Individual based movement estimation (Residuals and Petersen)

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Plan

- Reminder
- Residuals: conventional, archival, position and time
- Petersen: conditional on movement

Setup movement matrix

- Construct a giant matrix $M_{N\times N}^{\star}$, where N is the number of cells in the study area plus one cell for each of the ways a fish can die (natual or caught by fishing).
- Assuming we have one fishing fleet (one spatial field for its F), an advection field α , a diffusion field D, and one natural mortality field M, then we can setup the matrix as:

$$M_{i \to j}^{\star} = \begin{cases} D_i/\Delta^2 + 0.5\alpha_i^{(x)}/\Delta, & \text{if cell } j \text{ right of cell } i \\ D_i/\Delta^2 - 0.5\alpha_i^{(x)}/\Delta, & \text{if cell } j \text{ left of cell } i \\ D_i/\Delta^2 + 0.5\alpha_i^{(y)}/\Delta, & \text{if cell } j \text{ above cell } i \\ D_i/\Delta^2 - 0.5\alpha_i^{(y)}/\Delta, & \text{if cell } j \text{ below cell } i \\ F_i, & \text{if cell } j \text{ is 'caught'} \\ M_i, & \text{if cell } j \text{ is 'dead'} \\ -\sum_{j \neq i} M_{i,j}^{\star}, & \text{if } i = j \text{ (obviously calculated last in row)} \\ 0, & \text{otherwise} \end{cases}$$

 Δ is size of grid cell. Notice that e.g. F can depend on where the fish is.

• Solved with matrix exponential to find probability of each tag history.

Fields

• Fields are constructed from environmental inputs $I_1, I_2,...,I_m$ or effort E supplied on a grid, as:

$$h(i) = S_1^{(h)}(I_1(i)) + \dots + S_m^{(h)}(I_m(i)) \text{ used as } \alpha(i) = \nabla h(i)$$
$$\log D(i) = S_1^{(D)}(I_1(i)) + \dots + S_m^{(D)}(I_m(i))$$
$$F(i) = \lambda E(i)$$

- The S-functions are splines.
- Notice the discrete nature of these fields.

Archival tags

• Data:

$$o_0 = (x_0, y_0), o_1 = (x_1, y_1), \dots, o_n = (x_n, y_n)$$

First and last known, all others subject to observation uncertainty

• Model:

$$o_{t} \sim N(\psi_{t}, \Sigma_{o})$$

$$\psi_{t+\Delta t} \sim N(\psi_{t} + \alpha(\psi_{t}, t)\Delta t, 2D(\psi_{t}, t)\Delta t I_{2\times 2})$$

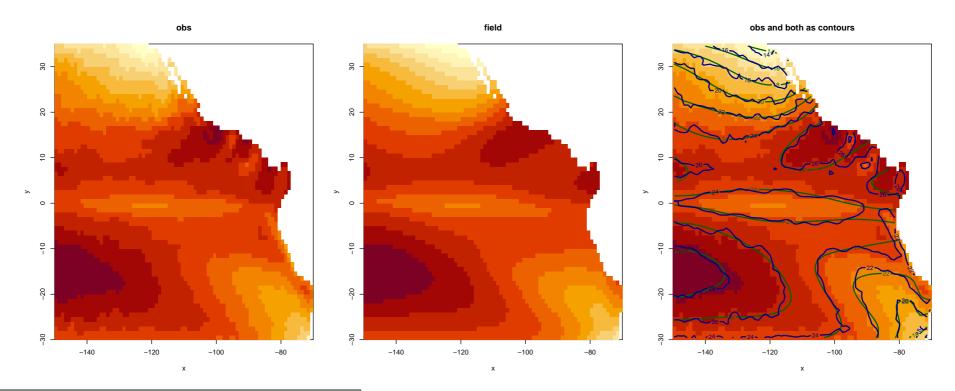
- Notice that next ψ depends on parameter values (via fields), so can switch to a new cell in grid ... $\mathring{\bullet}$
- Need a differentiable representation of each field

Differentiable representation of field

- Need to represent each field, so we evaluate anywhere $\alpha(x,y)$, differentiable in both x and y direction.
- Options: Splines, polynomials. neural network, (local regression), ...
- Splines and polynomials can be tricky to setup in 2D and can have erratic behavior near edges
- Tried first with a neural networks, but landed on local regression

