1 2	SUPPLEMENTARY MATERIAL	
3	Predictive analysis across spatial scales links zoonotic malaria to deforestation	
4		
5	Patrick M. Brock, Kimberly M. Fornace, Matthew J. Grigg, Nick M. Anstey, Timothy William,	
6	Jon Cox, Chris J. Drakeley, Heather M. Ferguson, Rowland R. Kao	
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10	Supplementary Material Content	
11	Figure S1: Top 16 predictors	
12	Figure S2: Marginal effect curves of full model including non-scalable variables	
13	Figure S3: Correlations across scales	
14	Figure S4: Variance with relative variable importance	
15	Figure S5: Case cluster profiles	
16	Figure S6: Single-scale model predictive ability by scale	

17 Table S1: Correlations of variance with relative variable importance

18 Figure S1. Relative variable importance (RVI) of the 16 most important variable-scale

19 combinations from a) the full BRT model fitted across the whole study site (176 predictors);

20 and b) the full BRT model fitted across the whole study site including two additional non-

Distance

Clearing P:A (previous year) 0.5 km Population density 2 km Mean slope 0.5 km Cover P:A (previous year) 0.2 km Cover P:A (previous year) 0.1 km Loss (previous year) 0.5 km Loss P:A (previous year) 4 km Loss P:A (previous year) 5 km

Environmental

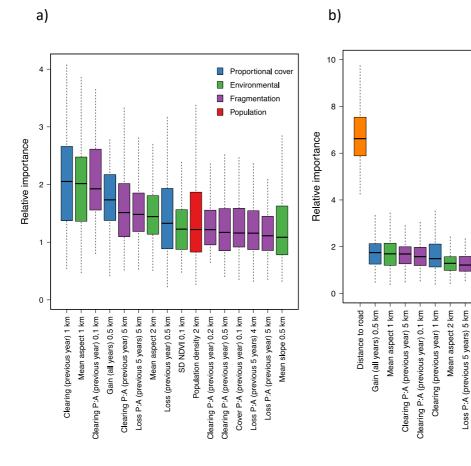
Fragmentation

Population

Proportional cover

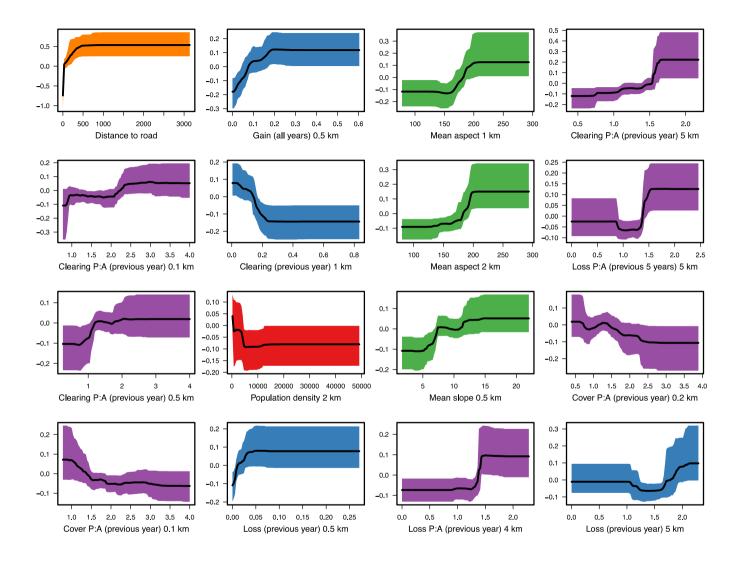
21 scalable variables – shortest distance to road and clinic – (178 predictors).

