

Ectomycorrhizal fungi diversity associated with small mammals in response to urbanisation

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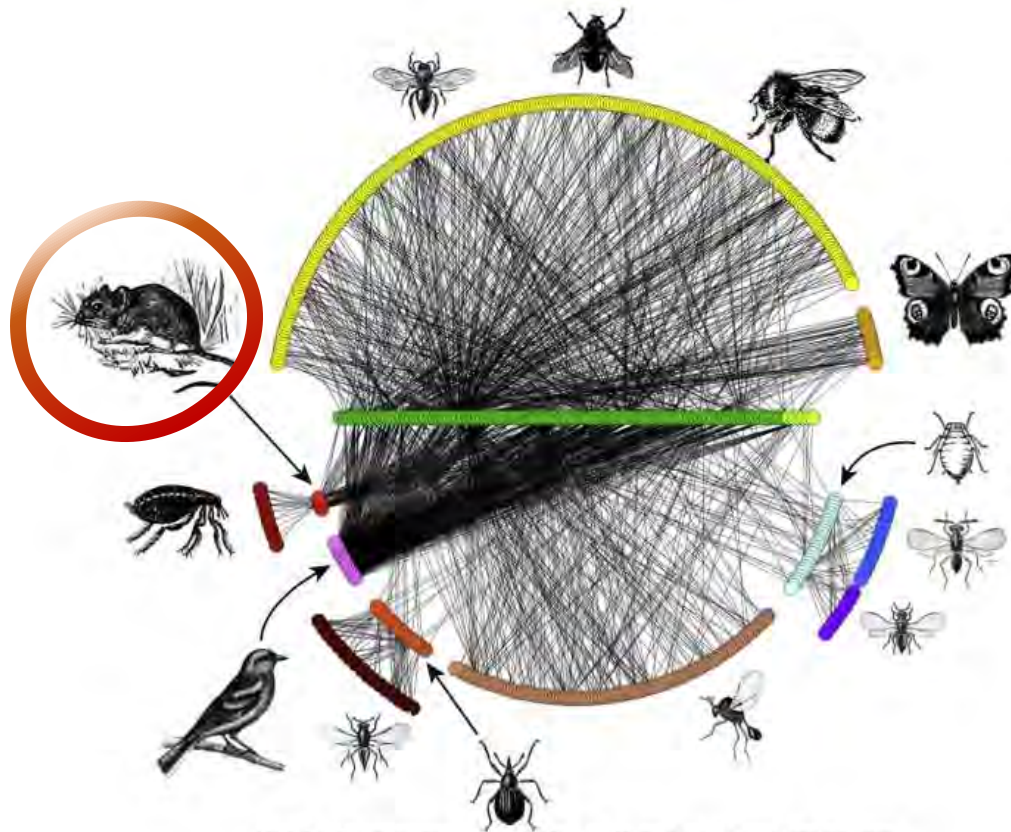
Overview

Integrative ecology of small mammals in response to urbanisation

Ectomycorrhizal fungi diversity associated with small mammals in response to urbanisation

Small mammal behaviour in response to urbanisation

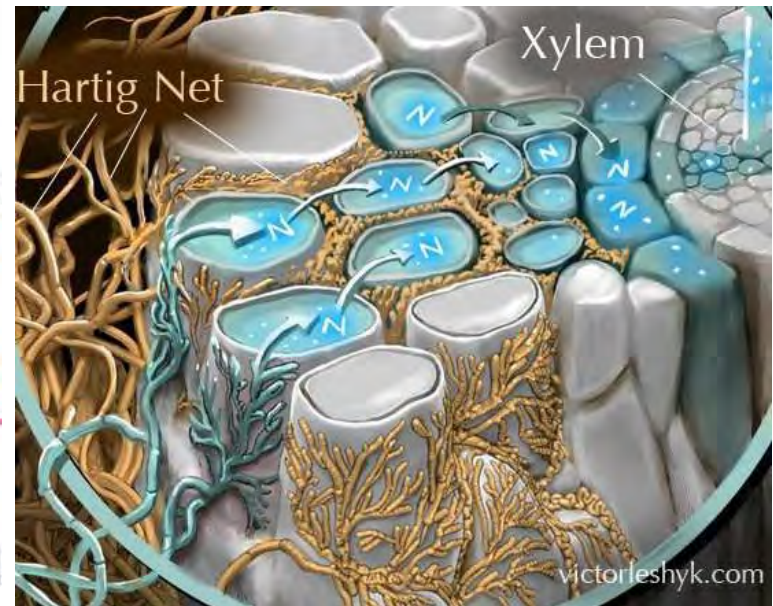
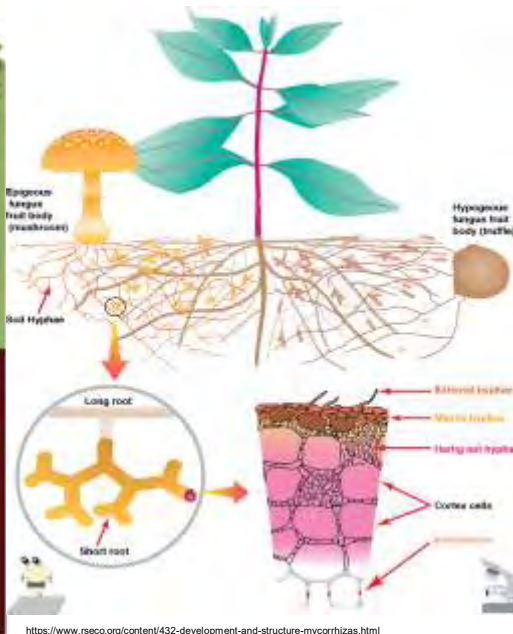
We are all interacting!