Neural network representation

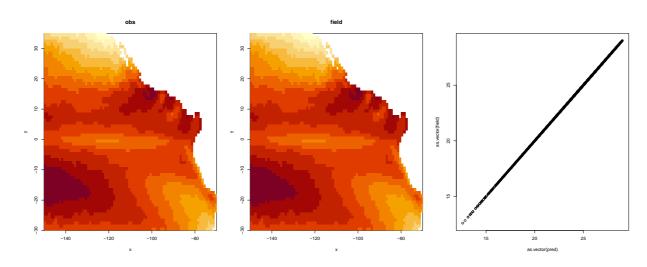
- A one layer network was fitted ("trained") externally
- 3 inputs x, y, and 1. 15 nodes in the hidden layer gives. 60 model parameters
- When the network has been trained we can represent the field by these 60 parameters
- Differentible and can easily be calculated: $f(x,y) = v_1\phi(w_{1,1}x + w_{1,2}y + w_{1,3}) + \cdots + v_{15}\phi(w_{15,1}x + w_{15,2}y + w_{15,3})$



^atwo layer also tested

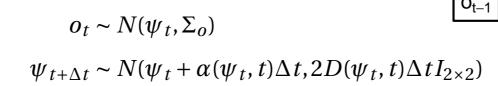
Local regression

- Use data as is
- Evaluate the field anywhere by weighted average of observations "nearby"
- This approach conflicted with a design principle in old TMB, but Kasper made it possible in new TMB!
- If we set the neaby-radius R = 1 (equal to distance between observations), then no smoothing is applied
- Nothing needs to be optimized
- The field is differentiable, but getting e.g. useful $\nabla h(x,y)$ needs some consideration
- Either apply smoothing, or define additional fields from Δx and Δy fields



Archival tags - now all OK

- Now we can do exactly what we want for archival tags
- Model:

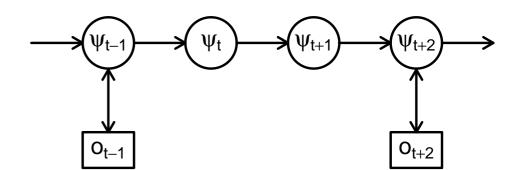


 O_{t+1}

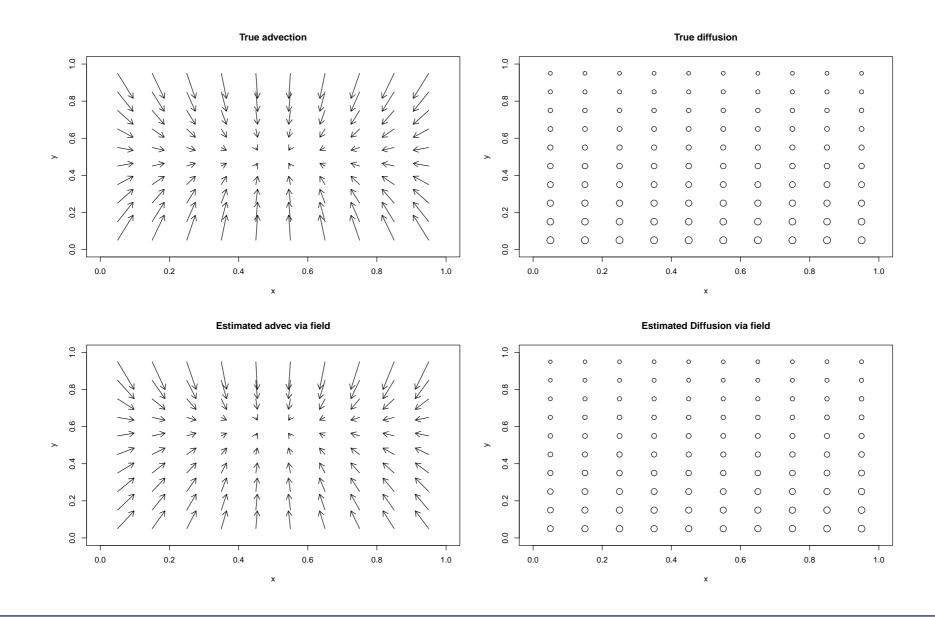
- Implemented via an old-style Kalman filter
- Fairly consistent with the matrix exponential approach for conventional tags, but notice that we did not need a grid for the archival tags...