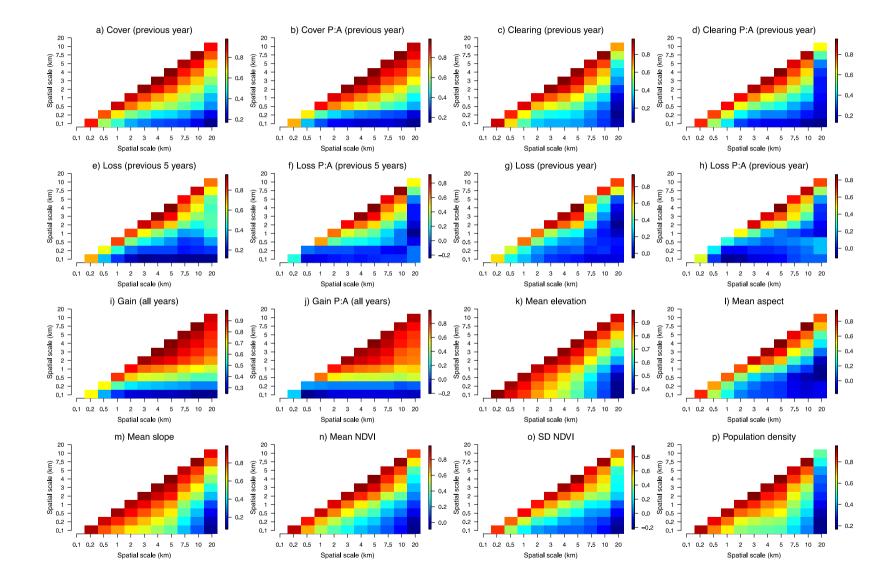






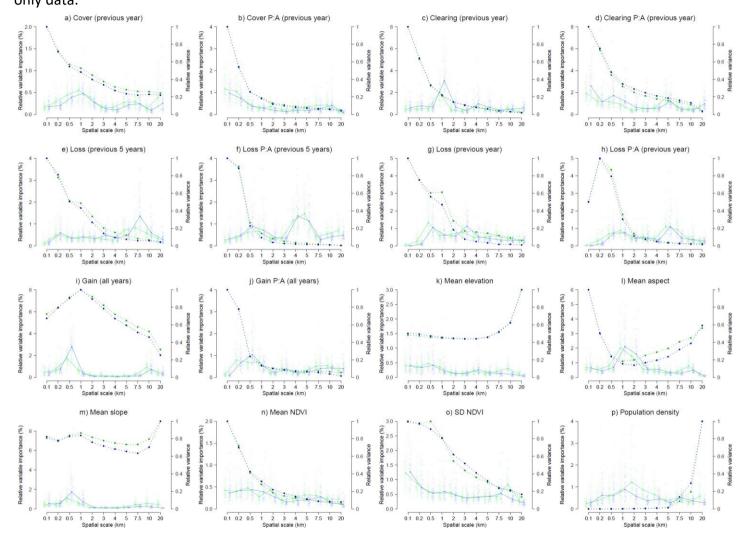
- 26 Figure S2. Marginal effect curves of the 16 variable-scale combinations with the highest relative variable importance (RVI) from the full BRT
- 27 model fitted across the whole study site including two additional non-scalable variables shortest distance to road and clinic (178
- 28 predictors).
- 29
- 30
- 31





34 Figure S3. Spearman rank correlations between scales for all 16 scalable landscape variables.

- 37 Figure S4. Relative variances superimposed on the traces of relative variable importance (RVI) of all variable-scale combinations (as shown in
- *Figure 2*) from full BRT models of *P. knowlesi* occurrence (176 predictors). Green points represent the whole study site, blue points mainlandonly data.



- 41 Figure S5. Variation in the 15 variable-scale combinations with the greatest differences in
- 42 median marginal probability of occurrence between clusters. Cluster A shown in purple,

43 cluster B shown in orange.

