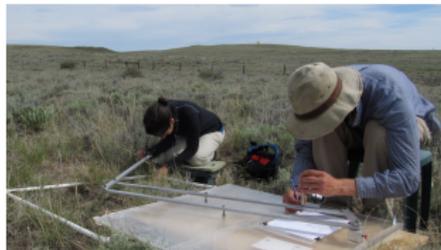
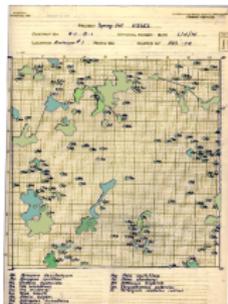


# IDEs as models for individuals: who gets into the 1%, and why?

Stephen Ellner  
Ecology and Evolutionary Biology, Cornell



# Collaborators



Peter Adler, Utah State



Robin Snyder, Case Western



Giles Hooker, Cornell



Brittany Teller, Utah State



Mark Rees, Sheffield



Dylan Childs, Sheffield



National Science Foundation  
WHERE DISCOVERIES BEGIN

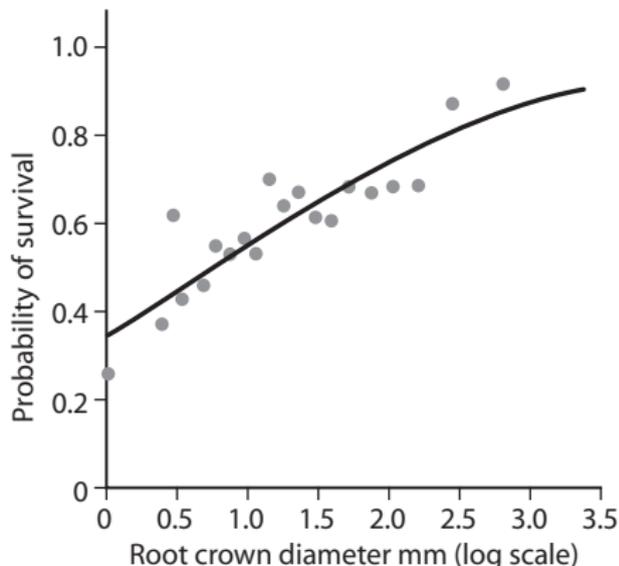
# IDEs for movement in “trait space”

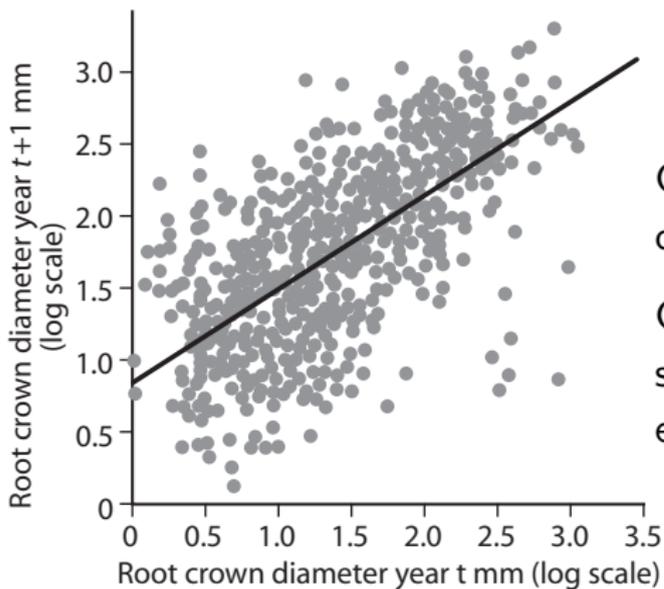
Plants & other organisms with indeterminate growth: size is most important trait.

Size varies continuously: effects of size described by regression models.



Platte thistle, *Cirsium canescens* (Rose et al. 2005)





Growth  $\sim$  size: dynamic model for changes in individual state.

Can be: nonlinear, non-Gaussian, size-dependent variance, random year effects, etc.

# Integral Projection Model (IPM)

$n(z, t)$  = distribution of individual state,  $z$

Transitions  $z$  (now) to  $z'$  (next census) described by

$$K(z', z) = \underbrace{P(z', z)}_{\text{Survival/growth}} + \underbrace{F(z', z)}_{\text{Reproduction}}$$

$$n(z', t + 1) = \int_{\mathbf{Z}} K(z', z) n(z, t) dz$$

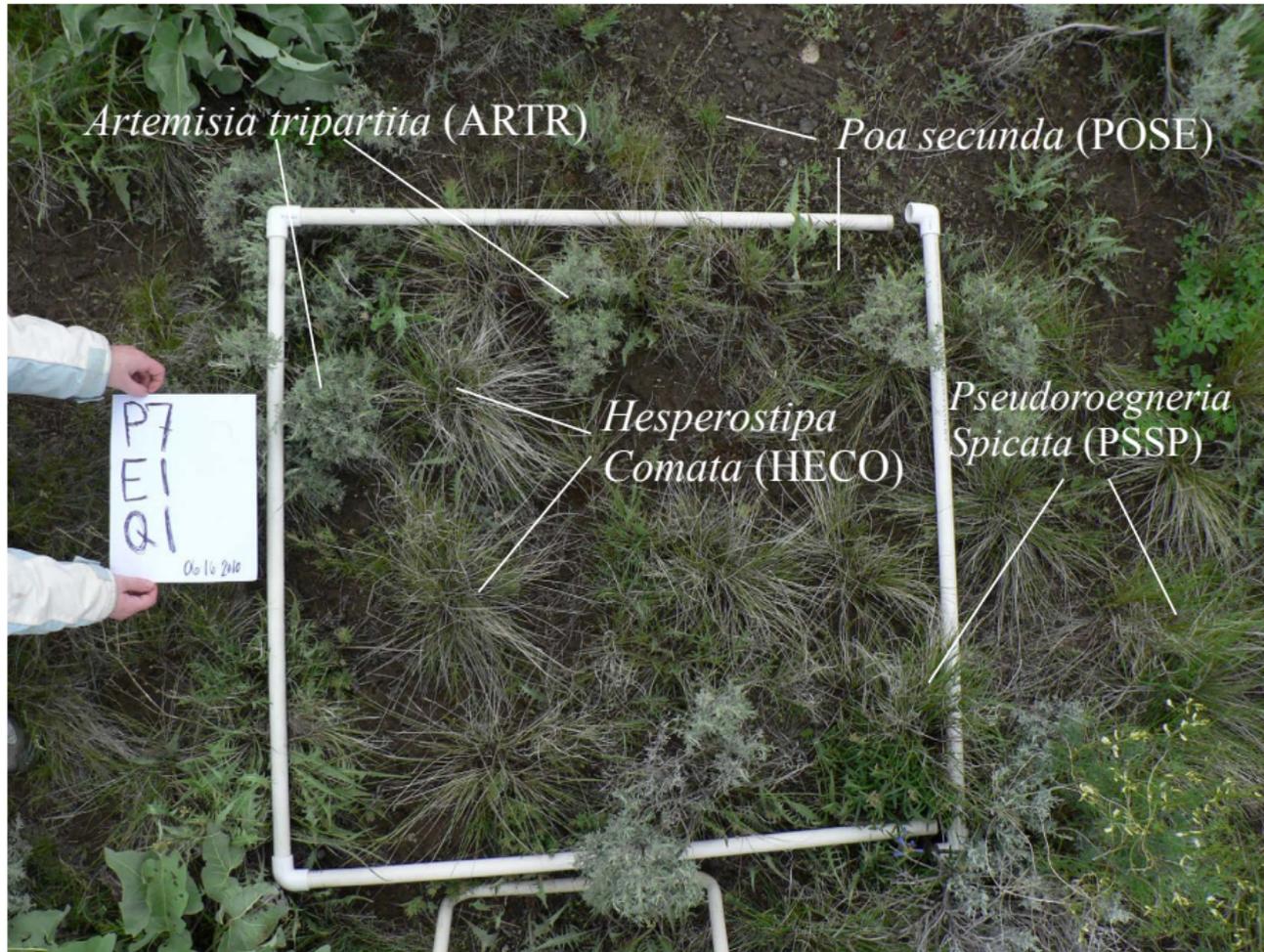
NOTE: ' always means "value at next census"