Conventional as archival

- Can we consider a conventional tag as an archival where all the observations between start and end are missing
- Yes! In a state-space model we can have time-steps without corresponding observations
- However, we need to decide how many steps we we want to resolve its path into
- After that a Kalman filter without data update is used
- Advantage: No grid needed
- Advantage: Only local calculations needed for each tag
- Advantage: Fast! This was actually a bit of a surprise
- Advantage: Calculations can be done in parallel
- Advantage: The movement matrix can be extracted if we need it (and at different resolutions)



Simulation test (1000 conventional, 100 Archival)



Survival and catch

• A fish that is recaptured is surviving until that, so a history of:

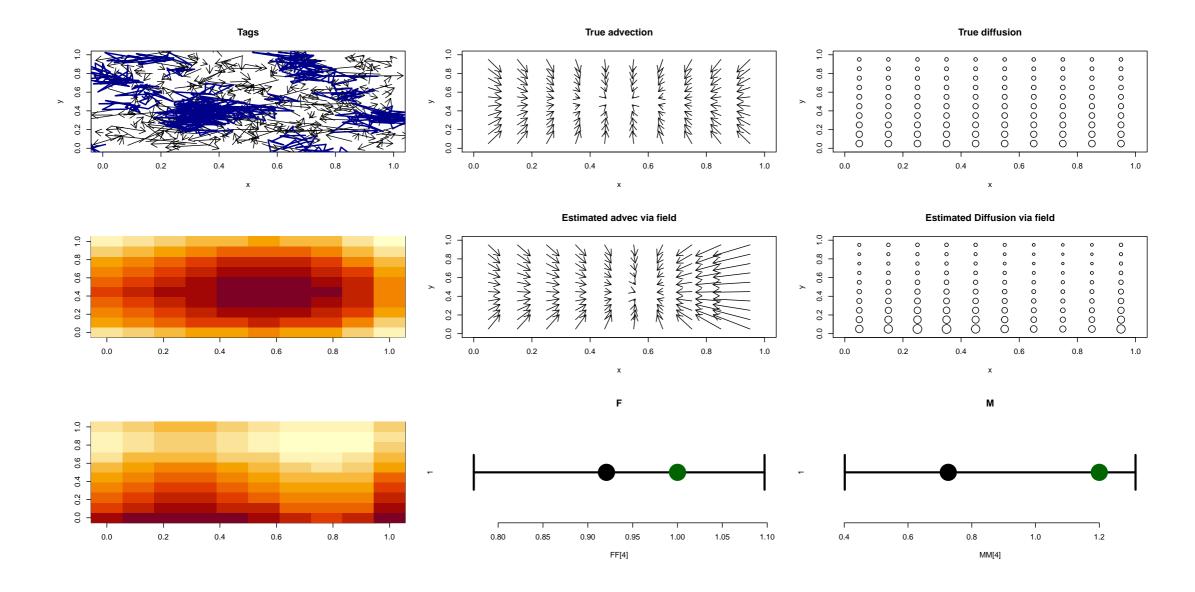
$$e^{-(M+F(x_1,y_1))\Delta t_1}e^{-(M+F(x_2,y_2))\Delta t_2}\cdots e^{-(M+F(x_{m-1},y_{m-1}))\Delta t_{m-1}}\frac{F(x_m,y_m)}{F(x_m,y_m)+M}(1-e^{-(M+F(x_m,y_m))\Delta t_m})$$

• A fish that is never recaptured is not caught first step, second step, ...

