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**Table S1. Process IDs for *Rattus* sp. *coi* barcodes deposited the Barcode of Life Data System (BOLD) database**

<b>Sampling site</b>	<b>Process ID</b>
Johor	UMNPA004-20 – UMNPA056-20
	UMNPA058-20 – UMNPA068-20
Perak	UMNPA069-20
	UMNPA071-20 – UMNPA076-20
	UMNPA078-20 – UMNPA080-20
	UMNPA082-20 – UMNPA083-20
	UMNPA085-20
	UMNPA087-20 – UMNPA091-20
	UMNPA093-20 – UMNPA102-20
	UMNPA161-20 – UMNPA194-20
	UMNPA196-20 – UMNPA216-20
	UMNPA218-20 – UMNPA223-20

**Table S2. Small mammals captured at different habitat types**

<b>Habitat type</b>	<b>n</b>	<b>Animal</b>	<b>n</b>
Plantation	121	<i>Rattus tanezumi</i> R3 mitotype	78
		<i>Tupaia glis</i>	21
		<i>Rattus tiomanicus</i>	12
		<i>Rattus exulans</i>	7
		<i>Rattus argentiventer</i>	3
Human dwelling	47	<i>Rattus tanezumi</i> R3 mitotype	26
		<i>Tupaia glis</i>	13
		<i>Rattus tiomanicus</i>	6
		<i>Rattus exulans</i>	2
Paddy field	26	<i>Rattus argentiventer</i>	22
		<i>Rattus exulans</i>	2
		<i>Rattus tanezumi</i> R3 mitotype	1
		<i>Rattus tanezumi</i> s.s.	1
Rubbish dumpsite	13	<i>Rattus tanezumi</i> R3 mitotype	10
		<i>Tupaia glis</i>	2
		<i>Rattus exulans</i>	1
Forest border	10	<i>Rattus tiomanicus</i>	4
		<i>Tupaia glis</i>	4
		<i>Rattus tanezumi</i> R3 mitotype	1
		<i>Rattus exulans</i>	1

**Table S3. Chigger infestation status based on animal host species, sex and maturity**

<b>Host species</b>	<b>Infestation rate</b>	<b>Mean chigger intensity</b>	<b>Median chigger intensity</b>	<b>Range</b>
<i>Rattus tanezumi</i> R3 mitotype	99/116 (85%)	282	192	5-1,511
<i>Tupaia glis</i>	36/40 (90%)	238	94	2-2,735
<i>Rattus exulans</i>	8/13 (61%)	201	139	17-746
<i>Rattus tiomanicus</i>	20/22 (91%)	117	80	3-477
<i>Rattus argentiventer</i>	5/25 (20%)	63	27	1-156

<b>Host gender</b>	<b>Infestation rate</b>	<b>Mean chigger intensity</b>	<b>Median chigger intensity</b>	<b>Range</b>
Female	82/103 (80%)	278	147	1-2,735
Male	87/114 (76%)	206	141	2-975

<b>Host maturity</b>	<b>Infestation rate</b>	<b>Mean chigger intensity</b>	<b>Median chigger intensity</b>	<b>Range</b>
Adult	119/149 (80%)	274	146	1-2,735
Juvenile*	48/65 (74%)	164	127	6-731

<b>Season</b>	<b>Infestation rate</b>	<b>Mean chigger intensity</b>	<b>Median chigger intensity</b>	<b>Range</b>
Dry	118/125 (94%)	292	175	3-2,735
Wet	51/92 (55%)	124	112	1-588

\*Sub-adults and juveniles were combined into a single category for this analysis

**Table S4. Number of small mammals with the indicated chigger species richness**

<b>Chigger species richness</b>	<b>Number</b>	<b>Percentage (%)*</b>
1	116	53.5
2	46	21.2
3	40	18.3
4	14	6.5
5	1	0.5

\*Out of total number of small mammals (n=217)