NOTE: trait space  $\mathbf{Z}$  compact.

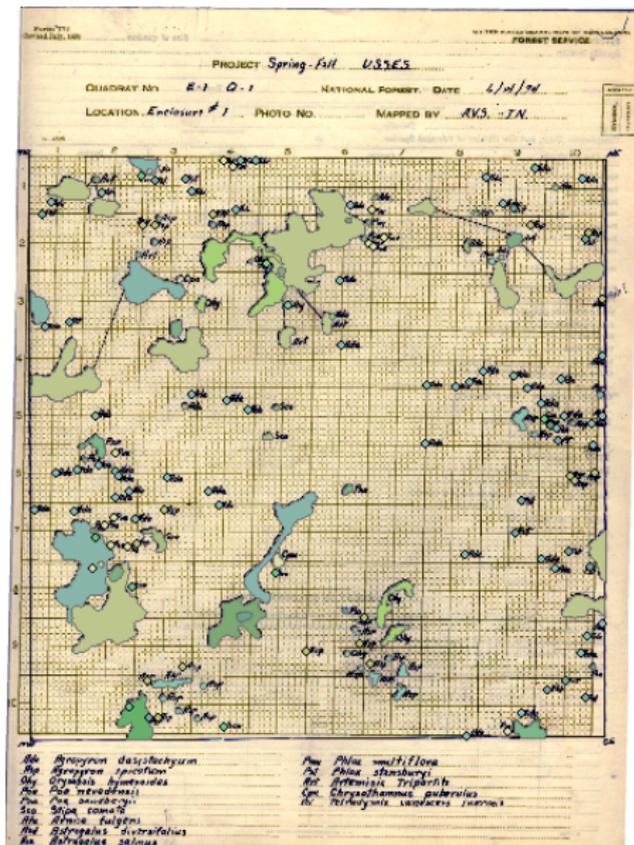
# Idaho sagebrush steppe

*That's where the data are:* 26 quadrats established 1926-1932 at US Sheep Expt. Station, mapped most years until 1957 (22 annual transitions)

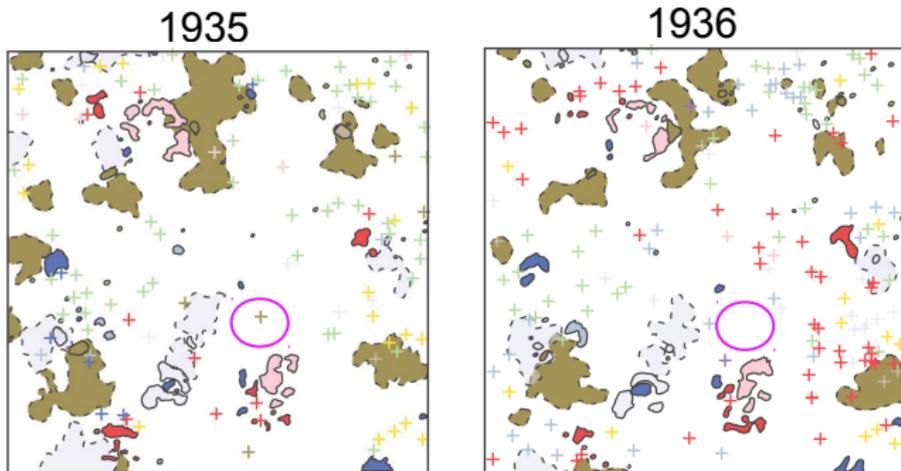




# Hand-drawn maps...



# ...digitized by Peter Adler to GIS shapefiles



For each plant, we know:

- size
- survival
- growth
- location, size, species of competitors
- daily temperature, rainfall, snowfall

Survival and growth fitted as regression models of individual area, site covariates (grazed/ungrazed, etc.), and competitive pressure  $W$ .

Competitive pressure  $W$  = the sum over all neighboring plants of

Size of neighbor

- × “competition coefficient”  $\alpha_{ij}$   
(effect of species- $j$  neighbor on species- $i$  focal plant)
- × “competition kernel” (near vs. far neighbors)

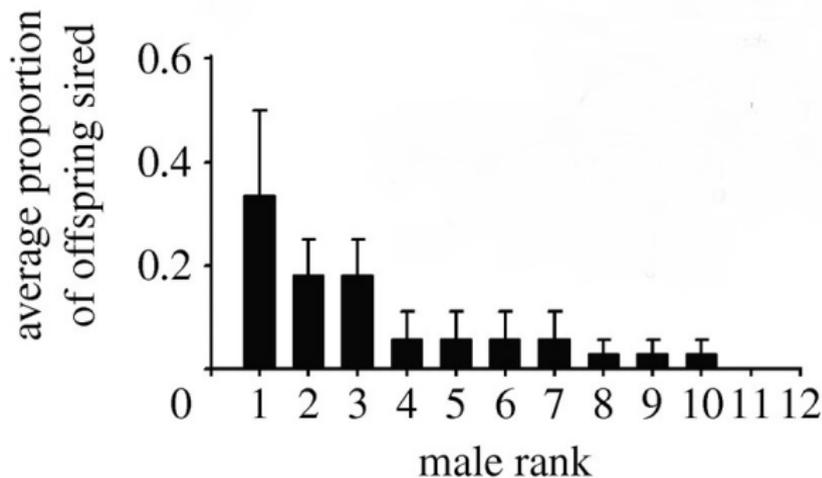
(B. Teller et al. 2016, *Methods in Ecology and Evolution*)

## Demographic variability among individuals

Reproductive skew:

A few parents have many offspring  
most parents have few offspring.

