

# Estimating environmental heterogeneity from spatial dynamics data

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IISC

1 July 2015

# Outline

- 1 Introduction
- 2 Endogenous vs exogenous: qualitative
  - Explanations for spatial patterns
- 3 Endogenous vs exogenous: quantitative
  - Spatial synchrony
  - Pines
- 4 Conclusions

# What do ecologists want?

- *Explicit* questions: explain observed patterns
- *Implicit* questions: explain outcomes of ecological interactions: persistence, coexistence, trait evolution, etc..
- *Qualitative* answers: presence/absence, persistence/extinction, coexistence/exclusion
- *Quantitative* answers: how many? how quickly? scale(s) of pattern?
- *Deductive* (forward) models: model  $\rightarrow$  outcome
- *Inductive* (inverse) models: data  $\rightarrow$  model

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## Typical examples

- Can spatial pattern allow coexistence of similar species?
- Is a particular example of coexistence spatially mediated?
- Do endogenous or exogenous drivers produce spatial clustering?
- What drives spatial clustering in a particular case?

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# Typical models

- Stochastic spatial point processes: continuous time & space, point individuals
- Usually assume isotropy and translational invariance
- Population structure and environmental heterogeneity both described by *correlation functions*

e.g. *spatial logistic*:

- constant *per capita* fecundity  $f$
- dispersal kernel  $D(r)$
- death rate increases with local density:  

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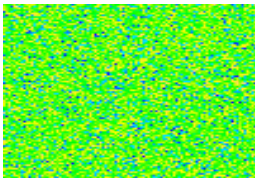
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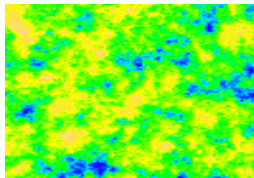
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# Spatial correlation

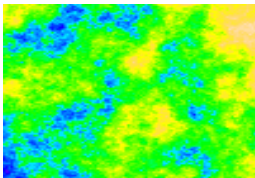
Random



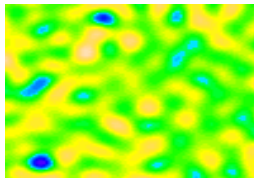
Short-range



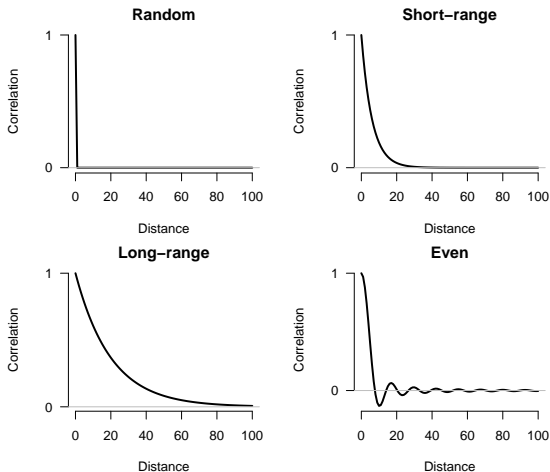
Long-range



Even

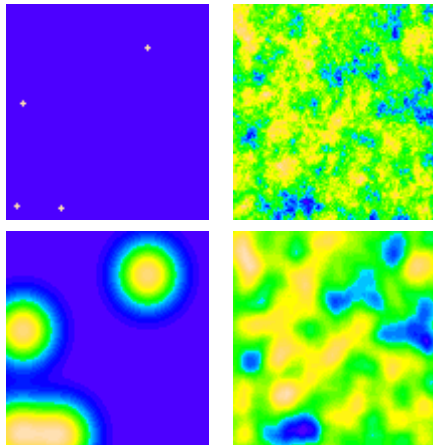


# Spatial correlation



# Spatial convolution

- distance-dependent spatial interactions  
(dispersal, competition, pollen flow ...)
- blurs out landscape according to scale of kernel
- relatively straightforward math
- easily described by *power spectra*:  
transformation of spatial correlation information



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# Templates

- assume habitat map reflects underlying spatial pattern more or less exactly: low diffusion (but  $> 0$ ), high growth rate



photo: Henry Horn





# Demographic noise-driven pattern

- Demographic stochasticity appears in correlation equations
- Drives pattern in homogeneous, stable systems (e.g. competition)
- Effects could be weak:
  - depend on scale of interaction neighborhood (Bolker, 1999):  
small if effective # neighbors > 10-20?
  - effects depend on  $R = f/\mu$ :  
for equal scale of dispersal & competition,  
no clumping if  $R > 2$  (Bolker & Pacala, 1999)
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# General (environmental) noise-driven pattern

