

Some issues in modelling biodiversity using spatially modelled covariates

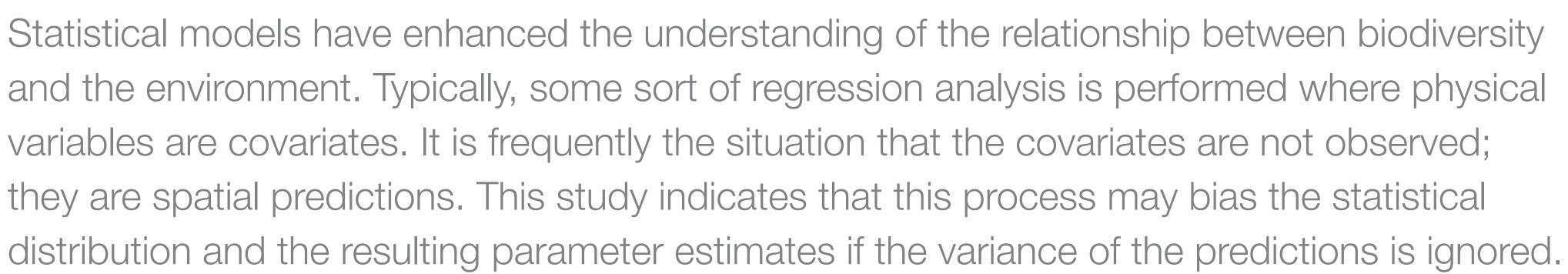
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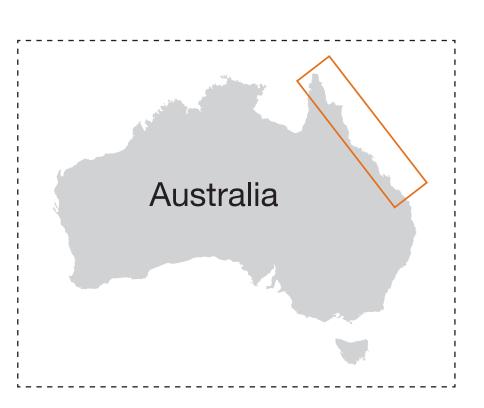


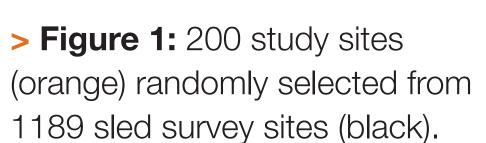




Great Barrier Reef data

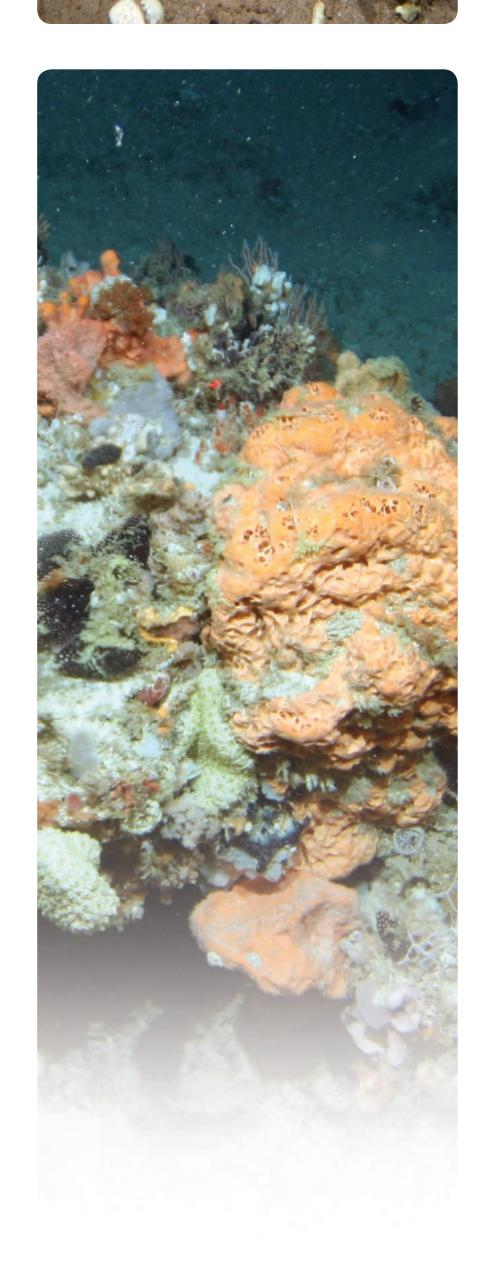
Data comprises of 1189 sites, where biological and physical variables were measured (Pitcher et al 2007, see Figure 1). The biological outcome variables were presence/absence of a particular species, and species richness (number of species) in a benthic sled sample. The physical variables used in this study were: depth, %carbonate and %mud. We randomly selected 200 sites from this complete set to mimic performing a biological survey (without measuring physical data).