Wire-tailed manakin *Pipra filicauda* (Ryder et al. 2009, *Proc. RSL B*)





“Canopy” plants are the lucky few.

MANY seedlings/yr,  
FEW become full-size adult.

R. Snyder and SPE (Am. Nat. 2016): who becomes one of lucky few plants, and why?

Survival/growth kernel  $P$  is Markov chain for individual paths through life (add death as absorbing state)

⇒ Many life-cycle properties can be computed!

R. Snyder and SPE (Am. Nat. 2016): who becomes one of lucky few plants, and why?

Survival/growth kernel  $P$  is Markov chain for individual paths through life (add death as absorbing state)

⇒ Many life-cycle properties can be computed!

$$\text{Var}(\text{lifespan}|\text{state at birth}) = \mathbf{e}(2N^2 - N) - (\mathbf{e}N)^2$$

where  $N = (I - P)^{-1}$ ,  $\mathbf{e} \equiv 1$ .

R. Snyder and SPE (Am. Nat. 2016): who becomes one of lucky few plants, and why?

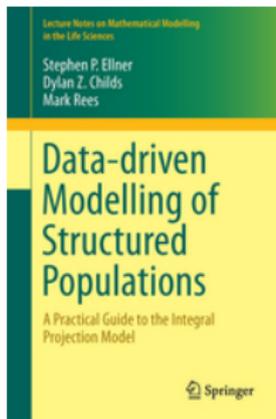
Survival/growth kernel  $P$  is Markov chain for individual paths through life (add death as absorbing state)

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where  $N = (I - P)^{-1}$ ,  $\mathbf{e} \equiv 1$ .

MANY more: download (free?)  
and see chapter 3.



Who becomes one of the lucky few, and why?

- 1 Compute the *conditional* transition kernels for the Lucky and Unlucky.
- 2 Compare these to ask: when and how do paths diverge?

Possible absorbing sets  $A_1, A_2, \dots, A_M$ :

- Process conditional on absorbing into  $A_j$  is Markov chain
- Easy to compute  $\text{Prob}(\text{absorb into } A_j), P(z', z | \text{absorb into } A_j)$

Define the “Lucky” absorbing set:

- Size at death is  $\geq z^*$
- Maximum size (at any age) is  $\geq z^*$
- Lifetime total # offspring<sup>1</sup>  $\geq T^*$
- Lifetime total # breeding times  $\geq T^*$
- and so on...

<sup>1</sup>state  $z = (\text{size}, \# \text{kids so far})$

# Examples

*Dacrydium elatum*, tropical tree

Deterministic growth.

Eelke Jongejans et al. (2010),  
Journal of Ecology



# Examples

*Artemisia ordosica* on  
Mongolian sand dunes

Variable growth.

S-L Li et al. (2011),  
Journal of Ecology



# Examples

*Cedrela odorata* (Spanish cedar),  
tropical tree

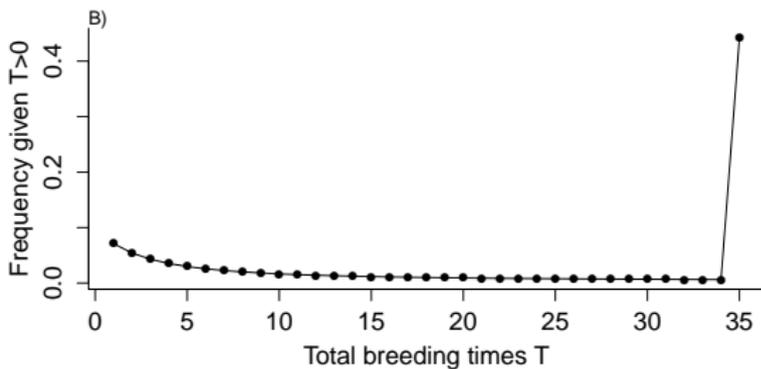
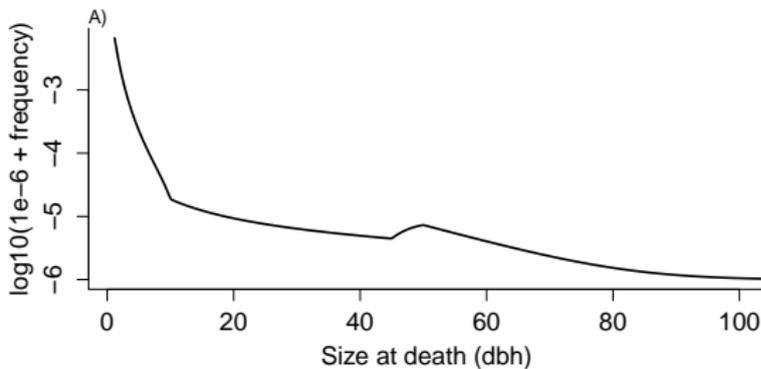
Variable growth; differences  
among individuals are persistent.

Zuidema et al. (2009): the  
“lucky few” are consistently  
fast-growing trees.