Credit: Quintessence Network DOI: 10.1016/j.tree.2015.12.003

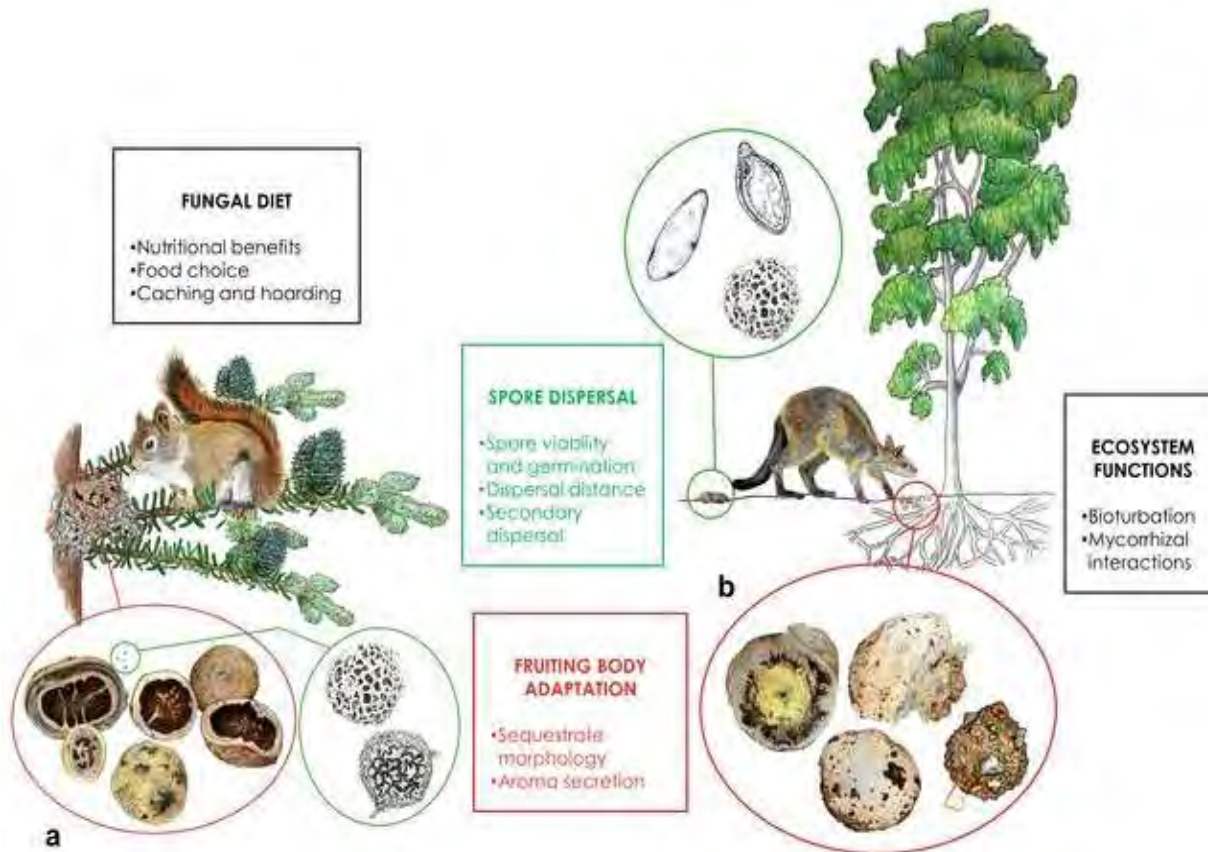
An outstanding set of interactions

- 90% of plants depend on mycorrhizal fungi
- Ectomycorrhizal fungi (EMF) are associated with woody plants
- EMF diversity is considered a strong indicator of forest health.

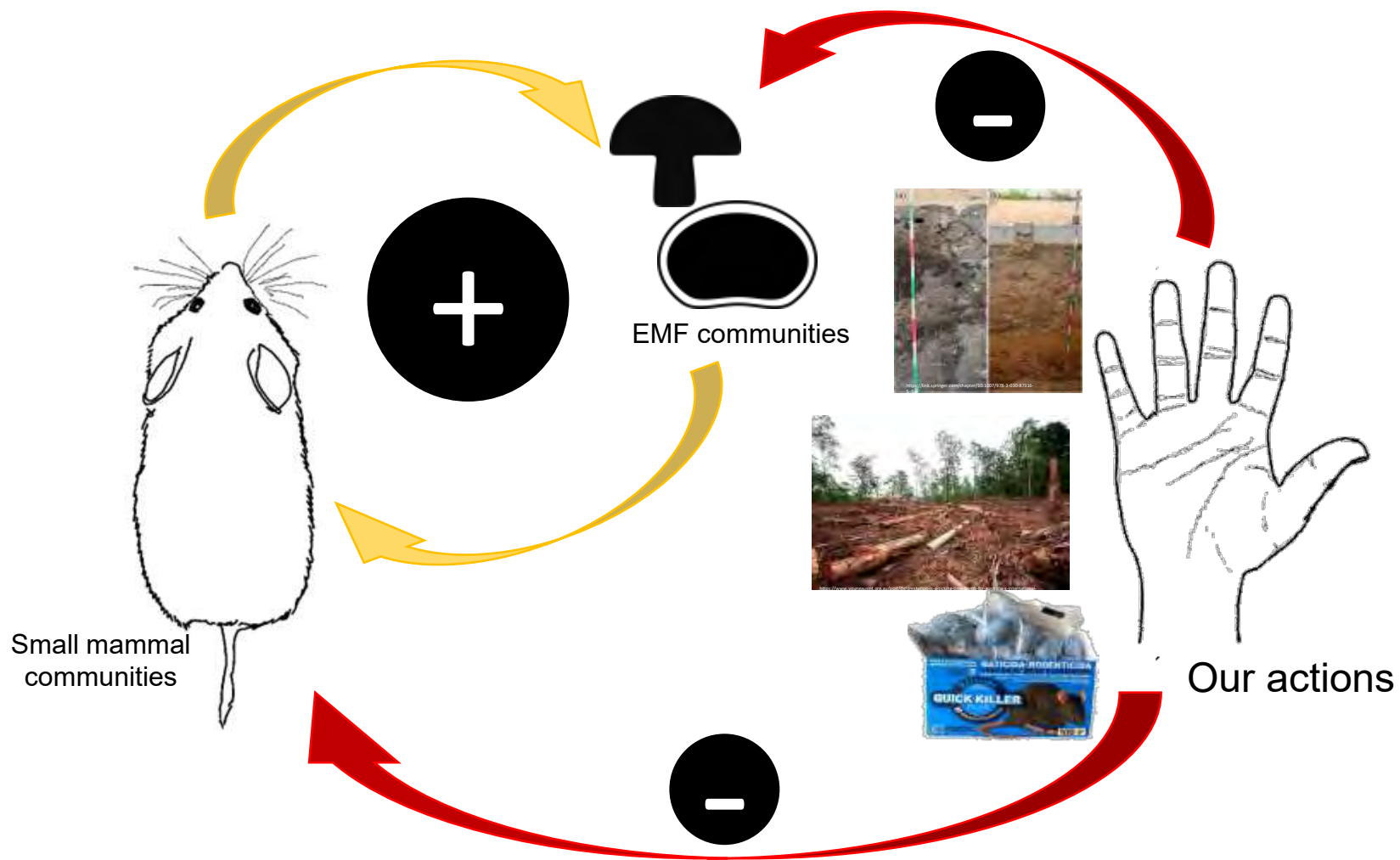


An outstanding set of interaction

- Small mammals are well-known fungivores
- Small mammals help in the dispersal of EMF spores