**Table S5. Number of each small mammal species with chigger coinfection**

<b>Host species</b>	<b>Total</b>	<b>Number coinfecting</b>	<b>Percentage (%)</b>
<i>Rattus argentiventer</i>	25	0	0
<i>Rattus exulans</i>	13	7	53.8
<i>Rattus phylogenetic R3</i>	116	66	56.9
<i>Rattus tanezumi</i>	1	0	0
<i>Rattus tiomanicus</i>	22	11	50.0
<i>Tupaia glis</i>	40	17	42.5

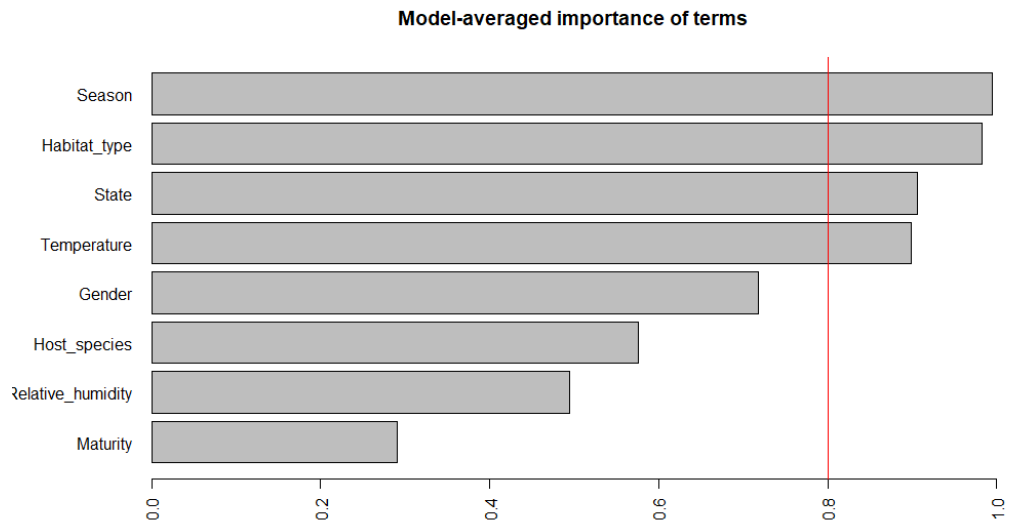
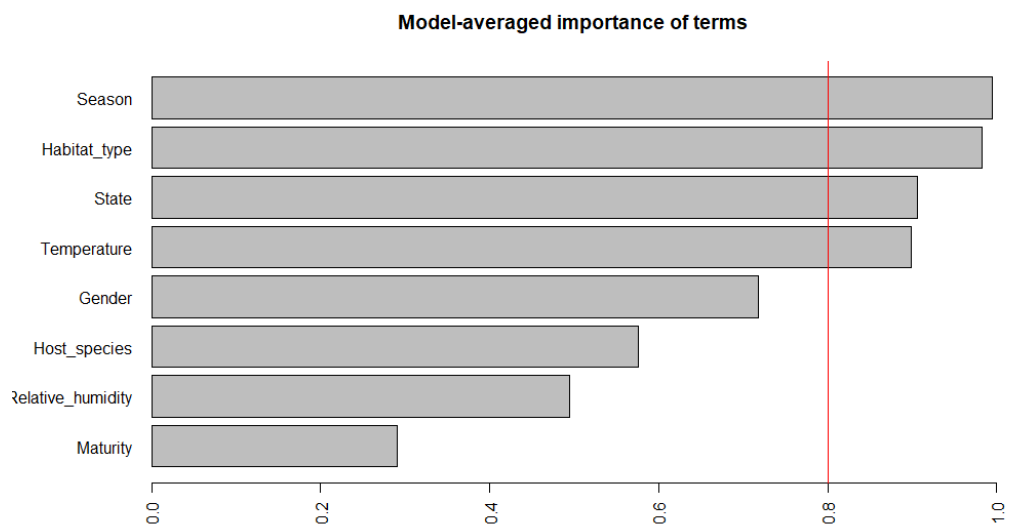
**A****B**

Figure S1. Model-average importance of terms of independent variables explaining chigger (a) abundance and (b) species richness. The variables with an importance score of >80% proportion support (x-axis, default in the “gmulti” package in R) were included in the final model.

