

Table S4. Model comparisons for Lyme disease and babesiosis. (We fitted a series of nested models and evaluate model fit with Δ DIC compared with the non-spatial process-based model. Process-based models include: (i) a non-spatial process model; (ii) a spatial model allowing for linear dependence of invasion mechanisms on local infection risk; (iii) a full covariate model including nonlinear dependence of invasion mechanisms on local infection risk and ecological covariates; (iv) a parsimonious model excluding any covariate from the full covariate model with a 95% credible interval that includes 0; and (v) a parsimonious model without the observation process. To justify modelling latent ecological processes, we also present results of fitting a ‘null’ phenomenological model described in the electronic supplementary material, text S1. All models estimate B , the coefficient for population size offset.)

	model description	observation parameter	ecological parameters	DIC	Δ DIC
Lyme disease	process-based model	r_s	$\mathbf{p}, \phi_0, \gamma_0, \theta_0$	17 462	—
	linear spatial dependence	r_s	$\mathbf{p}, \phi_0, \phi_1, \gamma_0, \gamma_1, \theta_0, \theta_1$	14 774	2688
	full covariate	r_s	$\mathbf{p}, \phi_0, \phi_1, \phi_2, \phi_3, \phi_4, \gamma_0, \gamma_1, \gamma_2, \gamma_3, \gamma_4, \theta_0, \theta_1, \theta_2, \theta_3, \theta_4$	14 321	3141
	parsimonious model	r_s	$\mathbf{p}, \phi_0, \phi_1, \phi_2, \phi_3, \gamma_0, \gamma_1, \gamma_2, \theta_0, \theta_1, \theta_3$	14 316	3146
	parsimonious no observation	r_s	$\mathbf{p}, \phi_0, \phi_1, \phi_2, \phi_3, \gamma_0, \gamma_1, \gamma_2, \theta_0, \theta_1, \theta_3$	14 333	3129
	null	r_s	τ_0, τ_1, τ_2	16 214	1248
babesiosis	process-based model	r_s	$\mathbf{p}, \phi_0, \gamma_0, \theta_0$	8691	—
	linear spatial dependence	r_s	$\mathbf{p}, \phi_0, \phi_1, \gamma_0, \gamma_1, \theta_0, \theta_1$	8038	653
	full covariate	r_s	$\mathbf{p}, \phi_0, \phi_1, \phi_2, \phi_3, \phi_4, \phi_5, \gamma_0, \gamma_1, \gamma_2, \gamma_3, \gamma_4, \gamma_5, \theta_0, \theta_1, \theta_2, \theta_3, \theta_4, \theta_5$	7293	1398
	parsimonious model	r_s	$\mathbf{p}, \phi_0, \phi_1, \phi_3, \phi_4, \phi_5, \gamma_0, \gamma_1, \gamma_2, \gamma_4, \gamma_5, \theta_0, \theta_1, \theta_4$	7287	1404
	parsimonious, no observation		$\mathbf{p}, \phi_0, \phi_1, \phi_3, \phi_4, \phi_5, \gamma_0, \gamma_1, \gamma_2, \gamma_4, \gamma_5, \theta_0, \theta_1, \theta_4$	7355	1336
	null	r_s	τ_0, τ_1, τ_2	8634	57