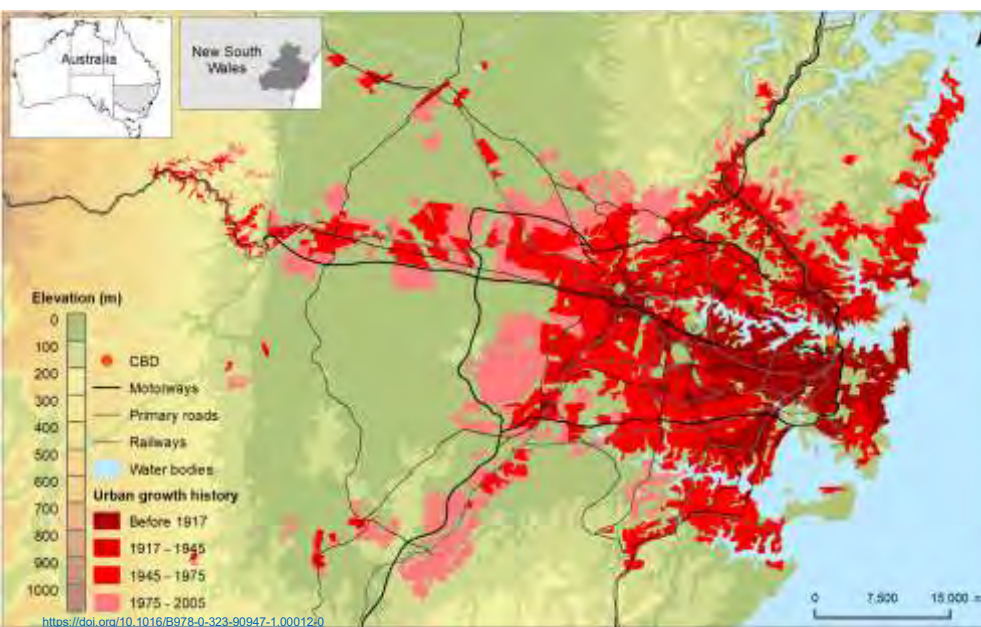


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Urbanisation

- One of the most prevalent and detrimental anthropogenic disturbances to forest ecosystems globally
- Some EMF and small mammals thrive in urban settings, but they face many challenges
- In this context, urban reserves represent a refuge from these challenges

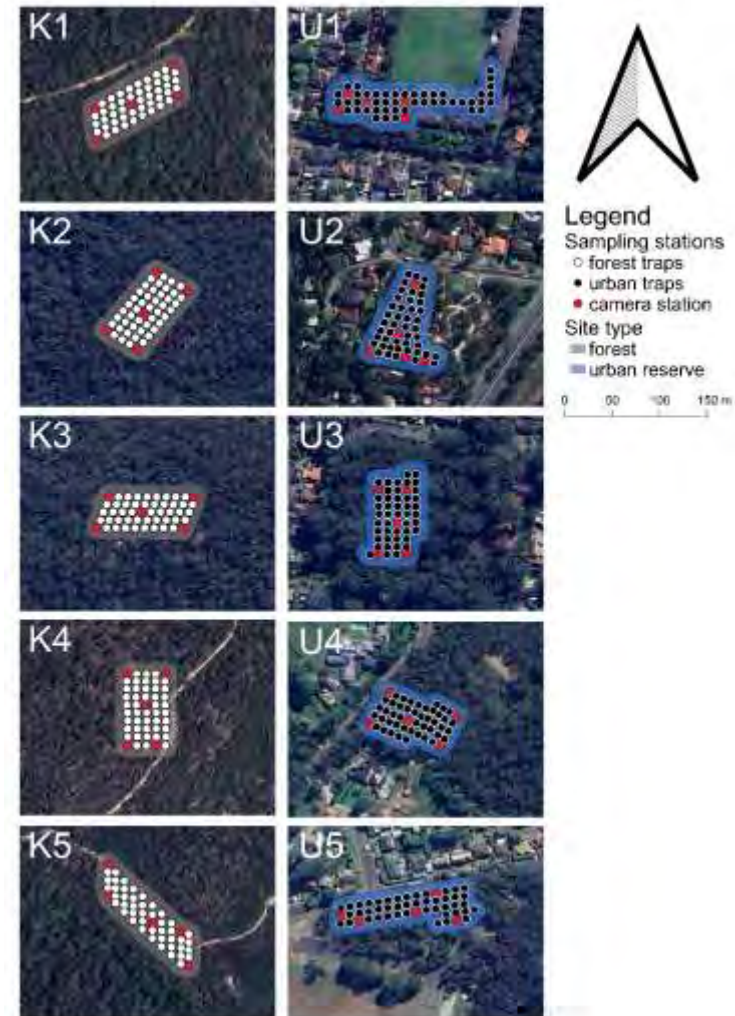
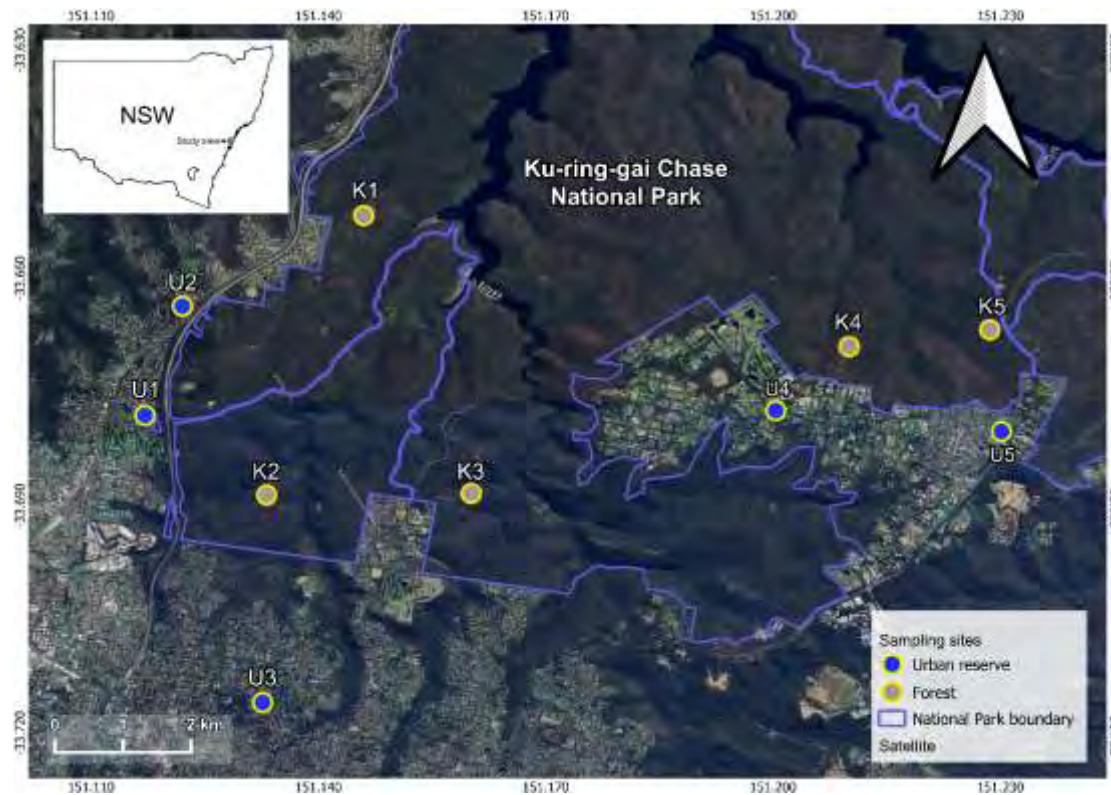