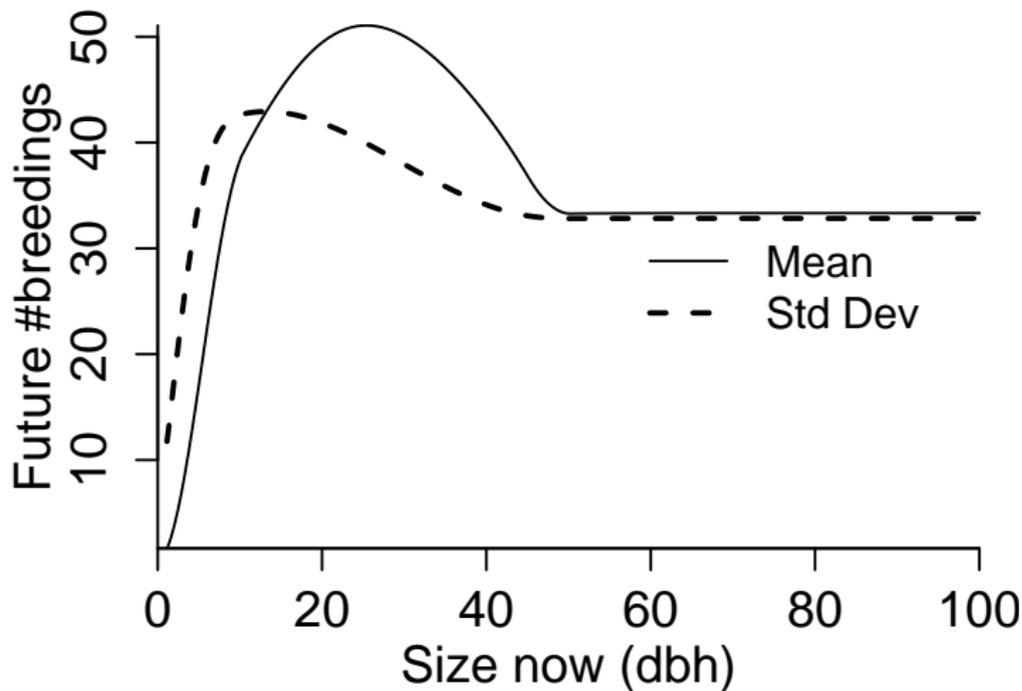


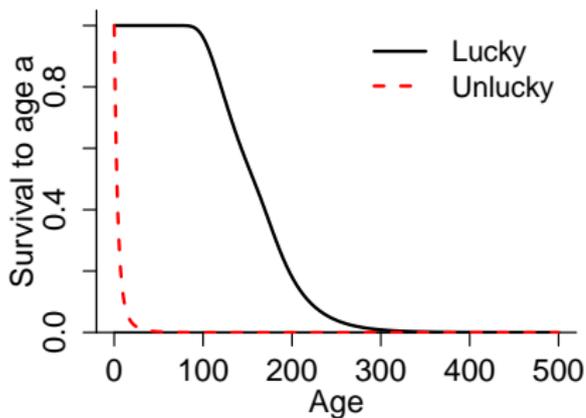
# Dacrydium elatum: the 1%

Size at death  
and LRS are  
bimodal

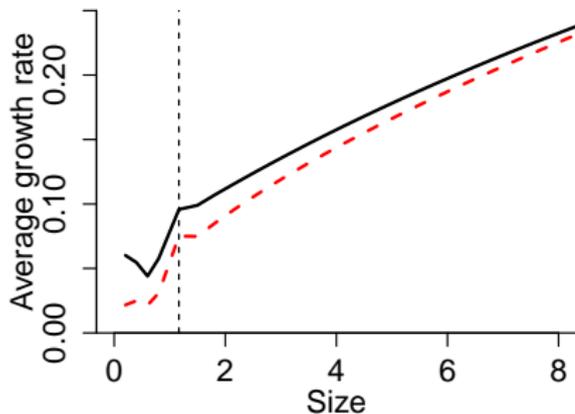


# Dacrydium elatum: Lucky=20cm dbh

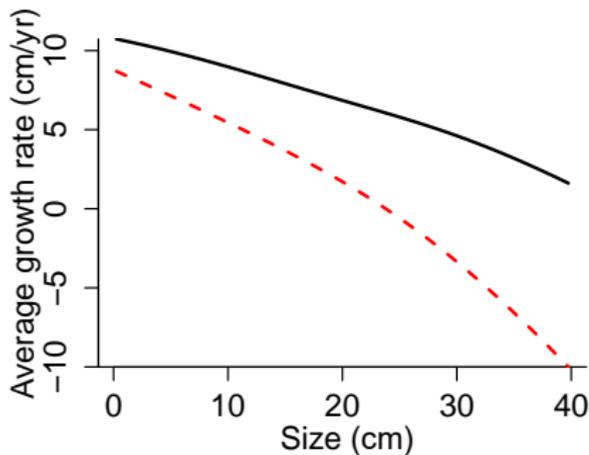
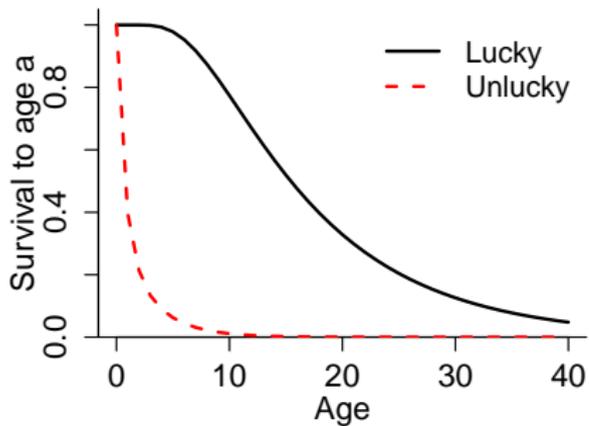




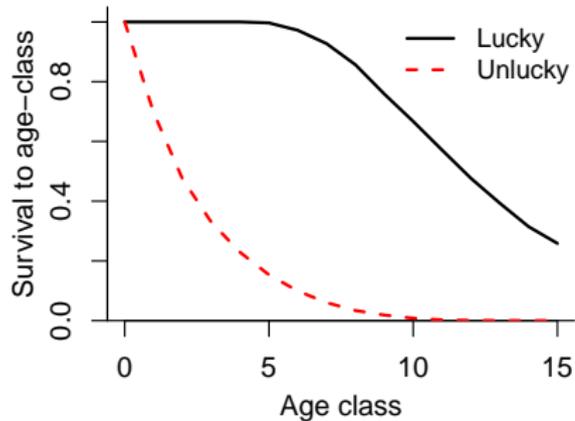
*Dacrydium*: Enormous survival differences.



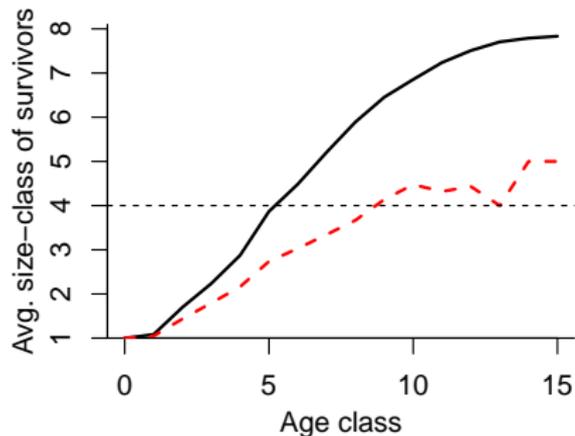
Tiny growth differences except in seedlings – where it aids survival.



*Artemisia* (Lucky = 40cm)  
Same story.