$$1 - \left(\frac{F(x_{1}, y_{1})}{F(x_{1}, y_{1}) + M} (1 - e^{-(M+F(x_{1}, y_{1}))\Delta t_{1}}) + e^{-(M+F(x_{1}, y_{1}))\Delta t_{1}} \frac{F(x_{2}, y_{2})}{F(x_{2}, y_{2}) + M} (1 - e^{-(M+F(x_{2}, y_{2}))\Delta t_{2}}) + e^{-(M+F(x_{1}, y_{1}))\Delta t_{1}} e^{-(M+F(x_{2}, y_{2}))\Delta t_{2}} \frac{F(x_{3}, y_{3})}{F(x_{3}, y_{3}) + M} (1 - e^{-(M+F(x_{3}, y_{3}))\Delta t_{3}}) + \cdots \text{ (or truncated)}$$

- Calculating these probabilities from the tracks is an approximation
- If focus was track's reconstructions, then these probabilities should be included in filtering

Simulation test (1000(283)) Conventional, 100(21) Archival)



Residual for position (conventional tag)

- Can only calculate these if the tag is recaptured
- According to the model the position follows a 2D independent normal, so

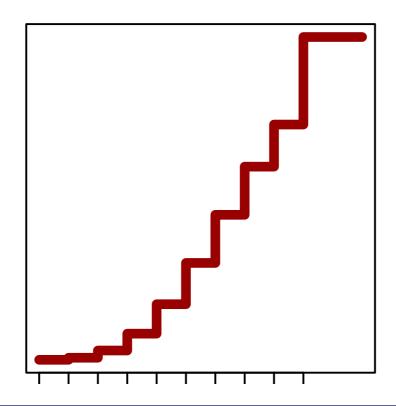
$$r_x = (x - \mu_x)/\sigma_x$$
 & $r_y = (y - \mu_y)/\sigma_y$

no problem.

Residual for recapture time (conventional tag)

- Model discrete in time, so recapture time is $t_1, t_2, ..., t_n$, or 'never'
- According to the model we can calculate probability of each
- Use randomized quantile residuals:

$$r_t = \Phi^{-1}(U), \quad U \sim \mathrm{unif}(P(T < t_{\mathrm{obs}}), P(T \le t_{\mathrm{obs}}))$$



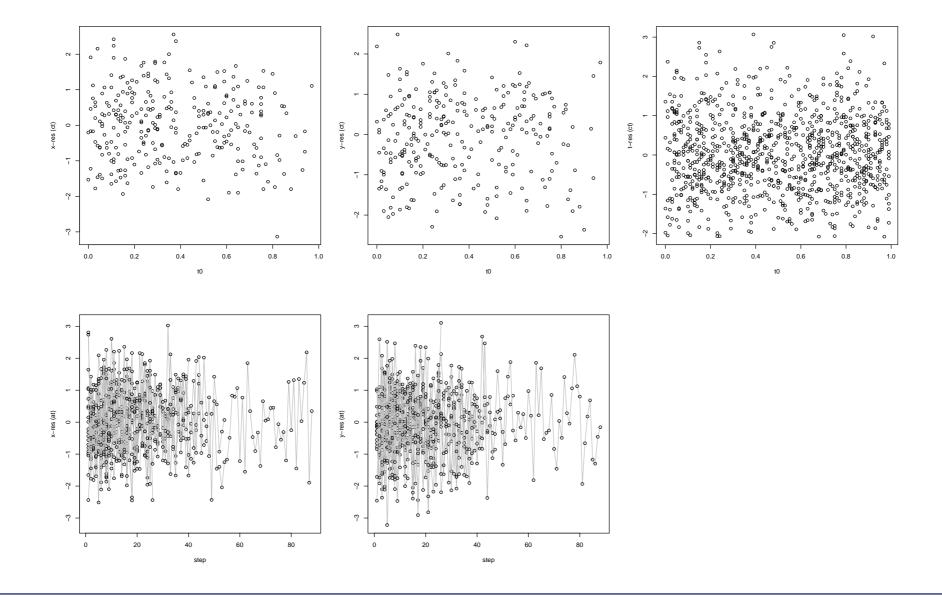
Residual for positions of archival tags

- Multiple residuals per tag
- Independent if we focus on prediction residuals

$$r_{x,i} = (x_i - \mu_{x_i|x_{i-}})/\sigma_{x_i|x_{i-}}$$
 & $r_{y,i} = (y_i - \mu_{y_i|y_{i-}})/\sigma_{y_i|y_{i-}}$