Aims

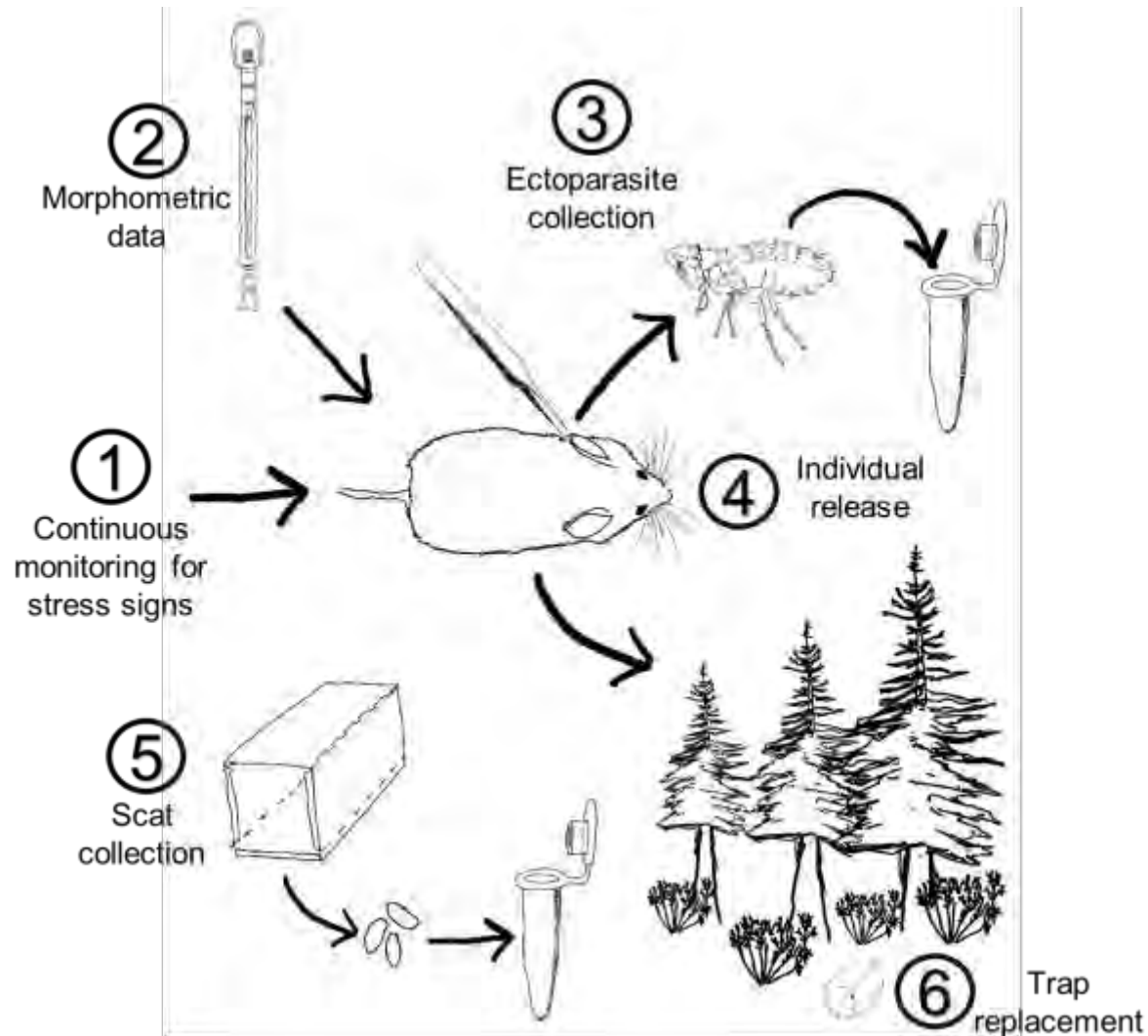
- Compare the **EMF community diversity** in small mammal scat and soil samples from **urban reserves** and **forest sites** around Ku-ring-gai Chase National Park, NSW, Australia.
- Assess the effects of **seasonality** (spring vs. autumn) and different species of small mammal **dispersal vectors** (i.e., *Antechinus stuartii* vs *Rattus fuscipes*) on EMF community diversity.

Methods

Study sites

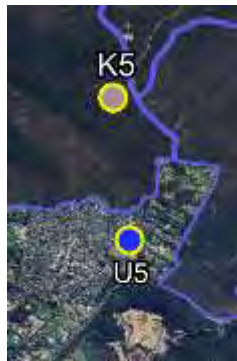


Methods



Methods

Three sampling
nights x site x
season



DNA extraction

- 110 scat samples
- 49 soil samples



NGS
barcoding

- ITS1-2 primers
- DADA2 pipeline



EMF fungal
diversity

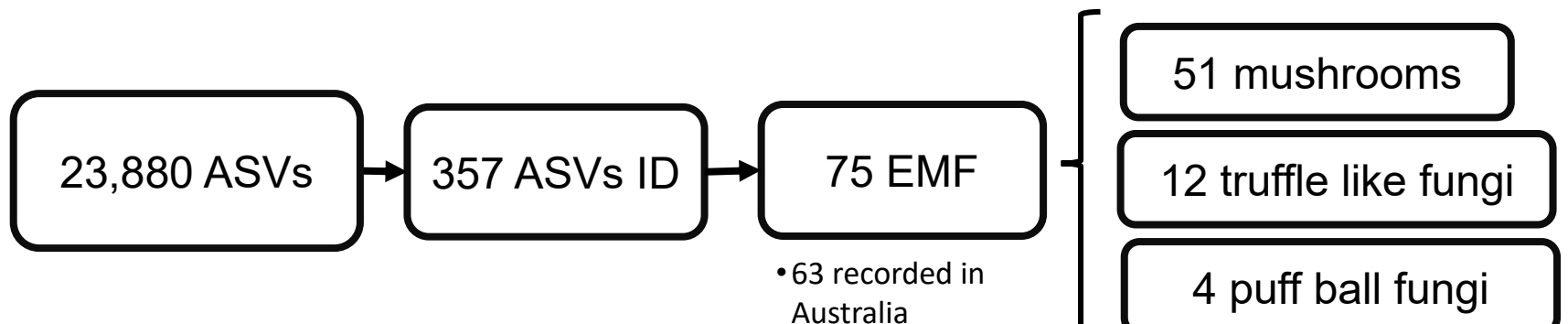
- Species richness
- Community composition
- Indicator species



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Results

- We obtained 100 scat samples in forest sites, while only 10 scat samples were collected in urban reserves.

