metric (Chang 2015).



Appendix S9. Number of abstracts with either of the terms Māori or mātauranga appearing in them over time.