Model and approximate bias

Let Y_i be a biological response such as species presence/ absence or richness, and let x_i be the vector of physical covariates at the survey site i. A generalised linear model is often used with

$$\mathrm{E}\left[Y_{i}|\boldsymbol{x}_{i}\right]=h\left(\eta_{i}\right)=h\left(\boldsymbol{x}_{i}^{\top}\boldsymbol{ au}\right),$$

where h is an inverse link function and τ is a px1 vector of unknown parameters. However, the physical covariates are commonly not observed but are estimated by spatial predictions, $\tilde{\boldsymbol{x}}_i$, based on observations at other sites. Allowing for variance from the spatially predicted covariate, the approximate mean and covariance of the outcomes are

$$\mathrm{E}\left[\mathrm{E}\left[Y_{i}| ilde{m{x}}_{i}
ight]|\left\{m{X}_{o}
ight\}
ight]=h\left(ilde{m{x}}_{i}^{ op}m{ au}
ight)+rac{1}{2}rac{d^{2}h}{d\eta_{i}^{2}}m{ au}^{ op}m{\Sigma}_{ii}m{ au},$$
 and

$$\operatorname{Cov}\left[Y_{i}, Y_{j} | \left\{\boldsymbol{X}_{o}\right\}\right] = \operatorname{E}\left[\operatorname{Cov}\left[Y_{i}, Y_{j} | \left\{\boldsymbol{X}_{o}\right\}\right]\right] + \frac{dh}{d\eta_{i}} \frac{dh}{d\eta_{j}} \boldsymbol{\tau}^{\top} \boldsymbol{\Sigma}_{ij} \boldsymbol{\tau},$$

where Σ_{ij} is the cross-covariance of prediction for the *i*th and *j*th physical covariates and $\{X_o\}$ are the observed physical covariates.

Simulation studies

To assess the size of relative bias a simulation study was performed. We simulated biological data given the observed data and analysed them using the spatially predicted covariates $(\tilde{\boldsymbol{x}}_i)$ and the observed covariates \boldsymbol{x}_i . A total of 1000 simulations were performed and are indexed by k in the equations below.

Simulation model 1

$\mathrm{E}\left[Y_i^{(k)}\right] = h\left(\tilde{\boldsymbol{x}}_i^{\top} \boldsymbol{ au}^{(k)}\right); \qquad \mathrm{E}\left[Y_i^{(k)}\right] = h\left(\boldsymbol{x}_i^{\top} \boldsymbol{ au}^{(k)}\right).$

Simulation model 2

$$\mathrm{E}\left[Y_i^{(k)}
ight] = h\left(oldsymbol{x}_i^ opoldsymbol{ au}^{(k)}
ight)$$

