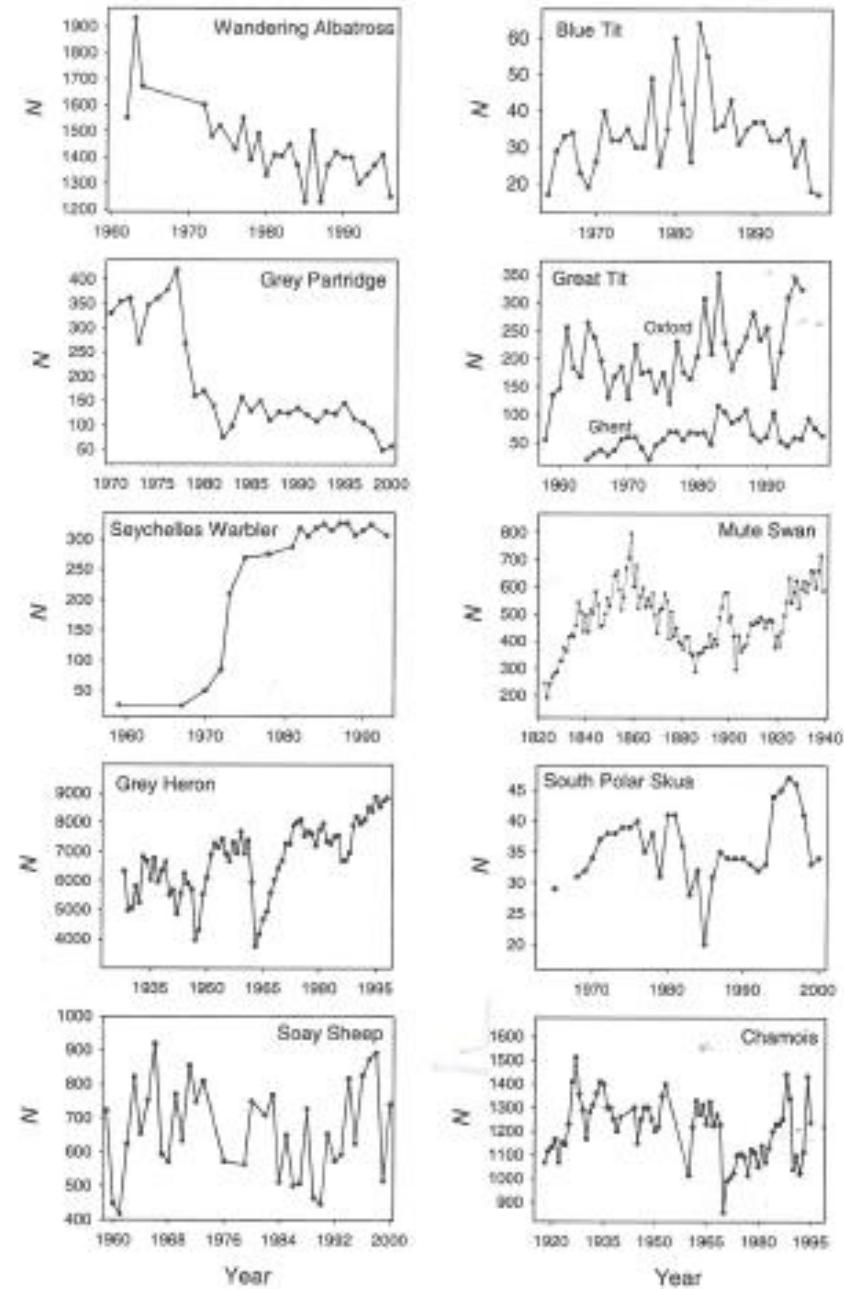




# Stochastic population modeling

Ola Diserud  
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# Some time-series



## 1.3 Lognormal distribution

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Multiplicative effects common in biological processes -> lognormal distribution important

Central limit theorem:

$\sum_{u=0}^{t-1} S_u$  approx. normally distributed

->  $X_t = X_0 + \sum_{u=0}^{t-1} S_u | X_0$  also approx. normal

->  $N_t = N_0 e^{\sum S_u} | N_0$  approx. lognormal

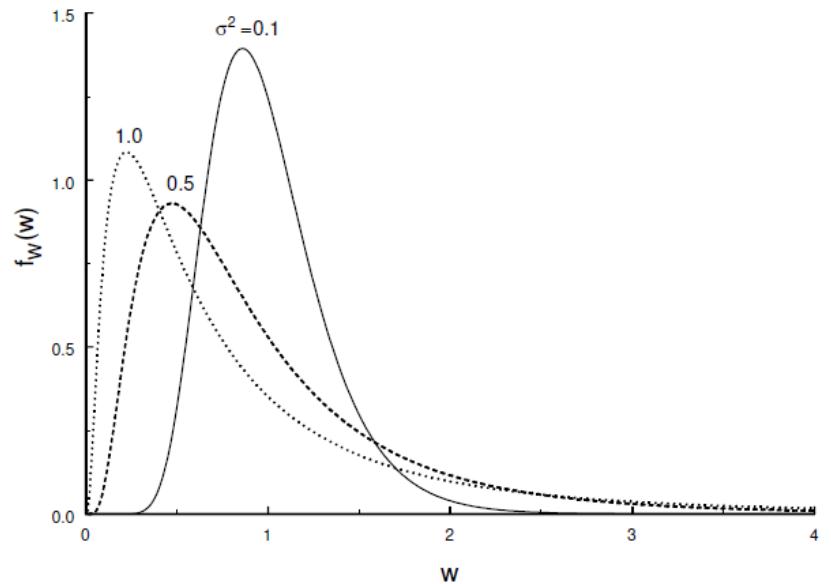
# Lognormal distribution

$$Y \sim N(\mu, \sigma^2)$$

$$Y = \ln V \quad V = e^Y$$

$$V \sim LogN(\mu, \sigma^2)$$

$$f_V(v) = \frac{1}{\sqrt{2\pi}\sigma v} e^{-\frac{1}{2}\left(\frac{\ln v - \mu}{\sigma}\right)^2}, v > 0$$



# Stochastic growth rate

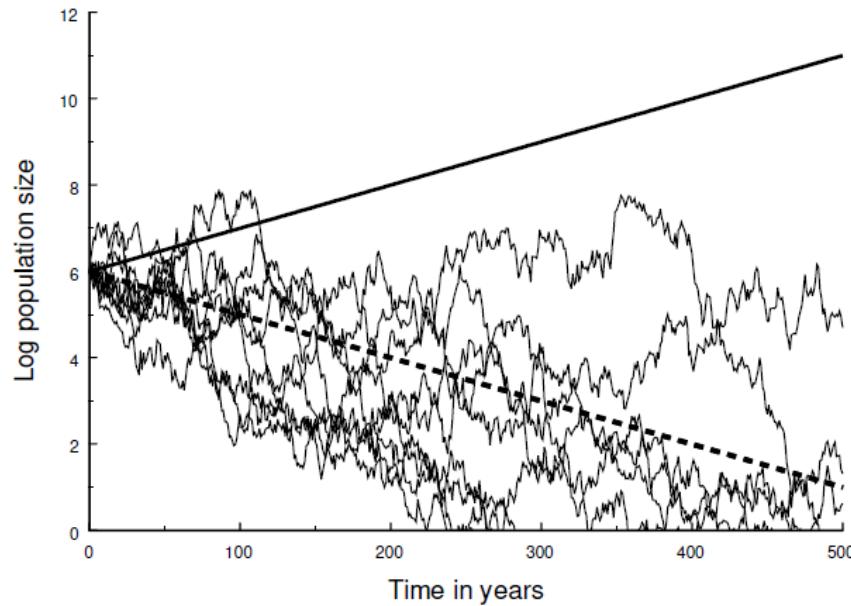


Figure 1.2: Simulation of 10 sample paths using the above stochastic model with  $r = 0.01$  and  $\sigma_e^2 = 0.04$ . The solid straight line shows the deterministic growth, while the dotted line is the mean stochastic growth.

