Improved state-space models for inference about spatial and temporal variation in abundance from count data:

Response to comments

Jeffrey A. Hostetler Richard B. Chandler

General Comments

In this submission, Hostetler and Chandler extend the so-called Dall and Madsen (DM) model for temporally and spatially-replicated animal point counts to include additional forms of process dynamics (e.g. exponential growth, density dependence) and to better cope with common features of ecological datasets (e.g. zero inflation, environmental noise, variation in observation error). The paper is well written, should be understandable by ecologists, and is reasonably motivated, the authors having situated their modeling approach within the context of state-space models for ecological time series (see Specific Comments for some musings on this topic). The authors also include requisite computing code to reconstruct the analysis within the R package unmarked, which is a nice touch and should make implementation of some of these methods easier for ecologists.

Thank you for your positive comments.

Despite these positives, I do have several reservations about the manuscript. First, the notion that one can estimate so many quantities with simple count data seems to me to be something like trying to squeeze blood out of a turnip. If distributional assumptions hold,

the authors have shown that a good deal of parameters (initial state distribution, population growth rates, immigration) can be estimated unbiasedly and have other good properties (coverage etc.). However, what if those distributional assumptions are violated?

In an ideal world, it seems to me that simulation studies should include two components: a 'proof of concept' type study to demonstrate parameter identifiability and to make sure the code isn't buggy (using the exact distributional forms to simulate data as are used in estimation), and (2) a robustness study to examine performance of the proposed model under suspected assumption violations. I think the ecological estimation literature is too often characterized by studies that employ (1) and not (2) (I am guilty of this myself). However, the need for a 'robustness' study seems particularly germane in this case, given the large number of assumptions to proceed with inference here. I suspect the authors must have similar feelings, given some of the statements in the discussion about the importance of including auxiliary data on detection probability, vital rates, etc.

We have added a robustness simulation section. We included three additional simulation models: DM with geometric-recruitment dynamics (underdispersed compared to Poisson), a Leslie matrix (underdispersed), and a Gompertz-logistic with two generations per year (overdispersed). We fit the first two sets of simulated data with the DM geometric-recruitment and our exponential version in unmarked, and the third with a Gompertz-logistic model with one generation per year (unmarked and JAGS) and two generations per year (JAGS). In addition to comparing model estimates, we also examined how well AIC did at selecting the more appropriate model in the first two cases.

I am also a bit troubled by the use of a Poisson process to encompass process error. This assumption in itself is a very strong one (see Specific Comments below), which seems to me

Table R1. Population dynamics options covered by improved state-space models.

Dispersion	Density-Independent	Density-Dependent
Underdispersed	DM	Not covered well yet
Poisson-distributed	Here	Here
Overdispersed	Here	Here

to be an additional argument for conducting a robustness study. What happens for instance if you use a Leslie-matrix type model on individual sites and then try to fit the Poisson process model? Do estimates come out near where they should? If not, is including an informative prior on detection probability (as if one had auxiliary data) sufficient to achieve good estimator properties?

In the Leslie matrix and geometric-recruitment simulations, estimates for Λ and p from the exponential model were biased, and in the Gompertz-logistic simulation, all estimates from the single generation models were biased. However, between the models presented in DM and here (including new overdispersion models), we cover almost all cases of density-dependence and dispersion (see Table R1). Furthermore, AIC does a good job of distinguishing between Poisson distributed and underdispersed data.

Specific Comments

1. Just pontificating here: I wonder if the use of the term "state-space" is becoming a little over-extended in the ecological literature. For instance, the generic state space model written in Eq. 1 certainly is a state space model in the sense it has Gaussian error and can be fitted using the Kalman filter. One could argue that the de Valpine and Hastings (2002)

model is analogous to a state space model, and they seem to have borrowed the terminology when naming their model. The models in the present paper share several features with state-space models: they are Markovian time series models with both process and observation error. On the other hand, I'm not sure it makes sense to talk of the Kery et al. (2009) model as a state space model at all. It doesn't have Gaussian error, can't be fit using a forward-backward type algorithm, and doesn't even have serial dependence. When does a "state space" model just become a hierarchical time series model??

We agree that some authors use the term "state-space" so broadly that its usefulness may be diminshed. However, we believe that the definition of state-space models that we provide in our introduction is sufficiently narrow to exclude many classes of hierarchical models.

Having said that, we agree that the Kéry et al. model doesn't fit within our definition because it isn't Markovian. This is now mentioned in the introduction (lines 106-108).

2. Is there a justification for using the Poisson process error model in Eqn 5 and throughout the remainder of the paper? This does not seem like an innocuous distributional choice, and it seems like real data could easily be over- or under-dispersed relative to the Poisson (e.g. with long-lived mammals or insects, respectively). Error associated with the binomial and Poisson distribution are drastically different unless the binomial success probability is really small and the index is large. I'm just not seeing any real justification other than modeling convenience.

When we have compared our models to the DM models with real data, the estimates from our models are more plausible and the AIC lower. That's our main justification - the data support the Poisson process error. In addition, we now present models that account for

overdispersion, and the original DM models account for some degree of underdispersion (see Table R1).

3. Simulation study. There needs to be more detail here. Was process error included in simulations? If so, was it induced with the same Poisson assumption as in estimation? I guess what I'm getting at here is that I'm dubious on whether the Poisson distribution is appropriate, so would be interested in seeing results where a different process error form is used in simulation that estimation. For instance, what does estimator performance look like when data are simulated with a binomial survival model and a Poisson immigration model as in the original DM model

Our "proof of concept" simulations did use Poisson process error. We have added one of the original DM models to our robustness simulation section.

Minor Comments/Typos

1. Equation 1. As written, lines 1 and 3 imply that $\eta_1 = 0$ (no sampling error in year 1). Given that there are other alternatives (e.g. the authors allude to using a stationarity assumption in year 1), would it make more sense to just specify an initial condition like N1 = x0 and state what the choices of x0 can be later (e.g. lines 95-97)?

Thank you for the suggestion! We have amended the equation and text (lines 66-68).

2. Lines 83-84. Instead of speaking of elusiveness, it may be better to distinguish between availability and detection probability given available as you do in the discussion- I think this way of looking at things is a little more general.

Thank you for the suggestion! We have amended the text (lines 81-83).

3. Line 91. Ignoring spatial variation (with respect to habitat covariates as well as spatial autocorrelation) has great potential for underestimating variances

This is a good point. We have amended the text (lines 90-92).

4. Line 132. Denoted what?

Denoted Ni,1. We have fixed this (line 132).

5. Line 150. I think these types of models would benefit from adding spatial autocorrelation via random effects at this stage

We agree that this would be a useful extension of these models, but believe it is beyond the scope of this manuscript.

6. Line 192 'to in'

We have corrected this (line 191).

7. Line 240. 'Three variations suggest themselves' - sounds like spontaneous generation!

We have taken ownership of these ideas (lines 241-242).

8. Line 294 simulations not simulation

We have corrected this (line 281).