Most general case: space(-time) noise  $\xi(x, t, N(t))$

- Scale may be  $> 0$  (non-white in space and time)
- Amplitude may scale differently from  $\sqrt{N}$
- Space-time correlations:
  - separable (Snyder & Chesson, 2004) ?
  - model correlations as their own sub-processes North et al. (2011)

Descriptions other than correlation functions? (Endler, 1986)

# Summary: what we need?

- Lots of interesting questions, but perhaps existing methods are good enough for ecologists?
- Importance of stochastic dynamics at various scales:
  - Is demographic (endogenous) noise really that important?
  - Perhaps a more traditional separation of scales (individual, patch, site, ...) is enough?
- **How do we estimate the (effects of) heterogeneity?**

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# Spatial synchrony

- Eastern spruce budworm, *Choristeroneura fumiferna*
- non-spatial dynamics: plant quality, climate, enemies (Kendall et al., 1999)?
- What generates large-scale spatial synchrony?
  - *Moran effect*: large-scale weather patterns
  - *Dispersal coupling*: movement of larvae



# Correlation equations: *via* continuous eqns

cf. Lande et al. (1999), Engen et al. 2002:

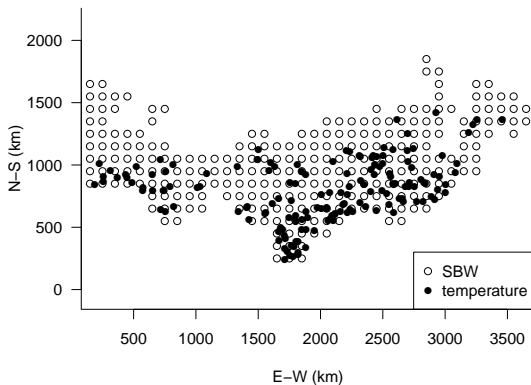
$$\frac{\partial N(\mathbf{x}, t)}{\partial t} = \underbrace{F(N(\mathbf{x}, t), E(\mathbf{x}, t))}_{\text{pop. growth}} - \underbrace{mN(\mathbf{x})}_{\text{emigration}} + \underbrace{m \int D(\mathbf{y}, \mathbf{x}) N(\mathbf{y}) d\mathbf{y}}_{\text{immigration}}$$

$$\frac{\partial n}{\partial t} \approx \underbrace{-rn(\mathbf{x}, t)}_{\text{regulation}} + \underbrace{m(D * n - n)}_{\text{redistribution}} + \underbrace{\sigma_E^2 e(\mathbf{x}, t)}_{\text{noise}}$$

$$2(r + m)c^* = m(D * c^*) + \sigma_E^2 \text{Cor}(e)$$

## Spatial synchrony

## Moth sampling locations

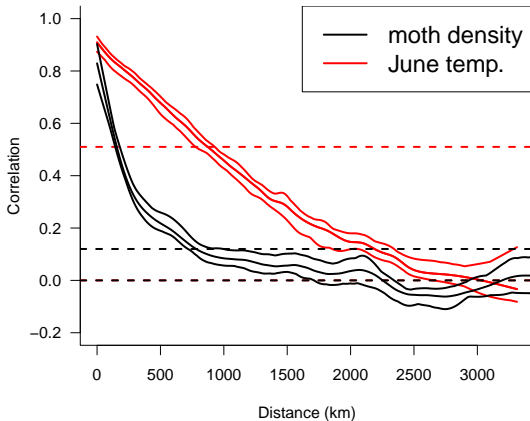




# Moth dynamics

## Spatial synchrony

## Spatial correlations of moth data



# Spatial logistic: solution

At equilibrium, the power spectrum of the population densities obeys

$$\tilde{S} = \left| \left( \tilde{N}^* \right) \right|^2 = \frac{\sigma_E^2 \tilde{e}}{2(r + m(1 - \tilde{D}))}$$

where  $\tilde{\cdot}$  denotes the Fourier transform. Therefore:

$$\sigma_P^2(p) = \sigma_E^2 + \frac{m}{r} \sigma_D^2$$

(Lande et al. 1999) where  $\sigma_X$  represents the standard deviation of the autocorrelation function

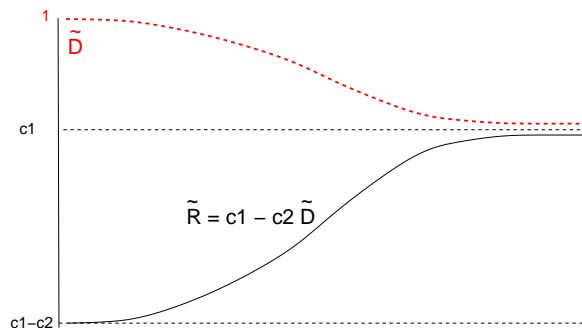
# Spectral ratios

To factor out dispersal, calculate the *spectral ratio*:

$$\tilde{R} = \frac{\tilde{e}}{\tilde{S}} = \frac{2}{\sigma_E^2}(r + m(1 - \tilde{D})) = c_1 - c_2\tilde{D}$$

*Deconvolve* the effects of environmental variability from the population pattern ...