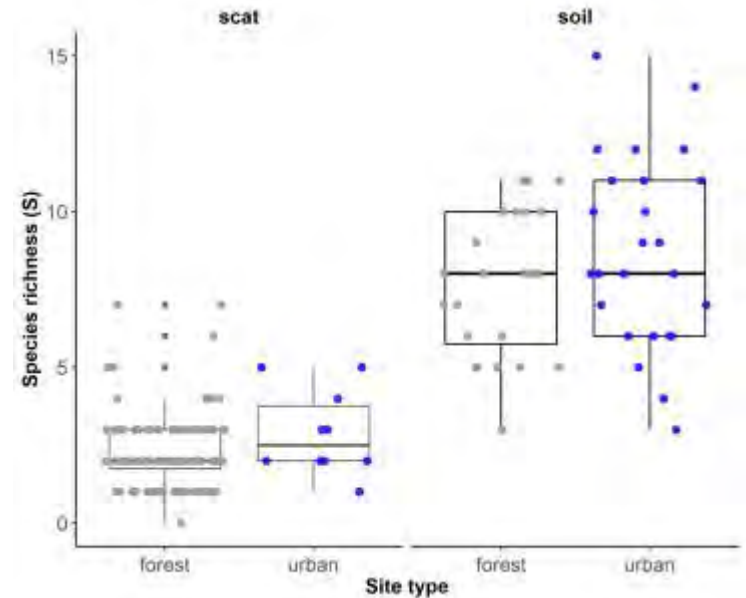
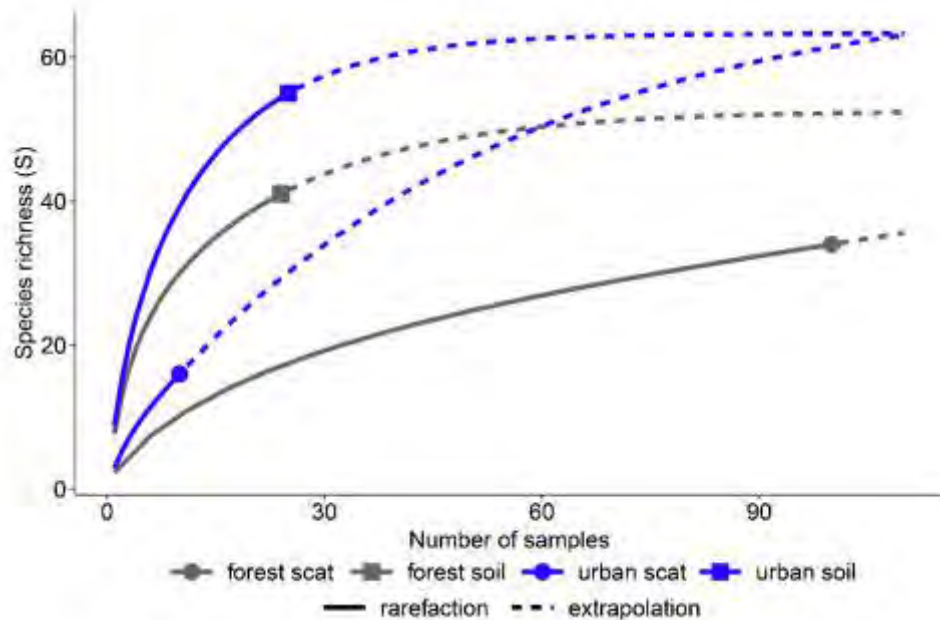


IDU	Site	RDP class	Kingdom	Phylum	Subdivision	Class	Subclass	Order	Family	Genus	Species	RDP conf	Species	Match	Match	E-1.1	E-1.1.1	E-1.1.2
OTU_9157	8	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	99.3	1	0	0	0
OTU_9608	10	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	100	2	0	0	0
OTU_4403	84	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.72	Agaricomycetes	100	1	0	0	0
OTU_1114	845	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	99.1	0	0	0	0
OTU_1608	152	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.98	Agaricomycetes	97.7	0	0	0	0
OTU_943	152	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	100	1	0	0	0
OTU_1807	261	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.61	Agaricomycetes	98.0	1	0	0	0
OTU_2862	142	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.61	Agaricomycetes	97.3	1	0	0	0
OTU_1229	151	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.57	Agaricomycetes	96.5	1	0	0	0
OTU_635	225	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.64	Agaricomycetes	99.1	1	0	0	0
OTU_4298	81	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	100	1	0	0	0
OTU_3057	14	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.60	Agaricomycetes	97.9	1	0	0	0
OTU_6063	11	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.92	Agaricomycetes	97.1	1	0	0	0
OTU_5812	28	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.55	Agaricomycetes	99.7	1	0	0	0
OTU_7900	12	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.57	Agaricomycetes	99.4	1	0	0	0
OTU_6209	26	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.78	Agaricomycetes	97.3	1	0	0	0
OTU_5888	80	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.90	Agaricomycetes	99.3	1	0	0	0
OTU_616	2579	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.74	Agaricomycetes	100	13	0	0	0
OTU_4325	84	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	97.3	1	0	0	0
OTU_7063	13	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	99.6	1	0	0	0
OTU_9717	4	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.96	Agaricomycetes	96.3	1	0	0	0
OTU_4239	58	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	99.3	1	0	0	0
OTU_2895	264	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	99.3	1	0	0	0
OTU_1018	250	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.88	Agaricomycetes	97.6	1	0	0	0
OTU_4662	71	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	100	1	0	0	0
OTU_645	4333	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	97.7	1	0	0	0
OTU_1028	4	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	98.3	1	0	0	0
OTU_666	4001	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.95	Agaricomycetes	96.7	1	0	0	0
OTU_139	14978	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.79	Agaricomycetes	100	3	0	0	0
OTU_3061	131	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.99	Agaricomycetes	97.3	1	0	0	0
OTU_6520	9	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	98.4	1	0	0	0
OTU_895	1331	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.64	Agaricomycetes	100	4	0	0	0
OTU_466	4311	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.74	Agaricomycetes	100	1	0	0	0
OTU_118	3944	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	1.000	Agaricomycetes	100	2	0	0	0
OTU_3884	88	Root	Fungi	Basidiomycota	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	Agaricomycetes	0.90	Agaricomycetes	99.6	1	0	0	0

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Diversity of EMF in response to urbanisation

- Soil samples are more diverse than scat samples
- Urban samples are more diverse than forest samples



Indicator taxa

Forest scat samples



© Roy Halling

Ionosporus australis

Urban scat samples



© julian_alonso

Descomyces albus

Forest soil (7 taxa)



© J. A. Cooper

Cortinarius durifoliorum

Urban soil (13 taxa)



