Patterns of firstpassage probabilities in population monitoring data



Confronting the theory with data (Holmes & Fagan 2002)

 141 chinook and 41 steelhead 30-70 year time series from ESUs in WA, OR, and CA



Corrupted Diffusion Approximation (CDA) really a 'Corrupted Random Walk Model'

 $log(N_{t+1}) = b log(N_t) + \mu + \varepsilon_{t,p}$ $log(y_{t+1}) = log(N_{t+1}) + \varepsilon_{t+1,np}$ $\varepsilon_{t,p} \sim Normal(0, \sigma_p^2)$ $\varepsilon_{t,np} \sim f(\beta, \sigma_a^2)$ b = 1

No density-dependence i.i.d. errors -> no auto-correlations Holmes (2001)

Theory makes a prediction about the distribution of mu_hat *random walk *N_t+1/N_t variance is related in a particular way to mu_hat variance



Similarly theory makes a prediction about the specific distribution of s_hat that we should observe



But...

- Problem: don't view the same population process over and over
- Actual data: many different processes with different underlying parameters (growth rates and variability)
- Solution: transform data to a standardized metric that has the same normalized statistical distribution for all processes

Standardized µ distribution



Standardized σ distribution

 $\left(\hat{\sigma}_{e}^{2}/\hat{\sigma}_{p}^{2}\right) \sim F(df_{slp}, df_{slp}) \times 1$

no trend in variance (variance is not a function of N)







First passage patterns: predicted versus observed

Let's look at some different data and a different study... This data has a lot less nonprocess error

117 Time series 20-50 yrs long72 are listed species



Distribution of process error estimates



IUCN Red List Criteria

- Criteria A2: "A reduction of at least xx%, projected or suspected to be met within the next xx years...."
- Criteria C1: "Population estimated to number less than xx and an estimated continuing decline of at least xx% within xx years...."
- Criteria E: "Quantitative analysis showing the probability of extinction in the wild is at least xx% within xx years..."

Cross-validation



Expected vs Observed Freq. of Hitting Particular Thresholds



REML

Proportional Declines













precision versus bias

- Once again 3 parameter model is successful at capturing number of extinctions observed in dataset = low bias....
- But maybe it's over-estimating low risks and under-estimating high risks (or visaversa) = precise??









These studies raise a larger question...

- Does there exist a general stochastic approximation to a very broad class of population trajectories?
 - Something like the stochastic Gompertz model in form
- How do we go about discovering that? And demonstrating that that form is indeed seen in real population trajectories?

Searching for general approximations for stochastic population trajectories...

- Building a theoretical foundation
 - Does there exist a general equivalent of (Tuljapurkar, Orzack, Heyde, Cohen)'s results but for population time series with density-dependence within a community web? Ives et al. 2004's result for derivation of Gompertz model from community models seems to be a start.
 - Can it be shown that the 'order' of this approximation is time dependent? Conjecture: short time (high order) → medium time (CDA) → long-time (DA)

Searching for general approximations for stochastic population trajectories...

- Building a statistical foundation
 - Need something akin to a sufficient statistic (parameter-free metrics) so that we can combine data from many different populations and study the distribution of those statistics.
 - Need to properly condition on observed data.
 - Cross-validation involves 'distribution of suff. statistic' times 'distribution of the estimates'. Theoretical pdfs of estimates are based on approximations, and on the CDA model. The CDA is merely an approximation for the real process. Do these approximations hold up with real data?
 - Can we delineate the set of equally plausible alternative explanations? Can we set up tests that can reject alternatives?



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