## Modelling dispersal, survival and trapping in SIT trials

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ModAH 2024 · Nantes, France

## Simulation of the process



- Thousands of insects (black dots) released from a single point
- They disperse independently from each other following a Brownian motion:
	- Changing direction continuously, with no preferential direction
	- **EXAMP** Same model of fluid diffusion at thermal equilibrium
- A few traps (orange squares) scattered over the area catch a fraction of the nearby individuals.
- As time progresses, individuals either eventually get caught or die (naturally, or from other causes such as predators)

## Target quantities

Diffusivity **Survival rate** 

The spread rate of individuals  $\begin{array}{|c|c|} \hline \end{array}$  The fraction of individuals still alive at a given time

- Diffusivity: the *rate of diffusion*. I.e. The rate at which the particules spread.
- Survival rate: the fraction of individuals still alive at a given time

## **Observations**

Counts of individuals caught during trap activation periods



- Each trap is set up and remains active during a few days
- At the end of the period, operators collect and count their catch

## Goal



- Develop a probabilistic model for the observations, given the target quantities (forward)
- Use observations to conduct statistical inference on the parameters (backward)

## We can only observe dispersal and survival

# indirectly

## through captures

Intuitively:

- Captures concentrated in the inner traps for a long time indicate small diffusivity and the converse
- Captures dropping quickly over time indicate low survival rate and the converse

## **Dispersal**

Brownian motion  $\rightarrow$  Diffusion equation:

 $\rho(\mathbf{x},t)$ : population density at position  $\mathbf{x}$  and time  $t.$ 

$$
\frac{\partial \rho}{\partial t} = D \nabla^2 \rho
$$

D: diffusivity coefficient

- Hypothesis: Brownian motion implies that the dispersion follows the diffussion equation, where
- $\nabla^2$ : laplacian operator (divergence of gradient)
- Where we find on of the parameters of interest: the diffusivity coefficient

## Dispersal: Closed-form solution

 $X_t$  ∣ *t*, *D* ∼  $\mathcal{N}$   $($  0, 2*tD***I**<sub>2</sub>)



• This simple dispersal model has a very simple analytical expression in terms of a bivariate Gaussian distribution centred at the release point and with dispersion proportional to *D* and *t*

## Effect of traps on density



Kernel density estimate of the empirical and theoretical distributions at day 15 with or without traps

- The capture of insects in traps cause a slight reduction in the density near the origin, with respect to the theoretical density (which ignores catch)
- In practice this is typically irrelevant. Only a problem if there is a significant share of insects caught after many days of capture to make it noticeable (15, in the example)
- However, this is a source of bias in the estimate of the diffusivity coefficient (wich is proportional to the variance of this density)

## Survival

Let  $T_0$  be the time of death of individuals.

Assuming a constant hazard rate<sup>1</sup>  $\lambda_0$ ,

$$
\begin{aligned} S_0(t) &= P(T_0 \leq t) \\ &= \exp\{-\lambda_0 t\} \end{aligned}
$$



- Hazard rate: instantaneous probability of death, given survival to that time.
- A simple (and common, although maybe not realistic) assumption is that the hazard rate is constant: if an individual survived until *t*, the probability of death in the next instant is independent of *t*
- Insects die, eventually, we don't know at which rate
- Furthremore, we can't **observe** their death directly
- When an individual gets caught, we know that they have survived until then

## Capture

Assumption: the hazard rate of capture decreases exponentially with the squared distance to the trap

$$
h_i(t\mid X_t) = \alpha e^{-\beta ||X_t - x_i||^2}
$$

Distance

- We need to model the capture process, even though its parameters are not our target quantities
- For trap *i* at  $x_i$ , conditional to position  $X_i$ .