*Cedrela* (Lucky = size class 6)



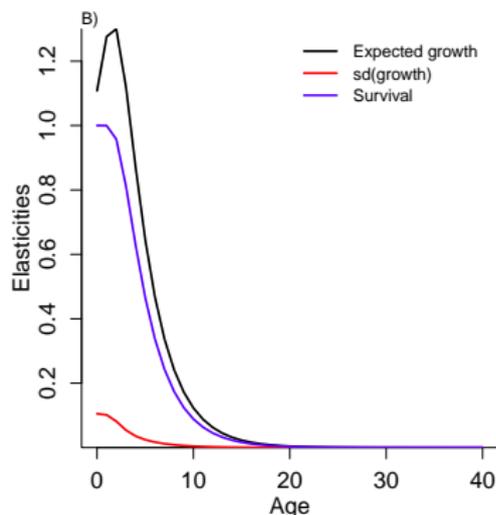
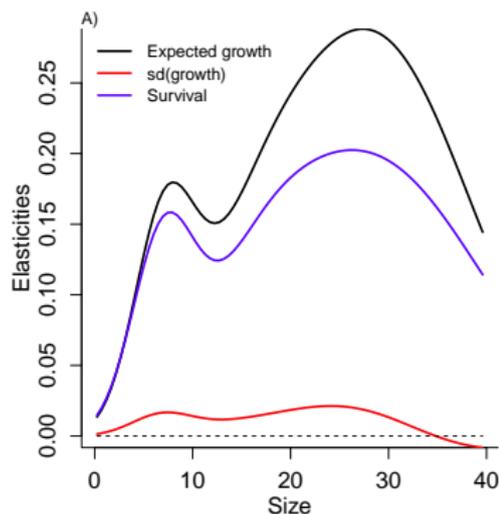
Big growth divergence:  
big enough to get  
Lucky in one time-step.

To a large extent, the Lucky are just those who do not die early.

# What kind of lucky break is most helpful?

Sensitivity of  $q_{\mathbb{L}} = P(\text{die Lucky})$  to perturbation at age  $a$ :

$$\frac{\partial q_{\mathbb{L}}}{\partial \phi} = \frac{\partial a_{\mathbb{L}}}{\partial \phi} P^a + a_{\mathbb{L}} N \frac{\partial P}{\partial \phi} P^a, \quad N = (I - P)^{-1}.$$



## Traits vs. luck (R. Snyder & SPE, in prep)

Ecologists want to believe that there is a reason for large differences in reproductive success.

- To what extent can trait differences override luck?
- Do these determine who joins the Lucky few, or is it still mostly luck?

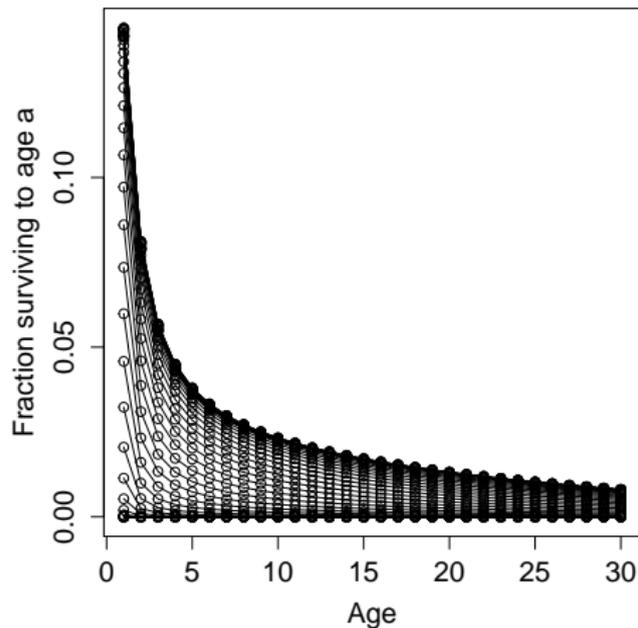
Motivating example (Idaho): how much does it matter if a seedling germinates in a good spot or a bad spot?

Good = no close conspecifics!

# Artemisia: site quality matters

Site quality:  $W_1$ , competition at first census

ARTR: survival for different  $W_1$



Good site: some chance of surviving to age 10, large and fecund.

Poor site: no.

## Partitioning variance in LRS: Trait vs. Luck

$R$  = Lifetime Reproductive Success (LRS)

$W_1$  = site quality at birth,  $Z$  = size at age 2.

## Partitioning variance in LRS: Trait vs. Luck

$R$  = Lifetime Reproductive Success (LRS)

$W_1$  = site quality at birth,  $Z$  = size at age 2.

$Var(LRS) =$

$$\underbrace{\mathbb{E}[s(W_1) Var(R^*|W_1, Z)]}_{\textcircled{1}} + \underbrace{\mathbb{E}[s(W_1)(1 - s(W_1))(\mathbb{E}(R^*|W_1, Z))^2]}_{\textcircled{2}} \\ + \underbrace{\mathbb{E}_{W_1}[s(W_1)^2 Var_Z E(R^*|W_1, Z)]}_{\textcircled{3}} + \underbrace{Var_{W_1}[\mathbb{E}(R|W_1)]}_{\textcircled{4}}$$

where  $R^*$  = LRS conditional on surviving to age 2.

$Var(LRS) =$

- 1 Effect of trait variation (site quality)
- 2 Components of luck
  - a Do you survive to age 2?
  - b How big are you, if you survive?
  - c Variation independent of state at age 2.

## Results: Luck dominates!

	<i>Artemisia</i>	<i>Pseudoroegneria</i>
Trait: $W_1$	0.2%	0.5%
Luck: survive to 2?	11%	12%
Luck(?): size at 2	0.8%	1.4%
Luck: later variation	88%	86%