**Table S6. Significant variables from best selected model for explaining chigger abundance**

Parameters/effects	Log ratio Chi-square	P-value	Variance inflation factor
Habitat_type***	67.220	<0.001	1.529
Temperature	0	0.9845	1.075
Season***	21.743	<0.001	1.112
State*	4.969	0.0258	1.313
<b>Observation</b>			216
<b>Log-likelihood</b>			-1191.987
<i>AICc</i>			2400
<b>The Goodness of Fit/Coefficient of determination (R<sup>2</sup>)</b>			0.9999761

**Table S7. Significant variables from best selected model for explaining chigger species richness**

Parameters/effects	Log ratio Chi-square	P-value	Variance inflation factor
Habitat_type***	56.053	<0.001	1.543
Relative_humidity*	5.341	0.0208	1.230
Season***	35.594	<0.001	1.093
State***	11.933	<0.001	1.286
<b>Observation</b>			216
<b>Log-likelihood</b>			-285.057
<i>AICc</i>			586.115
<b>The Goodness of Fit/Coefficient of determination (R<sup>2</sup>)</b>			0.9999894

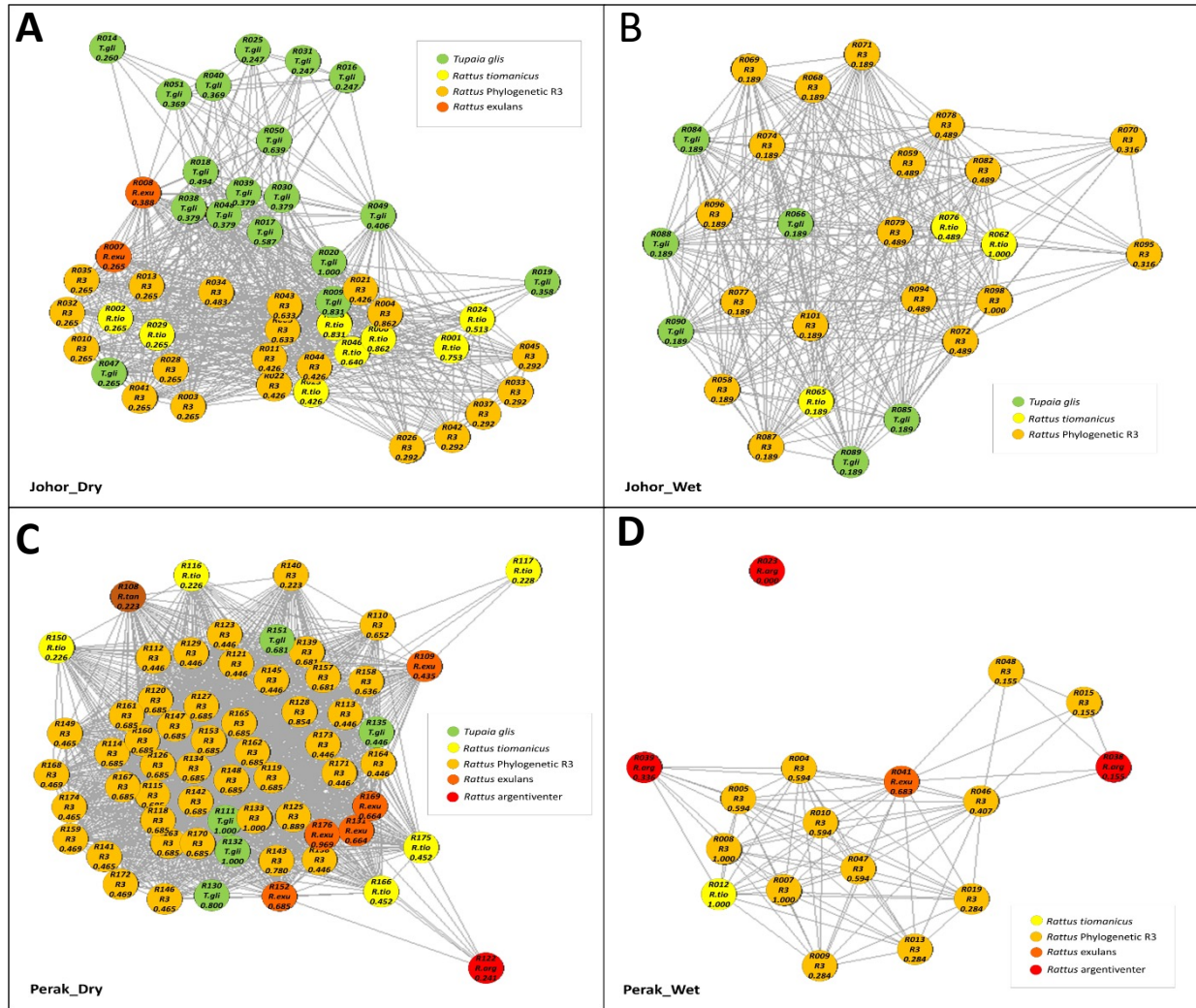


Figure S2. Unipartite network models at host individual levels by state and season. (a, b) Johor. (c, d) Perak. (a, c) Dry season. (b, d) Wet season.



**Table S8. Linkage density and V-ratio based on sampling site and season**

Network parameters	Johor/Dry (n = 48)	Johor/Wet (n = 27)	Perak/Dry (n = 62)	Perak/Wet (n = 17)