© N. Siegel

Russula pseudoareolata



© Alice Shanks

Rossbeevera pachydermis



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Hysteragium



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Hydnoplicata convoluta

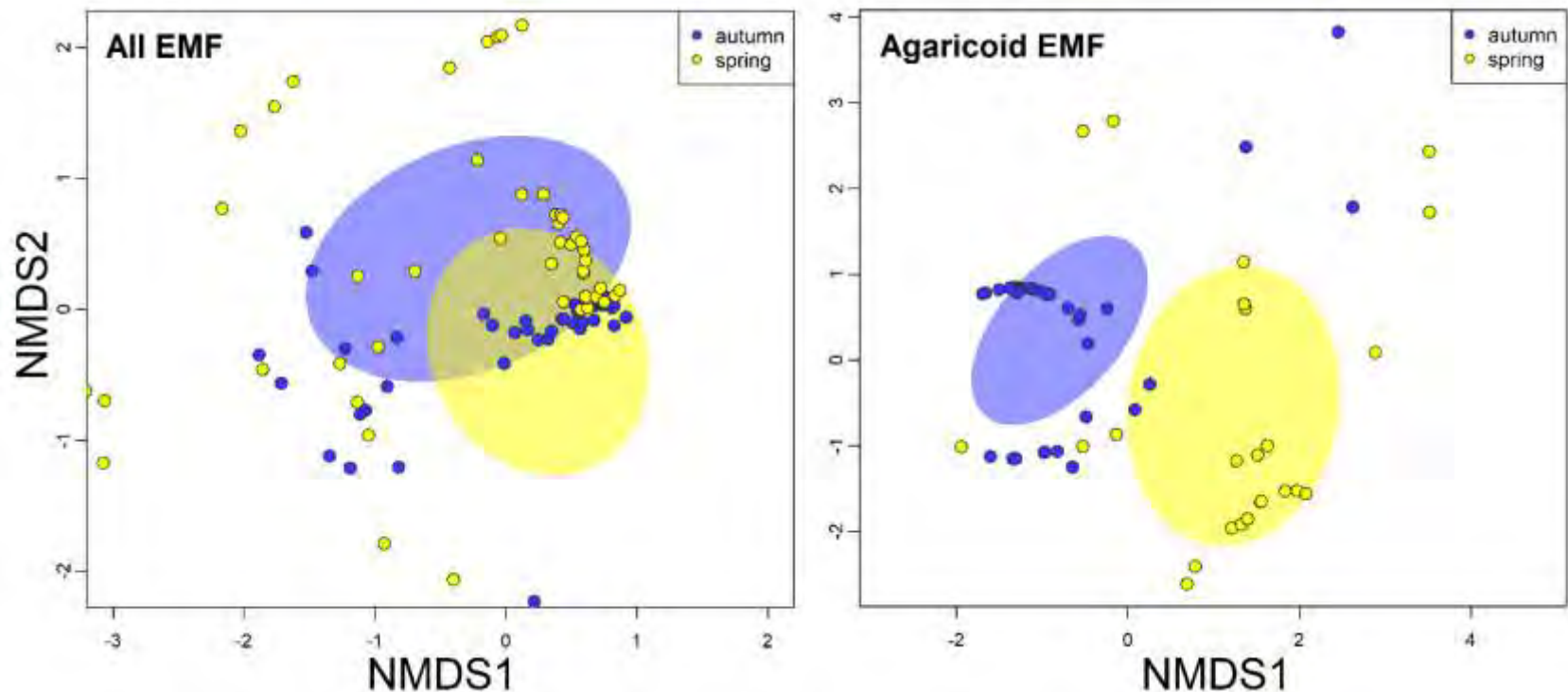


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Scleroderma albidum

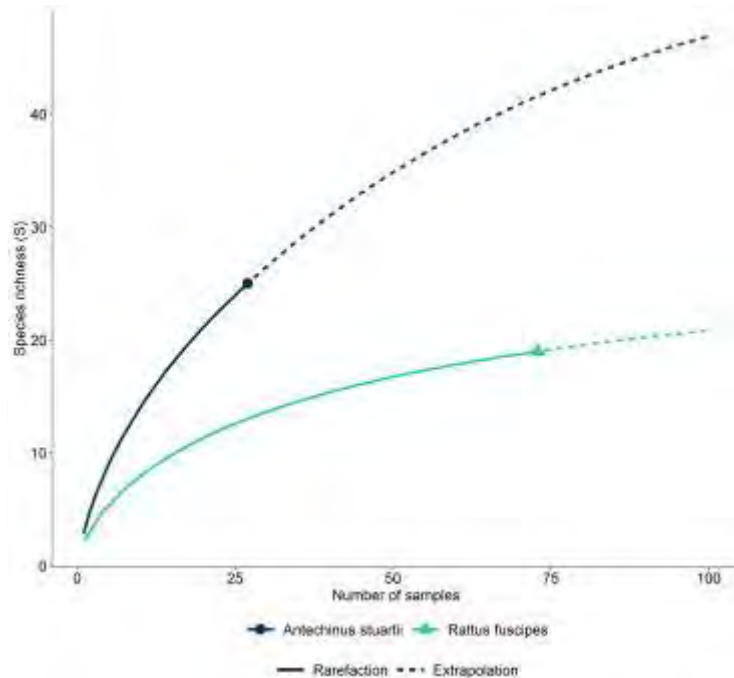
Seasonal variation in EMF diversity in scats

- Significant differences in diversity (H') but not in richness.
- Significant difference in the composition of agaricoid EMF.