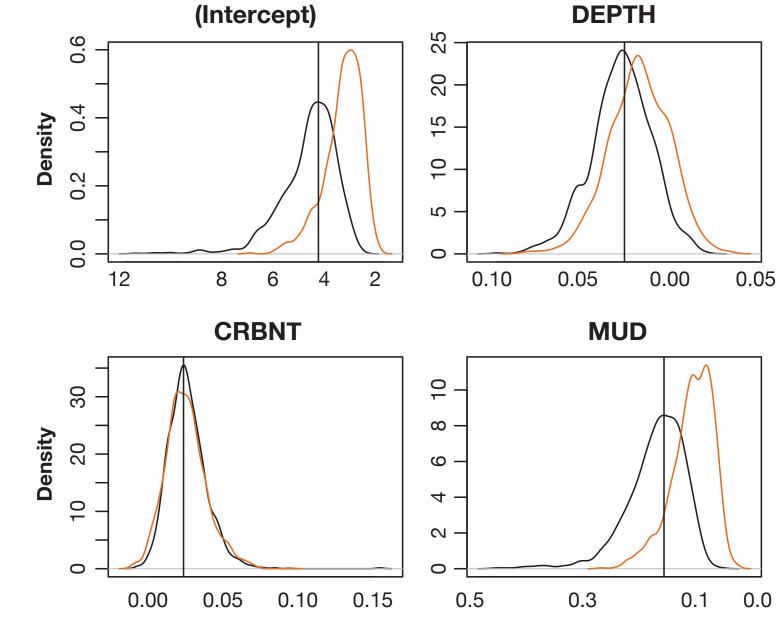
Results

Relative size of bias

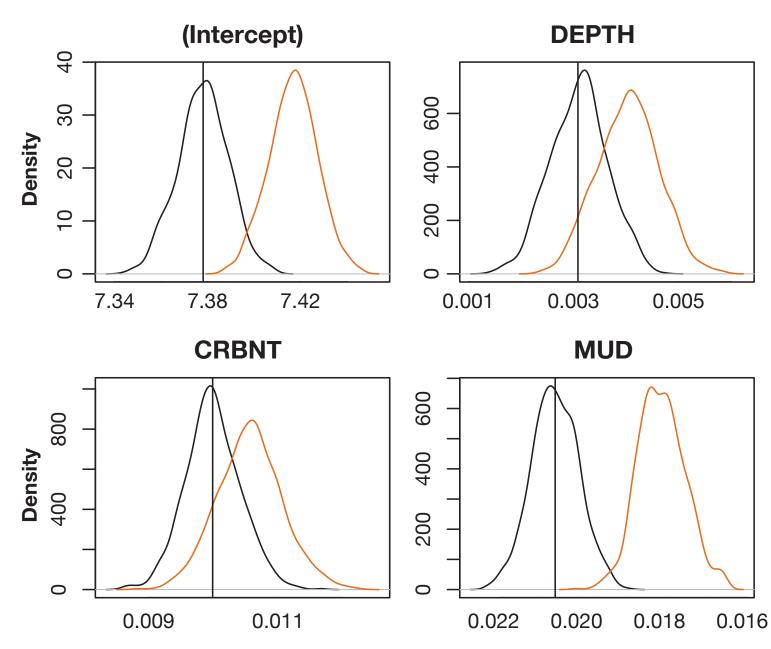
		Min.	Median	Max.
Presence/absence	bias/mean	-19.0%		839400.0%
(Cheilostomata Hippaliosina spp)	bias/variance	0.0%		1339000.0%
Richness	bias/mean	1.3%	3.5%	17.2%
	bias/variance	73.3%	361.6%	1200.0%
Summary statistics of the 200 sites				

Conclusions

- > Using spatially modelled covariates leads to bias in outcome distribution and parameter estimates
- > Simulation shows this is a real problem not just a theoretical one
- > Will need to account for uncertainty in future models



> Figure 2: Presence/absence - Empirical distribution of estimates from observed physical data (black) and predicted physical data (orange).



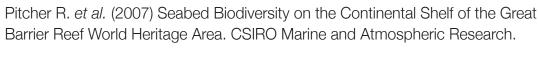
> Figure 3: Richness – Empirical distribution of estimates from observed physical data (black) and predicted physical data (orange).



data. Specific thanks go to Roland Pitcher, Zhi Huang, Kathy Haskard, Mark Palmer, Ross Darnell, Piers Dunstan, Jin Li, and Bill Venables.

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References













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