# Not just shrubs in Idaho!

## Kittiwake *Rissa tridactyla*



# Not just shrubs in Idaho!

Emmanuele Cam, University of Toulouse III



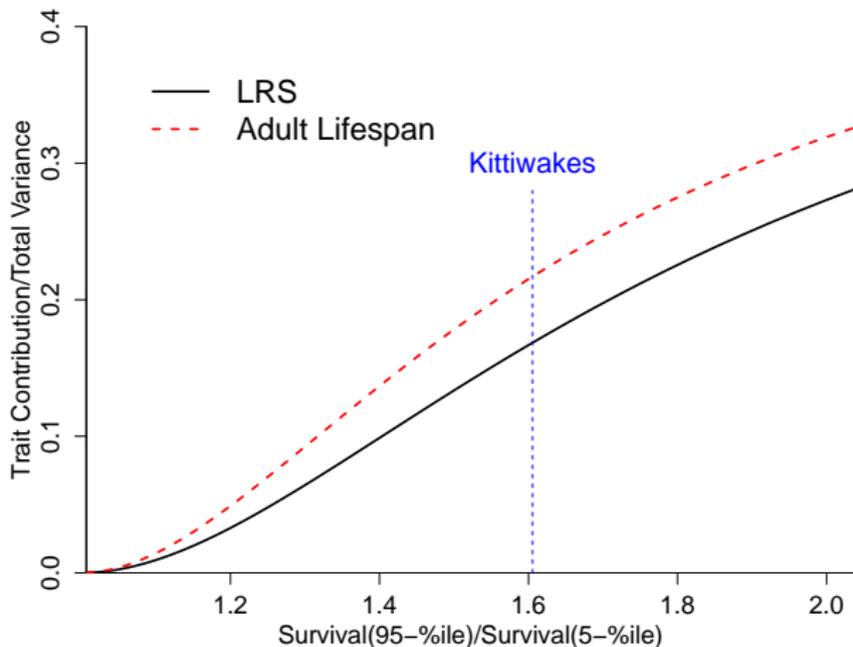
*The idea of differences in individual quality has been put forward ...to explain differences in lifetime production among individuals. (Cam et al. 2004, Oikos)*

In kittiwake adults (Cam et al. 2002, *American Naturalist*)

Survival probability,  $CV \approx 0.2$  ( $CV = \frac{\sigma}{\mu}$ )

Breeding probability,  $CV \approx 0.1$

## But variation in LRS is still mostly luck!



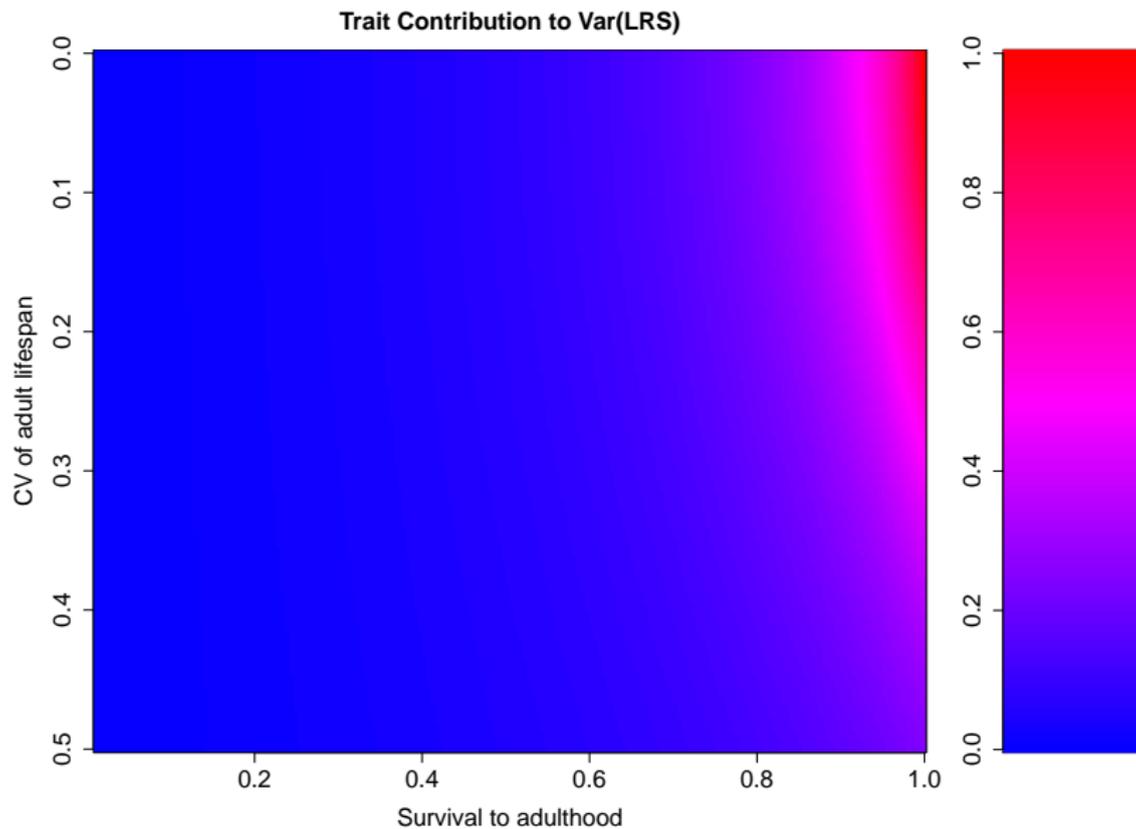
Trait variation +  
kittiwake model from  
Steiner & Tuljapurkar  
(2012)

Bonnet & Postma  
(2016), snow voles

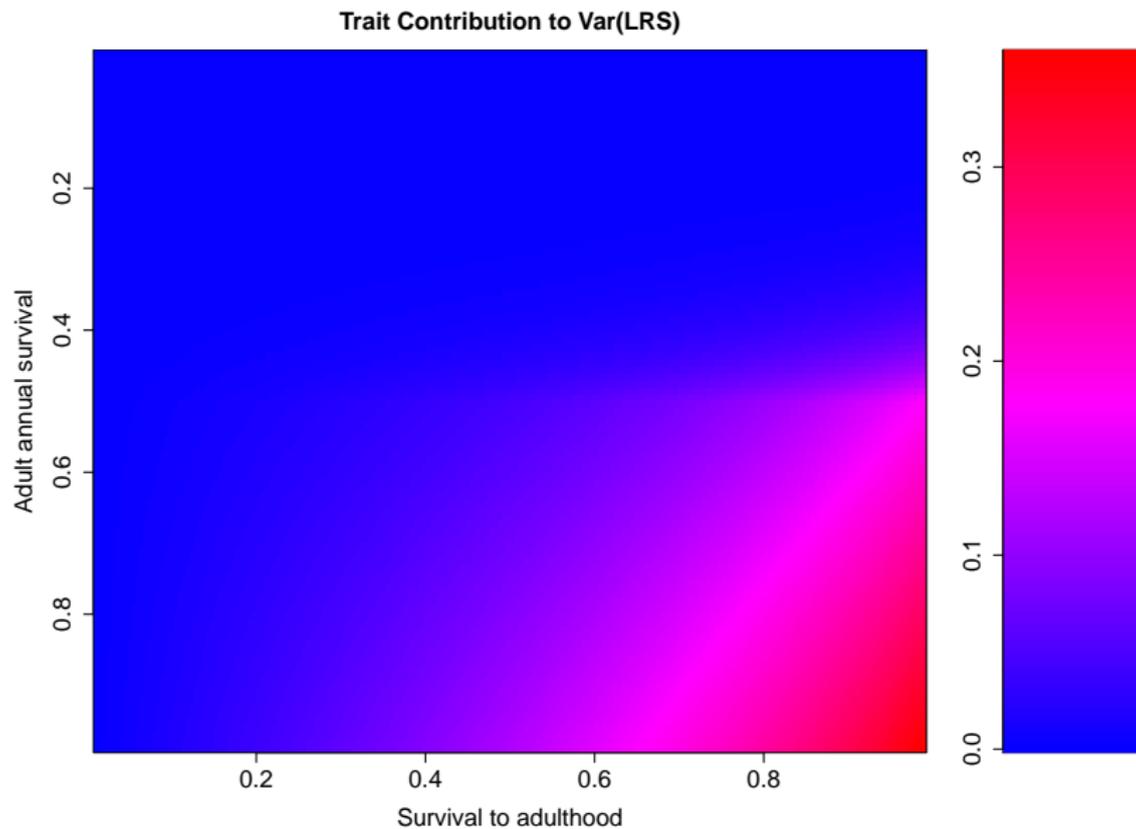
## Is LRS always dominated by luck? (R.Snyder & SPE)