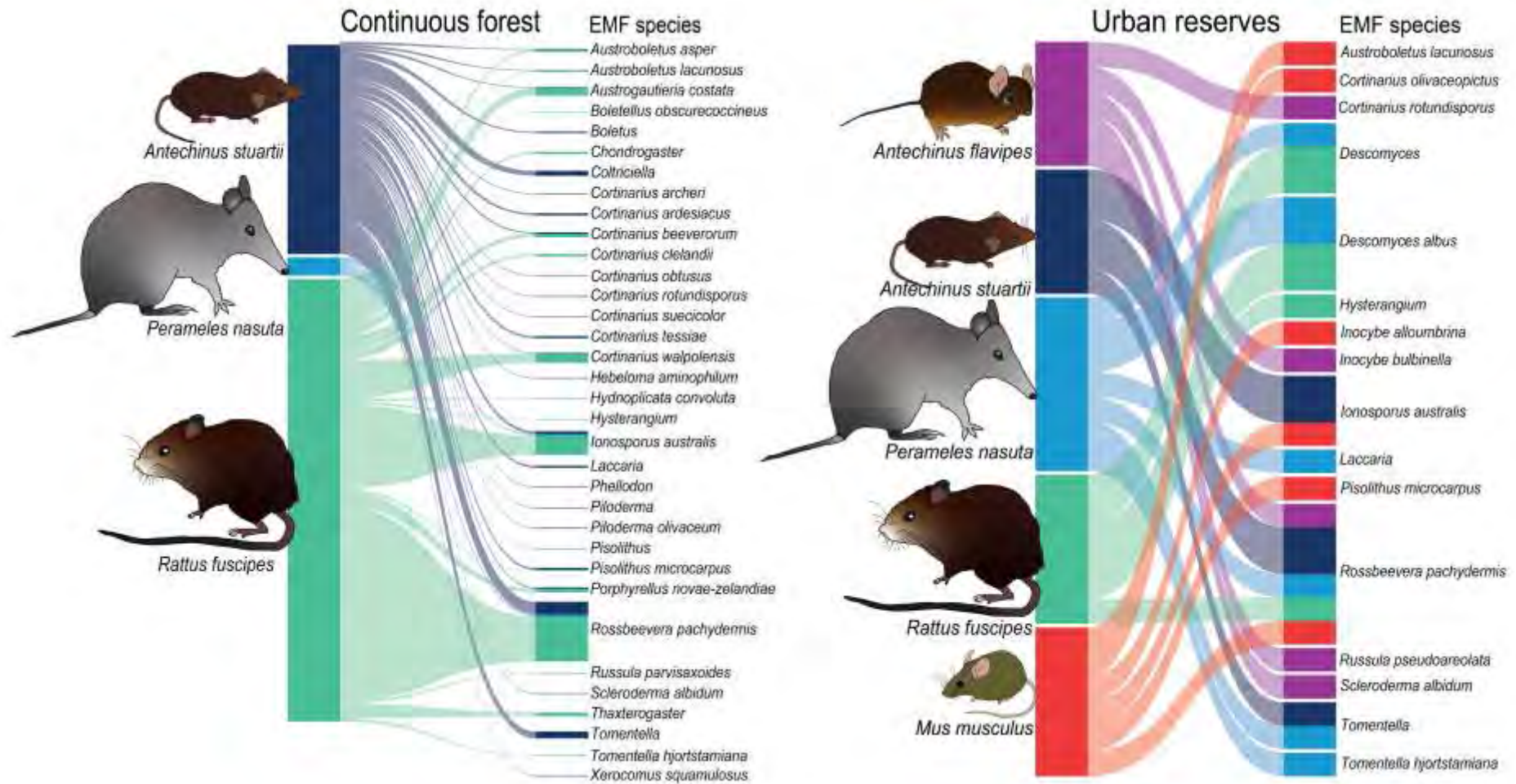


Small mammal vectors and EMF diversity

- Our sample: *Antechinus flavipes* (n = 1), ***A. stuartii*** (n = 27), *Perameles nasuta* (n = 5), ***Rattus fuscipes*** (n = 73), and *Mus musculus* (n = 2)
- Only one of the scat samples did not contain EMF
- *A. stuartii* had a higher diversity of EMF



Small mammal vectors and EMF diversity



Conclusions

- **Urban reserves** provide a refuge for EMF diversity
- Soil samples are highly informative, but scat samples tell us more about the **phenology** of EMF
- **Small mammals** are strongly associated with EMF
- Protecting soil and small mammals from disturbance can enhance EMF diversity, and **forest resilience**.

Integrative ecology of small mammals in
response to urbanisation

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graph TD; A[Integrative ecology of small mammals in response to urbanisation] --- B[Ectomycorrhizal fungi diversity associated with small mammals in response to urbanisation]; A --- C[Small mammal behaviour in response to urbanisation];
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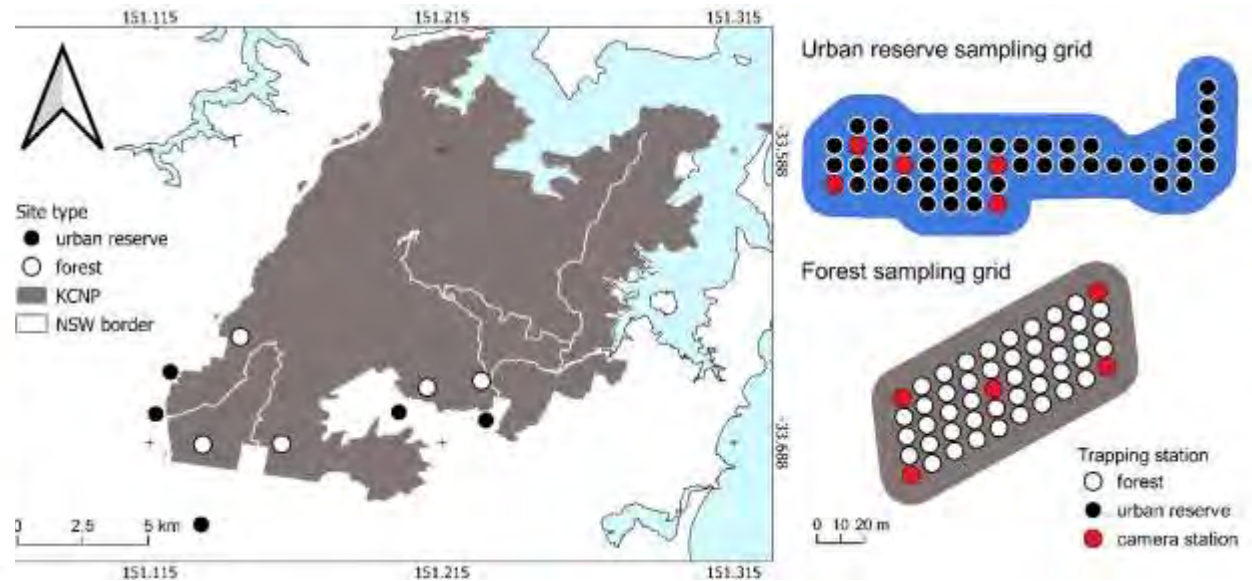
Ectomycorrhizal fungi diversity associated with small
mammals in response to urbanisation

Small mammal behaviour in response to
urbanisation

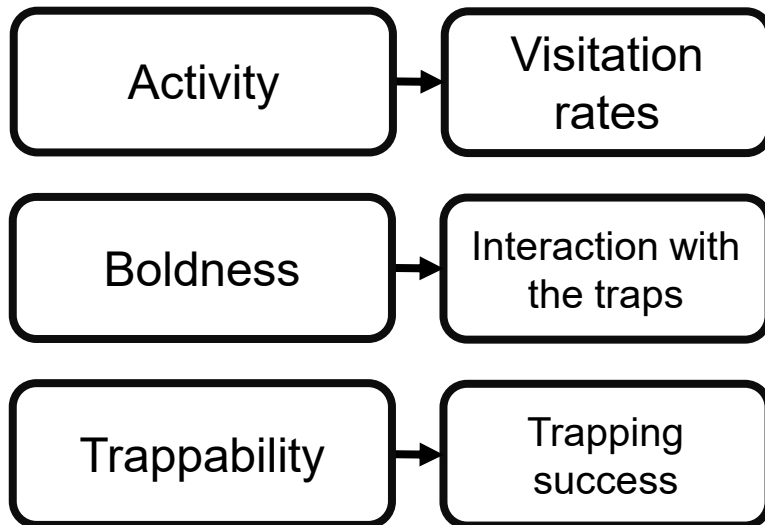
Small mammal behaviour in urban areas

- The behaviour of small mammals has a strong influence in the ecosystem services they deliver
- Small mammal diversity estimates obtained by live-trapping are strongly influenced by animal behaviour
- Small mammal face unique challenges in urban areas; they adapt through their behaviour
- We aimed to analyse the influence of urbanisation on **activity, boldness and trappability** of small mammals.