• A bit challenging how we plot them

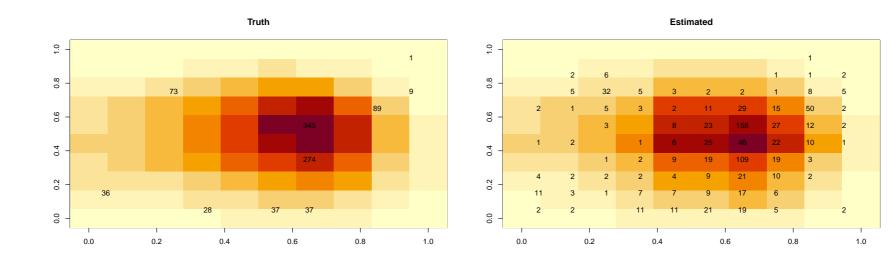
Residual example for the simulated case

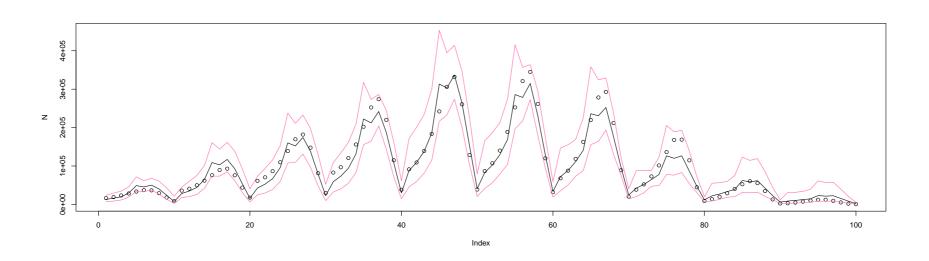


Exploration of Petersen type estimator

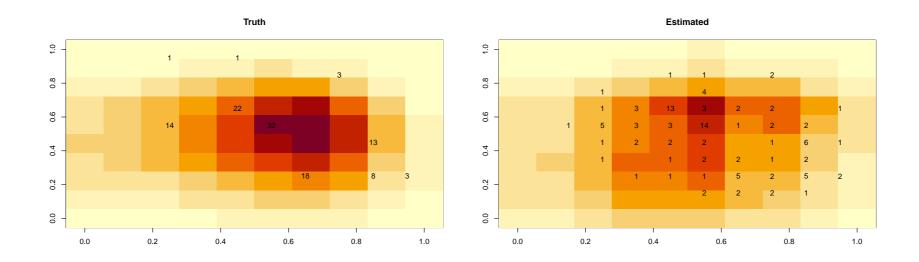
- Goal is to minimize reliance on the "effort" data
- We noticed that the movement pattern fairly constant with different effort patterns
- So can we treat movement as known, and use recovered tag-fraction to estimate abundance
- Condition on movement we can predict how many tags T_i will be in a given cell
- We know the catch taken C_i
- The expected number of tags observed will be $\frac{T_i}{N_i}C_i$
- So the observed number of tags may be $t_i \sim \text{Pois}\left(\frac{T_i}{N_i}C_i\right)$
- Coupled with a simple spatial model $N \sim \text{Gaussian}(\mu, \Sigma)$ (e.g. $AR(1) \times AR(1)$)
- Still very early, not all issues settled, but looks promising.

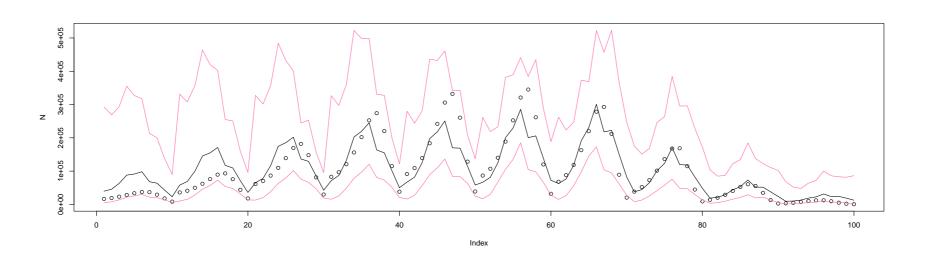
1000 tags released in 10 positions





100 tags released in 10 positions





100 tags released in 5 positions