#### Random variables associated to a released individual



• Use concepts and tools from time-to-event (survival) models with competing risks

## Specific hazard rates

$$
h_i(t)=\mathbb{E}_{X_t}\big[\alpha e^{-\beta||X_t-x_i||^2}\big],\quad \alpha,\beta>0
$$



- We don't know the position of every individual at time *t*, so we consider the expectation over its (known) probability density
- The calculation yields a closed-form expression for the specific hazard rates

### Hazard functions of the capture times for traps at increasing distances from the release location



(Dispersion and capture parameters as in the simulation)

## Capture times

#### Theoretical vs empirical distributions of capture times



Time (days)

(Traps with 10+ total catch)

## Mixed joint distribution

At any given time,

- �. Additive hazard rates
- �. Relative risks of death or capture are proportional to the specific hazard rates

$$
\mathbb{P}(C = i, T = t) = \mathbb{P}(C = i | T = t) f_T(t)
$$

$$
= h_i(t) S(t)
$$
where  $S(t) = \exp(-\int_0^t \sum_{i=0}^I h_i(u) du)$ 

Borrowing ideas from survival models with competing risks,

- At any given time, the hazard rates are additive
- Enabling the (numerical) computation of the overall survival (i.e. fraction of individuals that are still active at time *t*)
- We can integrate the distribution over the observation intervals for each trap to evaluate the likelihood!!
- $\dot{\boldsymbol{\omega}}$  Sadly, we lose the analytical expressions and need to evaluate the integrals numerically
- The exponential component multiplying the specific hazard rate is the overall survival function *S*(*t*)

## Distribution of end causes



Predicted vs. observed proportion of end events by cause (death / trap)

• We evaluated the expected and 95% quantile interval for the proportion of dead and caught individuals in each of the traps over all the simulation period and compared with the realised frequency (only traps with 1+ captures shown)

## Full likelihood



- We borrowed ideas from survival models with competing risks to combine the specific hazard rates and compute the joint distribution of end times and causes
- We integrate numerically over the observation periods to obtain the likelihood function
- The figure shows the likelihood function of the simulated scenario.
- The blue points indicate the maximum likelihood (which is almost flat over the diagonal)
- The true value is in the center of the central panel.

## SIT: sterile insect technique

Sequence of releases of sterile insects for pest control

• Released males mate with wild females producing sterile eggs

 $\vee$  Avoid pesticides (which contaminate the environment) [Introductory video presentation](https://www.youtube.com/watch?v=RVMAasW5dHc)

- Control disease transmission and reduce agricultural damage without insecticides
- Sterile males mate with wild females reducing offspring
- Need to quantify parameters such as number of individuals to release, temporal frequency, spatial extent, etc. to produce the desired effect of reduction or suppression of the wild population.
- These parameters depend, among other things on the **diffusivity** and the **survival** behaviour of the released sterile males in the context of the specific area.

## Issues

- Bias in dispersion (and diffusivity)
- Efficiency in numerical calculations (time consuming)
- Identifiability of capture parameters  $\alpha$  and  $\beta$ (reparameterise!)

## **Extensions**

- Non-isotropy
- Varying *attractiveness* of traps
- Integrate further sit parameters into the model
- Optimal designs

• Further parameters such as egg hatching rates, competitiveness…

## Conclusions

 $\blacktriangleright$  A formal probabilistic model for the dispersion, survival and capture of insects in sit experiments

**V** Enable inference with appropriate quantification of uncertainty in the target parameters

[WIP]

#### Thank you

- Press 's' to access speaker notes
	- Slides available at <https://astre.gitlab.cirad.fr/presentations/202408-modah-modelling-sit/>
	- An extended (23) version of the slides are at [https://astre.gitlab.cirad.fr/presentations/202408-modah](https://astre.gitlab.cirad.fr/presentations/202408-modah-modelling-sit/extended.html)[modelling-sit/extended.html](https://astre.gitlab.cirad.fr/presentations/202408-modah-modelling-sit/extended.html)

Slides made with quarto.

Basedon reveal.js and knitr.



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