PBG 200A Notes

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1 Continuous Structure in Populations

Canonical example: size or weight

1.1 Integral Projection Models (IPMs)

Individual states take values in an interval [a,b]. Write N(t,x) as the density of size x individuals at time (or year) t. For a fixed t, we can look at N(t,x) vs. x. Then the density of individuals in $[x,x+\Delta x]$ is approximately $N(t,x)\Delta x$. Here is a classic

$$N(t+1,y) = \int_{a}^{b} s(x)[g(x,y) + p(x)f(x,y)]N(t,x)dx$$
 (1)

where

$$s(x) :=$$
the percentage of size x individuals which survive (2)

$$g(x,y) :=$$
the infinitesimal probability of growing from x to y (3)

$$p(x) :=$$
the probability of a size x individual reproducing (4)

$$f(x,y) :=$$
the density of offspring of size y produced by individuals of size x (5)

2 Matrix Models with Temporal Heterogeneity

$$N(t) = \begin{pmatrix} N_1(t) \\ N_2(t) \\ \vdots \\ N_k(t) \end{pmatrix}$$

$$(6)$$

$$N(t+1) = A(t+1)N(t) \implies N(t) = A(t)A(t-1)...A(1)N(0)$$
 (7)

What is r?

2.1 Central Limit Theorem ++

Assume A(i) are stationary, ergodic, and primitive for $i = 1, 2, \ldots$ Then $\exists r$ and σ ($\sigma > 0$) such that

$$\log(N_1(t) + \dots + N_k(t)) \approx \log(N_1(0) + \dots + N_k(0))rt + \sigma\sqrt{t}Z$$
(8)

where Z is the standard normal.

2.2 Small Variance Approximation for r

For i.i.d. A(t),

$$A(t) = \underbrace{\mathbb{E}[A(t)]}_{\overline{A}} + \underbrace{A(t) - \overline{A}}_{B(t)} \tag{9}$$

$$r \approx \log \lambda - \frac{1}{2\lambda^2} \sum_{i,j,r,s} S_{ij} S_{rs} \text{cov}[B_{ij}(1), B_{rs}(1)]$$

$$\tag{10}$$

where S are the sensitivities $(\partial \lambda / \partial \overline{a}_{ij})$.