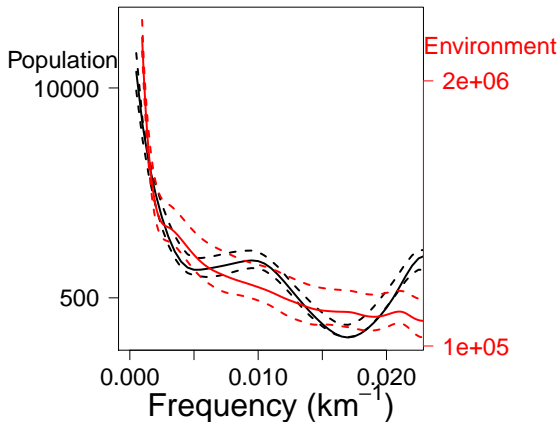
# Reconstructing the dispersal curve



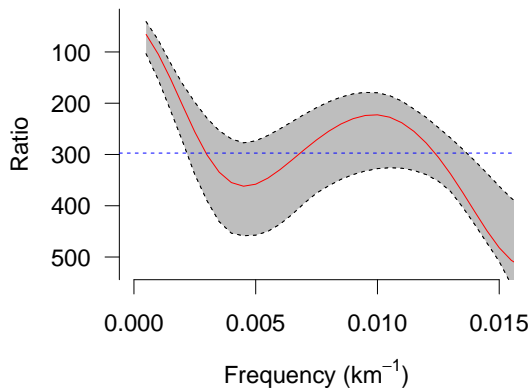
We know the limits  $\tilde{D}(0) = 1$ ,  $\tilde{D}(\infty) \rightarrow 1$ : thus

$$\tilde{D}_{\text{est}}(\omega) = \frac{\tilde{R}(\infty) - \tilde{R}(\omega)}{\tilde{R}(\infty) - \tilde{R}(0)}$$

# Moth data: spectra

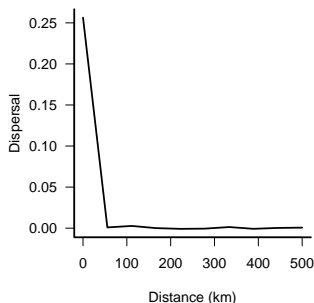


# Moth data: spectral ratios



## Spatial synchrony

## (Putative) moth dispersal curve



- Reconstructed moth dispersal kernel  $\approx 50\text{km}$
- Consistent with natural history



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# Pines

- Slash pine, *Pinus elliottii*
- Data on seed distribution, seedling distribution, but not collected on the same quadrats
- Sampling scheme highly irregular; sparse data



Wikipedia

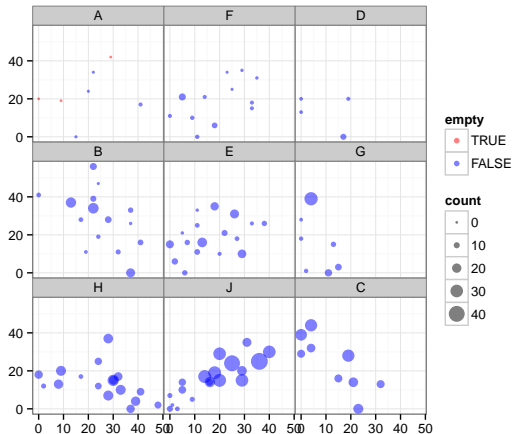
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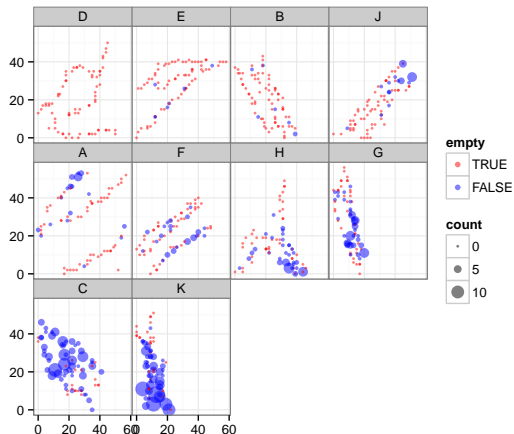
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Pines

