

A Quick and Dirty Bayesian Leslie Matrix Model of Norway HarpEast Data

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Population Values

A_t = Number of Nonpups in year t

H_t^A = Harvest of Nonpups in year t

P_t = Number of Pups in year t

H_t^P = Harvest of Pups in year t

$N_t = A_t + P_t$

A Simple Two-age Leslie Matrix Model

Boveng, P.L., Ver Hoef, J.M., Withrow, D.E., and London, J.M. 2018. A Bayesian Analysis of Abundance, Trend and Population Viability for Harbor Seals in Iliamna Lake, Alaska. Risk Analysis 38(9): 1988-209 DOI:10.1111/risa.12988

$$A_t = \delta_{t-1}(A_{t-1} - H_{t-1}^A) + \kappa_{t-1}(P_{t-1} - H_{t-1}^P),$$
$$P_t = c_{t-1}\phi_{t-1}(A_{t-1} - H_{t-1}^A),$$

$$\mathbf{n}_t = \begin{pmatrix} P_t \\ A_t \end{pmatrix}, \mathbf{h}_t = \begin{pmatrix} H_t^A \\ H_t^P \end{pmatrix}, \mathbf{M}_t = \begin{pmatrix} 0 & c_t\phi_t \\ \kappa_t & \delta_t \end{pmatrix}$$

$$\mathbf{n}_t = (\mathbf{n}_{t-1} - \mathbf{h}_{t-1}) \mathbf{M}_{t-1}$$

$$c_t = \exp(-\rho N_t/1000)$$

c_t = Exponential decay in fecundity with increased N_t

Parameters

P_1 : Abundance of Pups in Year $t = 1$

A_1 : Abundance of Nonpups in Year $t = 1$

δ_t : Adult Mortality Rate in Year t

κ_t : Pup Mortality Rate, from Survey to Survey, in Year t

ϕ_t : Fecundity Rate for all Adults, to Survey Time, in Year t

ρ : Controls Exponential Decay in Fecundity as Population \uparrow

Priors

$$[P_1] = \pi(P_1) \sim N(300,000, 50,000^2)$$

$$[A_1] = \pi(A_1) \sim N(1,000,000, 100,000^2)$$

$$[\delta] = \pi(\text{logit}(\delta_t)) \sim N(2.2, 0.5^2) \forall t; \text{logit}^{-1}(2.2) = 0.90$$

$$[\kappa] = \pi(\text{logit}(\kappa_t)) \sim N(1.4, 0.5^2) \forall t; \text{logit}^{-1}(1.4) = 0.80$$

$$[\phi] = \pi(\text{logit}(\phi_t)) \sim N(-0.4, 0.5^2) \forall t; \text{logit}^{-1}(-0.4) = 0.40$$

$$[\rho] = \pi(\log(\rho)) \sim N(-4.0, 0.5^2)$$

Data

H_t^A and H_t^P for all years

p_t an estimate of pup abundance in year $t \in \mathcal{S}$

s_t a standard error of pup abundance in year $t \in \mathcal{S}$

\mathcal{S} is an index set of sampled years from T total years

\mathcal{T} is the index set of years $t = 1, \dots, T$

Likelihood

$$[p_t \mid P_t, s_t, \boldsymbol{\delta}, \boldsymbol{\kappa}, \boldsymbol{\phi}, \rho, P_1, A_1, \{\mathbf{h}_t; t \in \mathcal{T}\}] \sim \text{N}(P_t, s_t)$$

where P_t is obtained from

$$\mathbf{n}_t = (\mathbf{n}_{t-1} - \mathbf{h}_{t-1}) \mathbf{M}_{t-1}; t \in \mathcal{T}$$

where $\{\mathbf{n}_t; t \in \mathcal{T}\}$ depends on $\boldsymbol{\delta}, \boldsymbol{\kappa}, \boldsymbol{\phi}, \rho, P_1, A_1$

Posterior

$$[P_1, A_1, \delta, \kappa, \phi, \rho \mid \{(p_t, s_t); t \in \mathcal{S}\}, \{\mathbf{h}_t; t \in \mathcal{T}\}] \propto \prod_{\mathcal{S}} [p_t \mid P_t, s_t, \delta, \kappa, \phi, \rho, P_1, A_1, \{\mathbf{h}_t; t \in \mathcal{T}\}] [P_1] [A_1] [\delta] [\kappa] [\phi] [\rho]$$

Posterior of $\{\mathbf{n}_t; t \in \mathcal{T}\}$ is constructed from posteriors of $P_1, A_1, \delta, \kappa, \phi$ and ρ .

Use MCMC with Metropolis sampling to obtain sample from posterior distribution.

Custom code written in R, uses batch sampling for δ, κ, ϕ , with a burnin of 10,000 samples, followed by 100,000 where only each 100th sample was saved, yielding 1,000 samples from posterior.