44

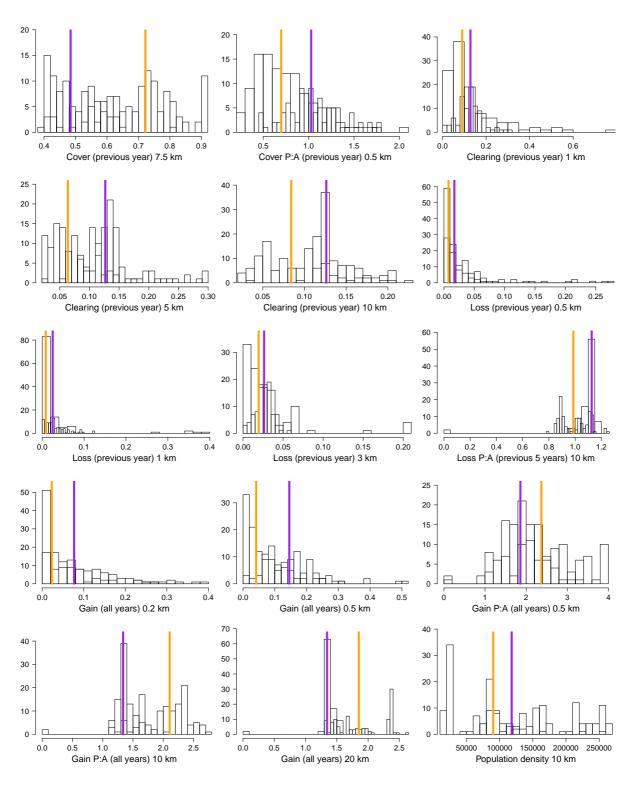
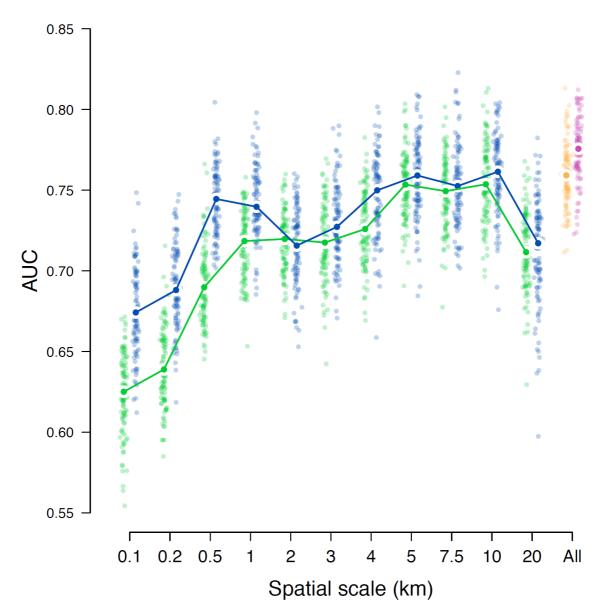


Figure S6. The predictive ability of BRT models of *P. knowlesi* occurrence across spatial
scales, with variables summarised over the same single scale (0.1-20 km; 16 predictors per
model) and simultaneously over all 11 scales (All; 176 predictors per model). Green and
yellow points represent models fitted across the whole study site, blue and purple points
models fitted to mainland data only. AUC denotes area under the receiver operator curve.



- 54 Table S1. Spearman rank correlations comparing relative variance with median relative
- *variable importance (RVI) across spatial scales from the full BRT model fitted across the*
- whole study site (176 predictors).

Variable	Spearman's rank correlation	Spearman's rank correlation test
	coefficient	p-value
Cover (previous year)	0.02	0.539
Cover P:A (previous year)	0.91	0.061
Cleared (previous year)	0.05	0.248
Cleared P:A (previous year)	0.76	0.094
Loss (previous 5 years)	-0.49	0.286
Loss P:A (previous 5 years)	-0.44	0.087
Loss (previous year)	-0.17	0.86
Loss P:A (previous year)	-0.22	0.734
Gain (all years)	0.26	0.755
Gain P:A (all years)	0.03	0.734
Mean elevation	-0.31	0.989
Mean aspect	-0.38	0.061
Mean slope	0.36	0.094
Mean NDVI	0.63	0.034
SD NDVI	0.6	0.052
Population density	-0.09	0.924