- Two life stages, Juvenile and Adult
- Juvenile survival  $s_J$ , adult lifetime  $\tau$  (random),  $F$  offspring/yr on average.
- Stable population at trait mean:  $R_0 = 1$ .
- $CV = 0.3 \Rightarrow$  3-fold ratio between 95<sup>th</sup> and 5<sup>th</sup> percentiles of a Gaussian trait distribution with positive mean.

Trait is offspring/yr (deterministic),  $CV=0.3$



Trait is adult survival ( $CV=0.3$ ), constant clutch size

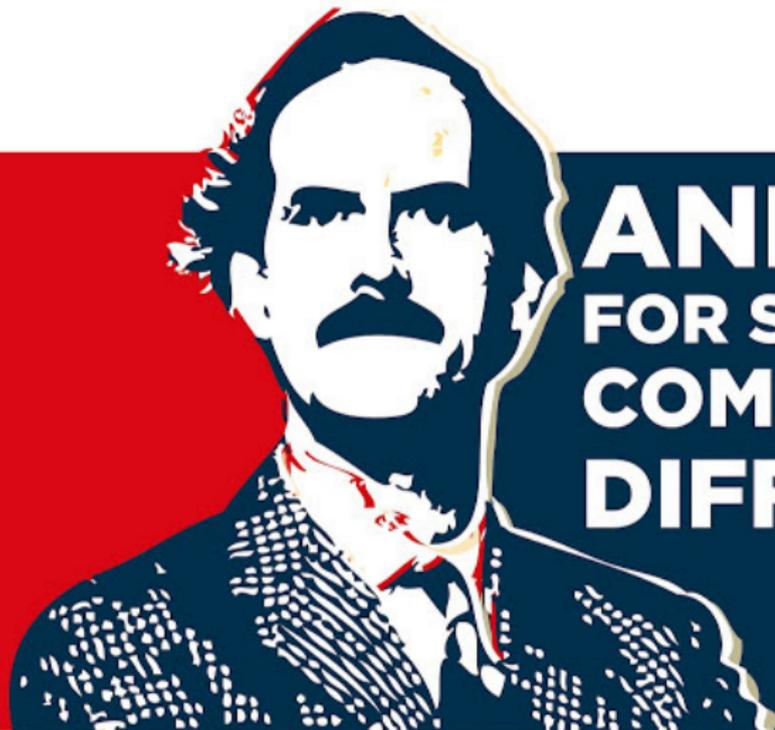


Trait variation accounts for under 10% of total variation in LRS, so long as trait  $CV < 1$ .

Trait variation dominates  $\text{Var}(\text{LRS})$  if survival to adulthood is high, and trait effectively is adult LRS.

- $\text{LRS} \approx \text{Lifespan} \times \text{Mean clutch size}$ .
- If one of those is  $\approx$ constant, and the other one is the trait, then trait variation dominates LRS.

Otherwise, so many forking paths through life that becoming big and fecund is mostly *sheer dumb luck*.



**AND NOW  
FOR SOMETHING  
COMPLETELY  
DIFFERENT.**

Typically we model transitions using densities: transition probabilities are absolutely continuous w.r.t. underlying measure.

For trait space, not always reasonable:

- Deterministic transitions of some trait components (genotype, breeding value).
- Constraints: allocations to growth + reserves = total energy intake.

Resulting models are very different, **very little theory**.

Simplest possible case:

- $z = (x, y) \in \mathbf{Z} = \mathbf{X} \times \mathbf{Y}$
- $x$  has deterministic transitions  
 $x' = \omega(x)$ ,  $\omega$  smooth with smooth inverse  $\alpha$
- $y$  has smooth stochastic transitions:  
 $P(y'|x, y) = s(x, y)G(y'|x, y)$

$$n(x', y', t + 1) = \int_{\mathbf{Z}} F(x', y', x, z)n(x, y, t) dx dy$$
$$+ 1_{\omega(\mathbf{X})}(x') |\alpha'(x')| \int_{\mathbf{Y}} G(y'|x'_*, y)s(x'_*, y)n(x'_*, y, t) dy$$

where  $x'_* = \alpha(x')$ ,  $n \equiv 0$  off  $\mathbf{Z}$

Example:

Individuals classified by size  $y$ , site quality  $x \geq 0$  that decreases over time.

$$x' = \delta x, \quad 0 < \delta < 1$$

$$\text{Growth: } y' \sim \text{Normal}(\mu = x + 0.9y, \sigma = \sigma_g)$$

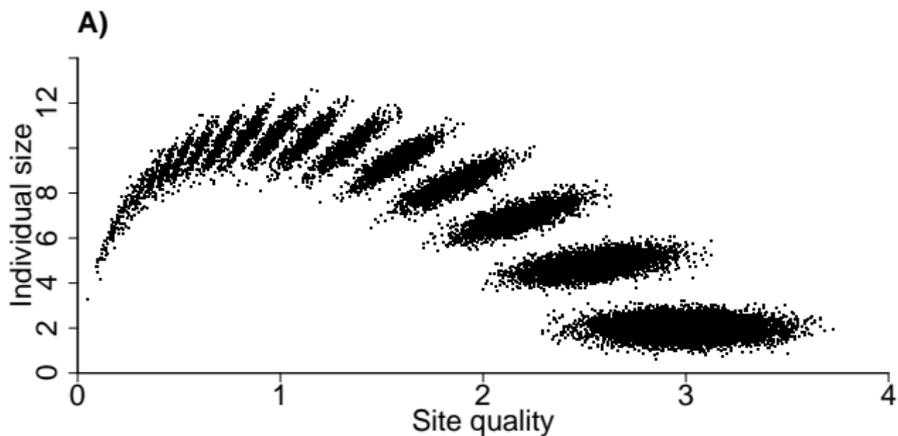
$$\text{Survival: } \text{logit } s(y) = A + By, B > 0$$