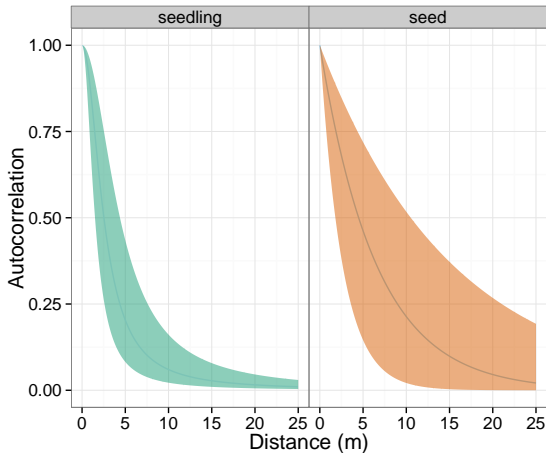
# Seed data



# Sapling data



# Correlation functions



## Analytical framework (2)

- assume *cross-covariance*  $C_{EN} = 0$   
no correlation between seeds and environment,  
e.g. long-distance dispersal



$$C_{SS}(r) = \bar{N}^2 C_{EE}(r) + \bar{E}^2 C_{NN}(r)$$

(where  $\bar{N}$ =mean seed density,  $\bar{E}$ =mean establishment probability)

- or (switch to correlation  $c$ )

$$C_{SS} \propto \frac{\sigma^2 E}{\bar{E}^2} c_{EE} + \frac{\sigma_N^2}{\bar{N}^2} c_{NN}$$

- ...a weighted mixture of the two correlation functions

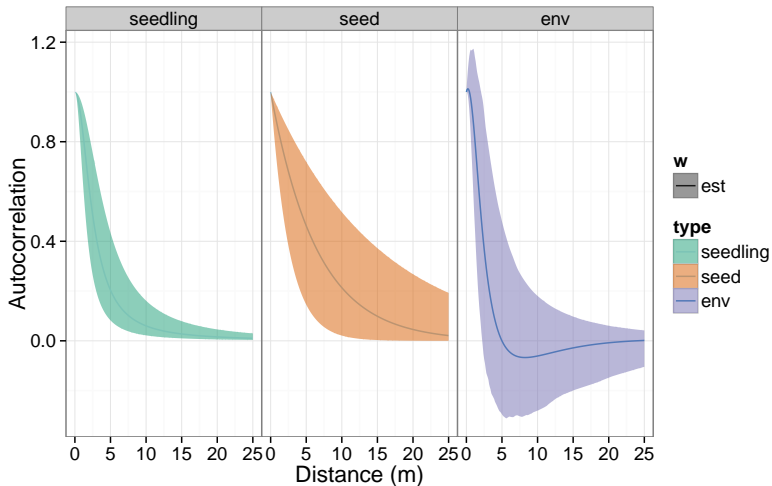
# Solving for $c_{EE}$

Therefore,

$$c_{EE} \propto \sigma_S^2 c_{SS} - \bar{E}^2 \sigma_N^2 c_{NN}$$

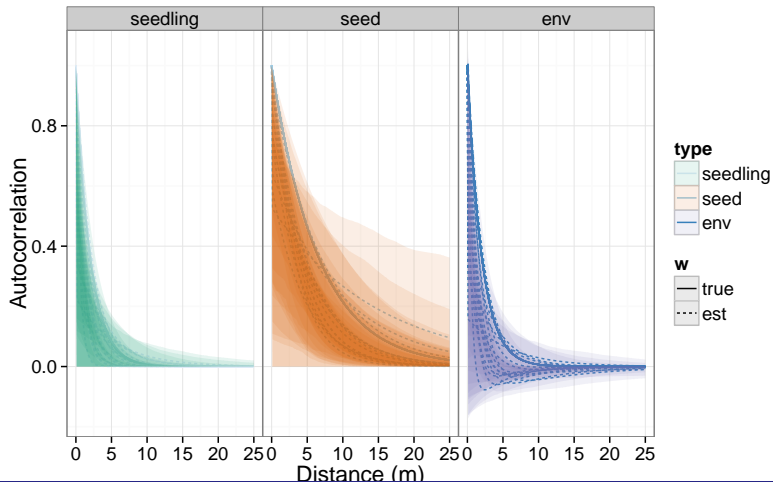
Can we really use this?

# Results: observed/inferred correlation equations





# Simulation results



# Caveats/assumptions

- Assumes linearization/moment truncation
- Assumes isotropy/homogeneity
- Estimating spectra of small, irregular, noisy data sets is difficult (!)
- Advantages vs. direct estimation (e.g. via MCMC) ?

- Simple, light-weight (!!), non-parametric (??) approach to spatial estimation
- leverage “unreasonable effectiveness” of linearization (Gurney & Nisbet, 1998)
- Use all available information:
  - snapshots, before/after, time/series
  - non-matching spatial samples

# Acknowledgements

**People** Ottar Bjørnstad, Sandy Liebhold, Aaron Berk, Gordon Fox, Stephen Cornell, Mollie Brooks, Emilio Bruna

**Funding** NSF, NSERC (Discovery Grant)

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