# MLE of $s$ ?

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$$\hat{s} = \bar{D} = \frac{1}{n} \sum_{i=1}^n D_i = \frac{X_n - X_0}{n}$$

$$\hat{\sigma}_s^2 = \frac{1}{n-1} \sum_{i=1}^n (D_i - \hat{s})^2$$

Distribution?

$$\frac{\hat{\sigma}_s^2 (n-1)}{\sigma_s^2} \sim ? \quad \chi^2_{df=n-1}$$

# Prediction

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$$\frac{X_{n+m} - X_n - m\hat{s}}{\sigma_s \sqrt{m + m^2 / n}} \sim N(0, 1)$$

$$\frac{\hat{\sigma}_s^2 (n-1)}{\sigma_s^2} \sim \chi^2_{n-1}$$

$$\Rightarrow T = \frac{X_{n+m} - X_n - m\hat{s}}{\hat{\sigma}_s \sqrt{m + m^2 / n}} \sim T_{df=n-1}$$

# Prediction interval

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$$P\left(-t_{n-1,\alpha/2} < T_{n-1} < t_{n-1,\alpha/2}\right) = 1 - \alpha$$

$(1 - \alpha)$  prediction interval for  $X_{n+m}$

$$X_n + m\hat{s} \pm t_{n-1,\alpha/2} \hat{\sigma}_s \sqrt{m + m^2 / n}$$

# Figure 1.3

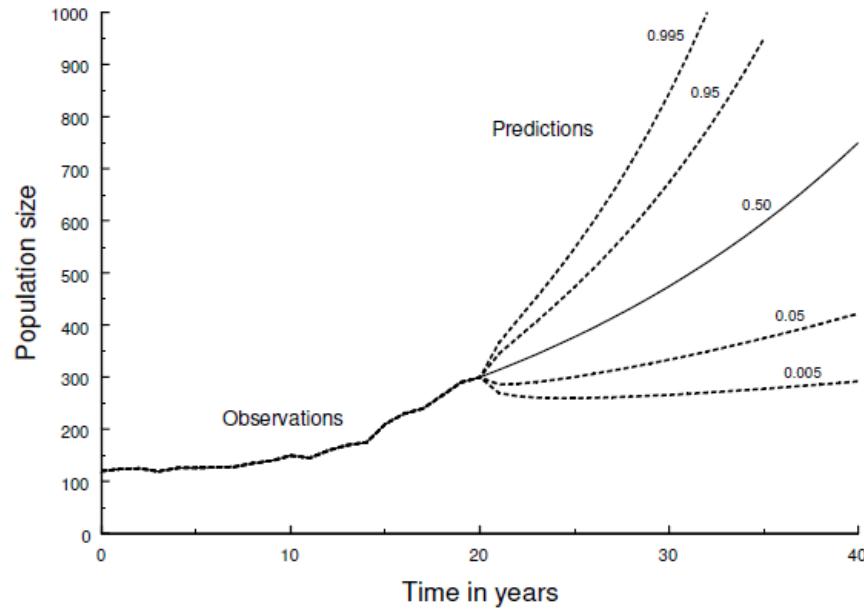


Figure 1.3: Time series observations of a population of Sea Eagles over 21 years together with 20 prediction intervals (90 and 99 %) and predicted median 20 years ahead.

# 1.6 Demographic stochasticity

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- ▶ Assume environment ( $Z$ ) constant – just look at differences between individuals, and
- ▶ populations with only adult individuals with the same stochastic properties
- ▶  $N$  individuals (females) then contribute with  $w_1, w_2, \dots, w_N$  to next year, where
  - ▶  $w = \# \text{offspring} + I$ ,  $I = \begin{cases} 1 & \text{survive} \\ 0 & \text{die} \end{cases}$
  - ▶  $w = \text{individual fitness}$
- ▶ Demographic variance  $\text{Var}(w_i) = \text{Var}(d_i) = \sigma_d^2$

# Conditionality

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1.  $E_Y \left[ E_x (X | Y) \right] = E_X (X)$
2.  $Var(X) = E_Y Var_X (X | Y) + Var_Y E_X (X | Y)$
3.  $Cov(X, Y) = ECov(X, Y | Z) + Cov(E(X | Z), E(Y | Z))$