$$\text{Number of offspring: } b(x, y) = by$$

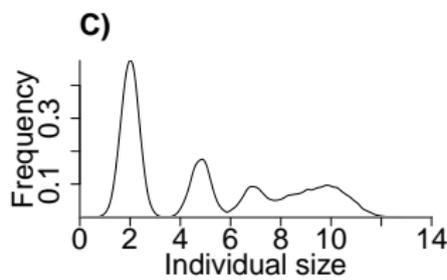
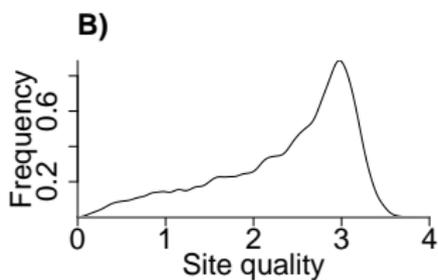
$$x \text{ at birth} \sim \text{Normal}(\mu_x, \sigma_x)$$

$$y \text{ at birth} \sim \text{Normal}(\mu_y, \sigma_y)$$

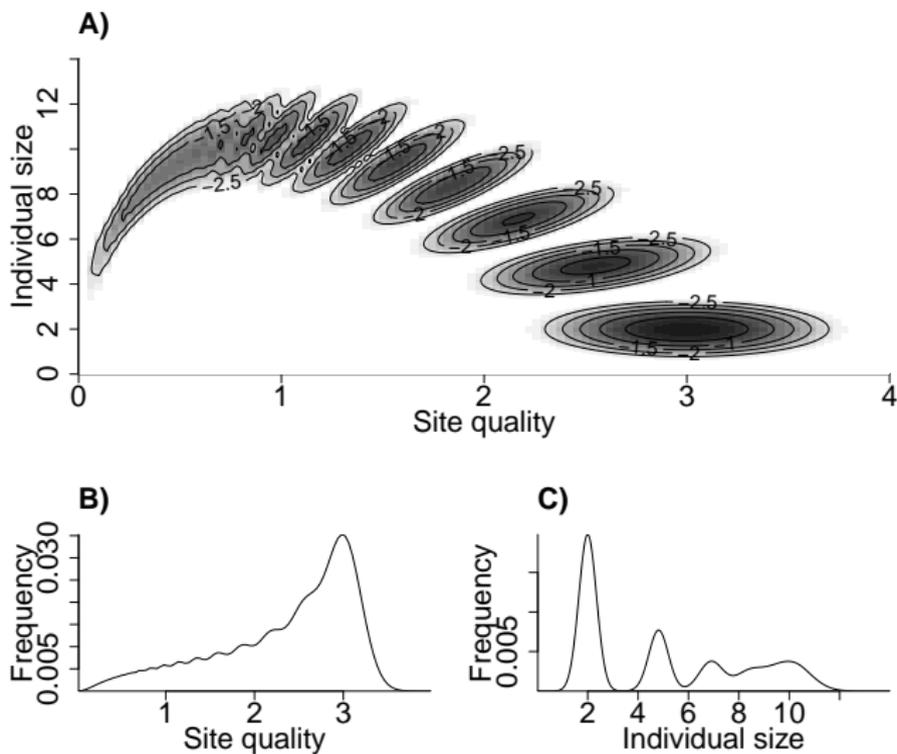
# Agent-based simulations (to stable distribution)



Individuals grow at first but then shrink, because their site quality has dropped.



# Numerical iteration of IPM



Individuals grow at first but then shrink, because their site quality has dropped.

$$n(x', y', t + 1) = \phi(x'; \mu_x, \sigma_x) \phi(y'; \mu_y, \sigma_y) \int_{\mathbf{Z}} byn(x, y, t) dydx \\ + \frac{1}{\delta} \int_{\mathbf{Y}} \phi(y'; x'/\delta + 0.9y, \sigma_g) s(y) n(x'/\delta, y, t) dy$$

- What function space does this “live on”?
- Does  $n$  stay smooth, or can it develop “shock waves”?
- All the basic theory: stable stage distribution, eigenvalue sensitivity formulas, etc.

- ① IDEs are models for the lives of individuals. We can extract from them much more information than we have been.
- ② IDEs based on individual-level processes may not have transition densities. We know very little about such models.



Questions?

Graduate diversity recruitment weekend, E&EB Cornell and NB&B Cornell  
April 21-23, 2007

An event to connect students from under-represented backgrounds with faculty, before they apply for graduate school.

For college Junior or Seniors, or students who have graduated and are considering graduate school.

Application at [cudiversityrecruitment.weebly.com](http://cudiversityrecruitment.weebly.com)

Application deadline December 1, 2016.

Attendees will get \$400 for travel, housing/meals for the weekend.

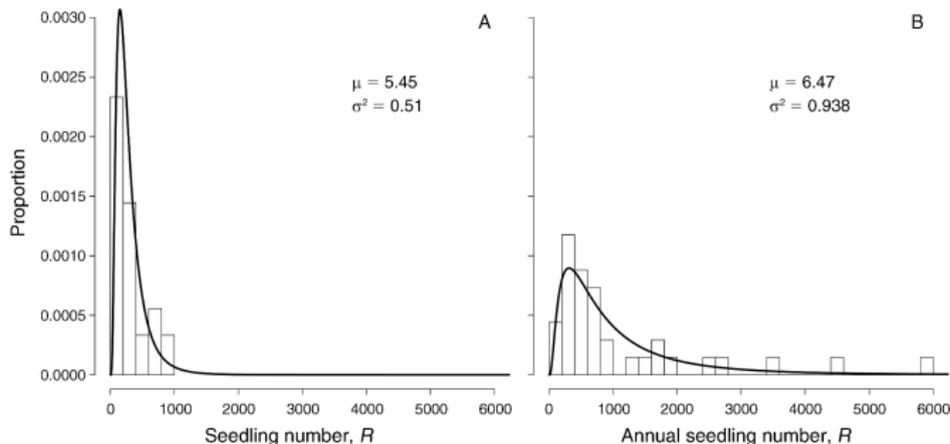


# Even among the lucky few, LOTS of skew

1088

EMILY V. MORAN AND JAMES S. CLARK

Ecology, Vol. 93, No. 5



Seedlings/yr, trees (Moran & Clark 2012)

(A) Duke Forest: top 5% make 29% of seedlings

(B) Coweeta: top 5% make 47% of seedlings

Competition kernel: splines (B. Teller et al. 2016, Methods in Ecology and Evolution)