Methods



Methods



Results

- 317 camera-trap nights; 448 independent visits; 20 species



Results

Site type	Species	Mean trap success ¹ ± SD	Visitation rate ² ± SD	Trap interaction (s) ± SD
Forest	<i>Antechinus stuartii</i>	2.33 ± 2.29	0.16 ± 0.12	14.79 ± 20.09
	<i>Cercartetus nanus</i>	0	0.01 ± 0.02	1
	<i>Perameles nasuta</i>	0.2 ± 0.3	0.08 ± 0.08	20 ± 32.31
	<i>Rattus fuscipes</i>	➔ 8.67 ± 5.23	0.68 ± 0.42	➔ 48.84 ± 74.16
Urban reserve	<i>Antechinus flavipes</i>	0.07 ± 0.15	0.01 ± 0.02	2.5 ± 2.12
	<i>Antechinus stuartii</i>	0.46 ± 0.87	0.03 ± 0.07	8.4 ± 17.13
	<i>Perameles nasuta</i>	0.27 ± 0.6	0.24 ± 0.22	23.53 ± 56.67
	<i>Rattus fuscipes</i>	0.13 ± 0.18	0.31 ± 0.43	3.1 ± 6.13
	<i>Rattus rattus</i>	➔ 0	➔ 0.64 ± 0.59	5.55 ± 18.46
	<i>Mus musculus</i>	0.27 ± 0.6	0.06 ± 0.06	23.1 ± 27.55
	<i>Oryctolagus cuniculus</i>	0.13 ± 0.3	0.03 ± 0.04	0

Results



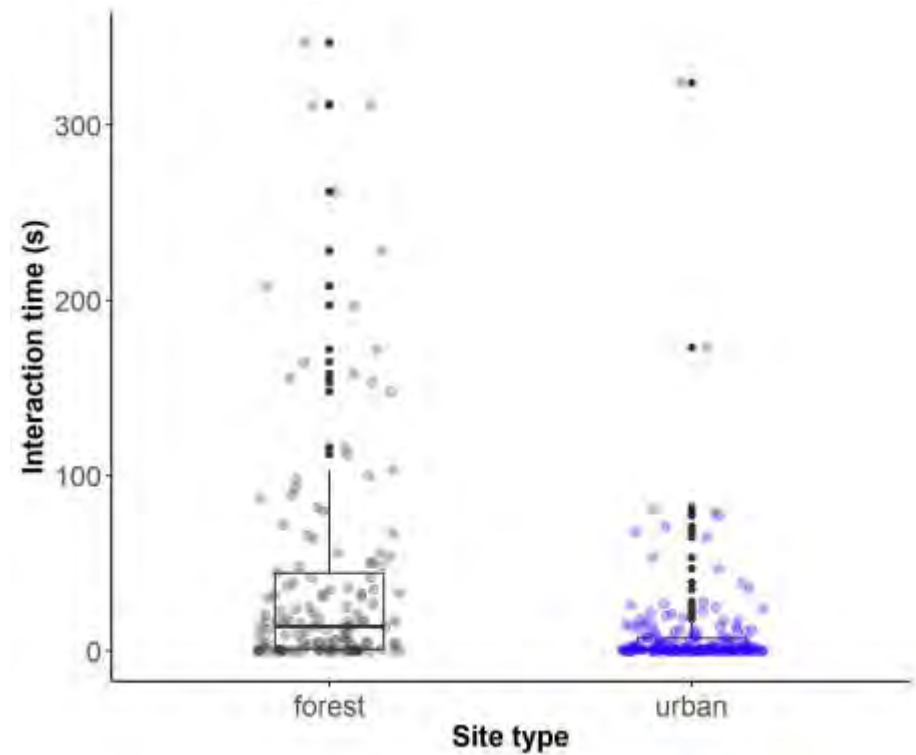
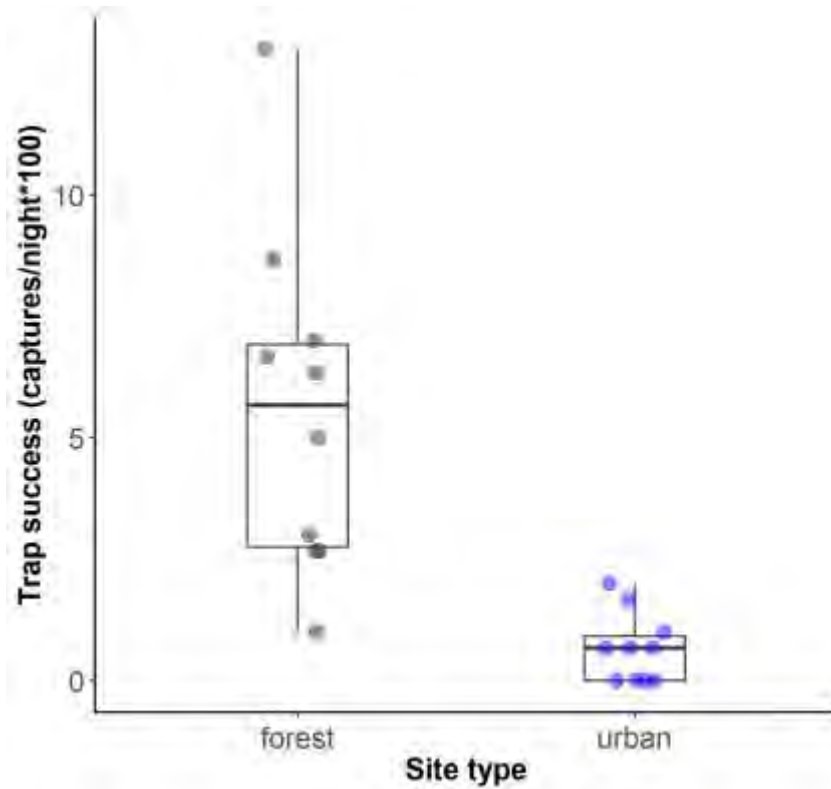
Results



Results



Results



Conclusions

- Small mammals in urban reserves were less bold and had **lower trappability** (i.e., trap success) than their forest counterparts. However, this was not a result of lower small mammal activity in the urban reserves, which had similar visitation rates as the forest sites.
- Camera-trapping in conjunction with live-trapping helps assessing small mammal **behaviour towards the trapping devices**

Collaborators



Prof Jaco Le Roux



Dr Christian A. Delfin



Dr Alexandra Carthey



Dr Ryan Stephens



Dr Margarita Vargas



Dr Cristina MacSwiney

Acknowledgements



CONAHCYT
CONSEJO NACIONAL DE HUMANIDADES
CIENCIAS Y TECNOLOGÍAS



northern
beaches
council



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Thank you!