# Birth and death processes

- ▶  $W=I+B$ ,  $I$  and  $B$  independent

$$\sigma_d^2 = p(1-p) + \sigma_b^2$$

$$\lambda = p + b$$

$$r = \ln(p + b)$$

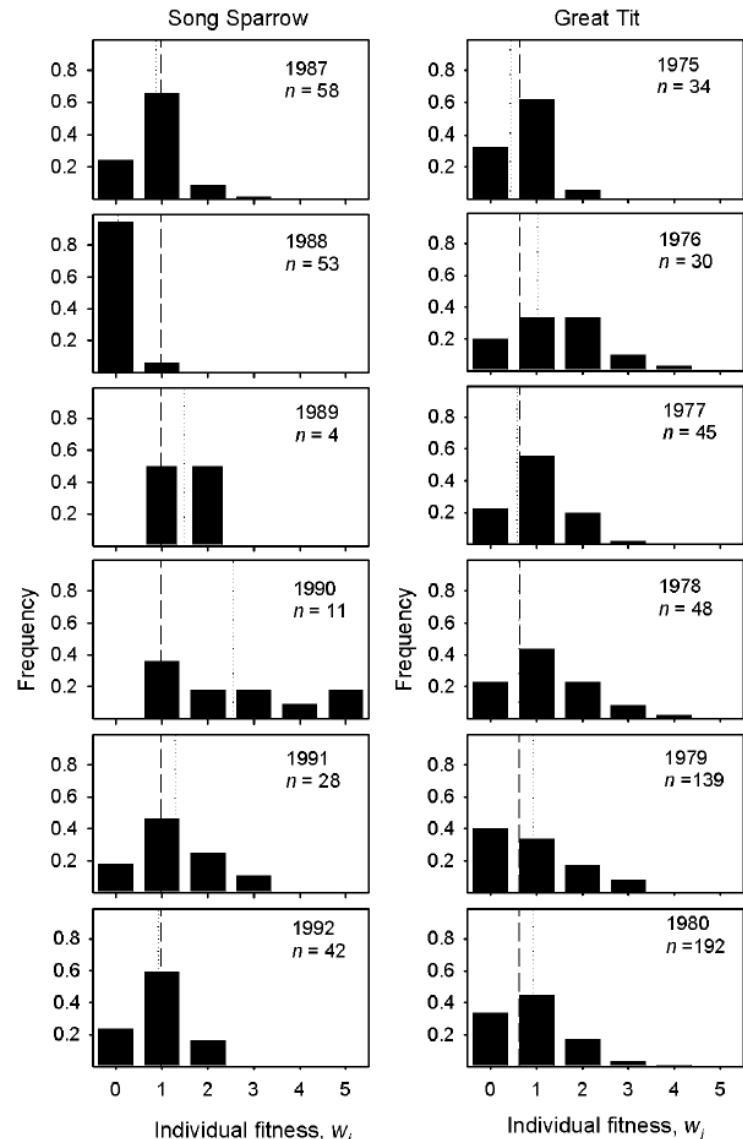


Figure 1.4: Annual variation in the distribution of contributions  $w$  to the next generation for two passerine species, the Song Sparrow *Melospiza melodia* on Mandarte island and the Great Tit in Wytham Wood. The dashed line indicates the mean values across all years and the dotted line the mean contribution a single year.

# Quick repetition

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- ▶ **Z** constant -> only demographic stochasticity

$$Var(\Delta N | N) = \sigma_d^2 N \quad Var(\Lambda) = \frac{\sigma_d^2}{N}$$

- ▶ Multiplicative model, large pop., **Z** stochastic

$$Var(\Delta N | N) = \sigma_e^2 N^2 \quad Var(\Lambda) = \sigma_e^2$$

# 1.7 Demographic and environmental stochasticity together

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If  $Z$  fluctuates between years -> individual contributions no longer independent