Linkage density	10.9255	10.7105	21.6172	5.2812
V-ratio (chigger-host)	12.3597	16.1200	33.9133	5.4729

Bipartite network parameters were calculated using ‘bipartite’ package in R freeware.

**Table S9. Prevalence of *Orientia tsutsugamushi* in small mammals from the Johor and Perak sites**

Category	n	Number of <i>Orientia</i> infection	Prevalence of infection (%)	[95% CI]
<b>Host species</b>				
<i>R. argentiventer</i>	25	2	8.00	1.4 to 27.5
<i>R. exulans</i>	12	2	16.7	2.9 to 49.1
<i>R. tanezumi</i> R3 mitotype	115	18	15.7	9.8 to 23.9
<i>R. tanezumi</i> s.s.	1	0	0	0
<i>R. tiomanicus</i>	21	1	4.8	0.3 to 25.9
<i>T. glis</i>	40	2	5.00	0.9 to 18.2
<b>Host gender</b>				
Female	102	13	12.8	7.2 to 21.2
Male	112	12	10.7	5.9 to 18.3
<b>Maturity</b>				
Adult	149	18	12.1	7.5 to 18.7
Sub-adult	41	6	14.6	6.1 to 29.9
Juvenile	24	1	4.2	0.2 to 23.1
<b>Habitat type</b>				
Forest border	9	0	0	0
Human dwelling	47	4	8.5	2.8 to 21.3
Paddy field	26	2	7.7	1.3 to 26.6
Plantation	119	18	15.1	9.5 to 23.1
Rubbish dumpsite	13	1	7.7	0.4 to 37.9
<b>State</b>				
Johor	99	10	10.1	5.2 to 18.2
Perak	115	15	13.0	7.7 to 20.9
<b>Season</b>				
Dry	123	14	11.4	6.6 to 18.7
Wet	91	11	12.1	6.5 to 21.0
<b>Total</b>	<b>214</b>	<b>25</b>	<b>11.7</b>	<b>7.9 to 16.9</b>