$$\rightarrow w_i = Ew + e + d_i$$

where

$$e = E(w | z) - E(w)$$

and

$$d_i = w_i - E(w | z)$$

# Fig 1.5

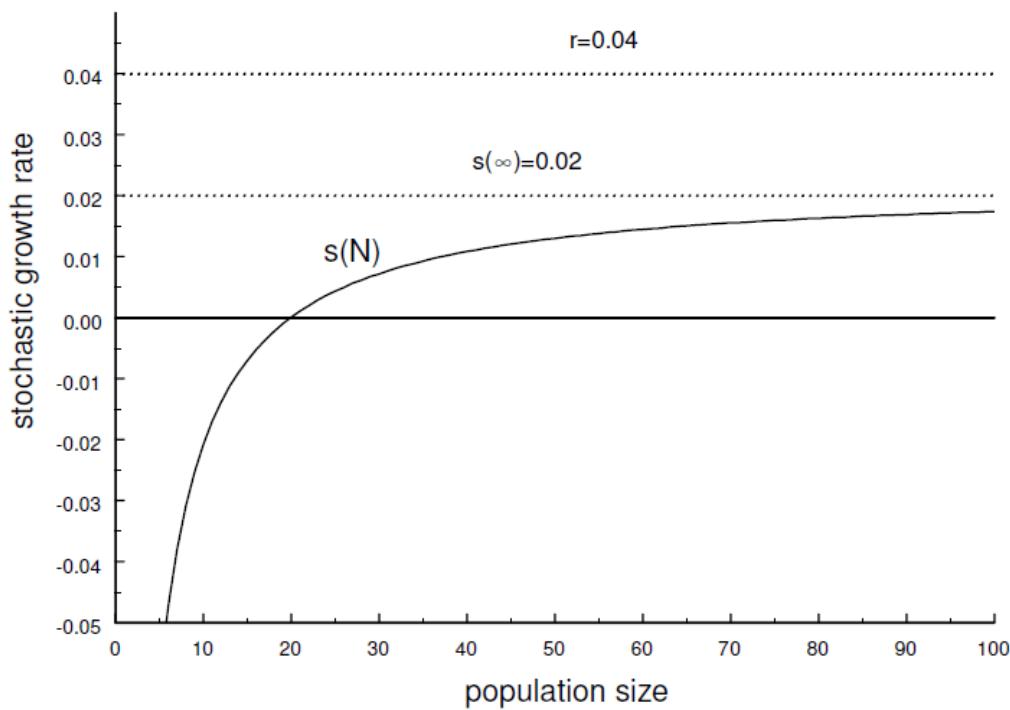


Figure 1.5: The stochastic growth rate as function of the population size. Parameter values are  $r = \ln Ew = \ln \lambda = 0.04$ ,  $\sigma^2 = 0.04$ , and  $\sigma_d^2 = 1$ , giving  $N^* \approx 20$ . The dotted lines show the value of  $r$  and  $s(\infty)$ .

# Fig 1.6

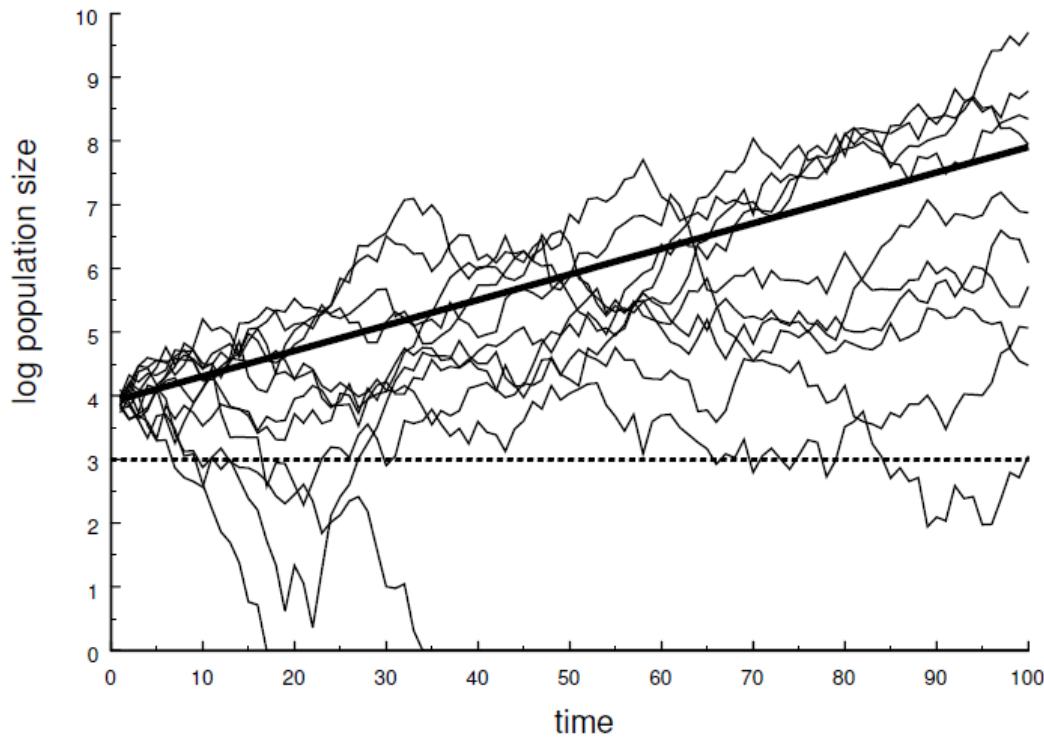


Figure 1.6: Simulations of the process described in Fig.1.5 with initial population size  $N_0 = 50$  giving  $\ln N_0 \approx 4$ .

