

# Inferring moth emergence from abundance data: A novel mathematical approach using birth-death contingency tables

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HJ Andrews Experimental Forest  
Long Term Ecological Research

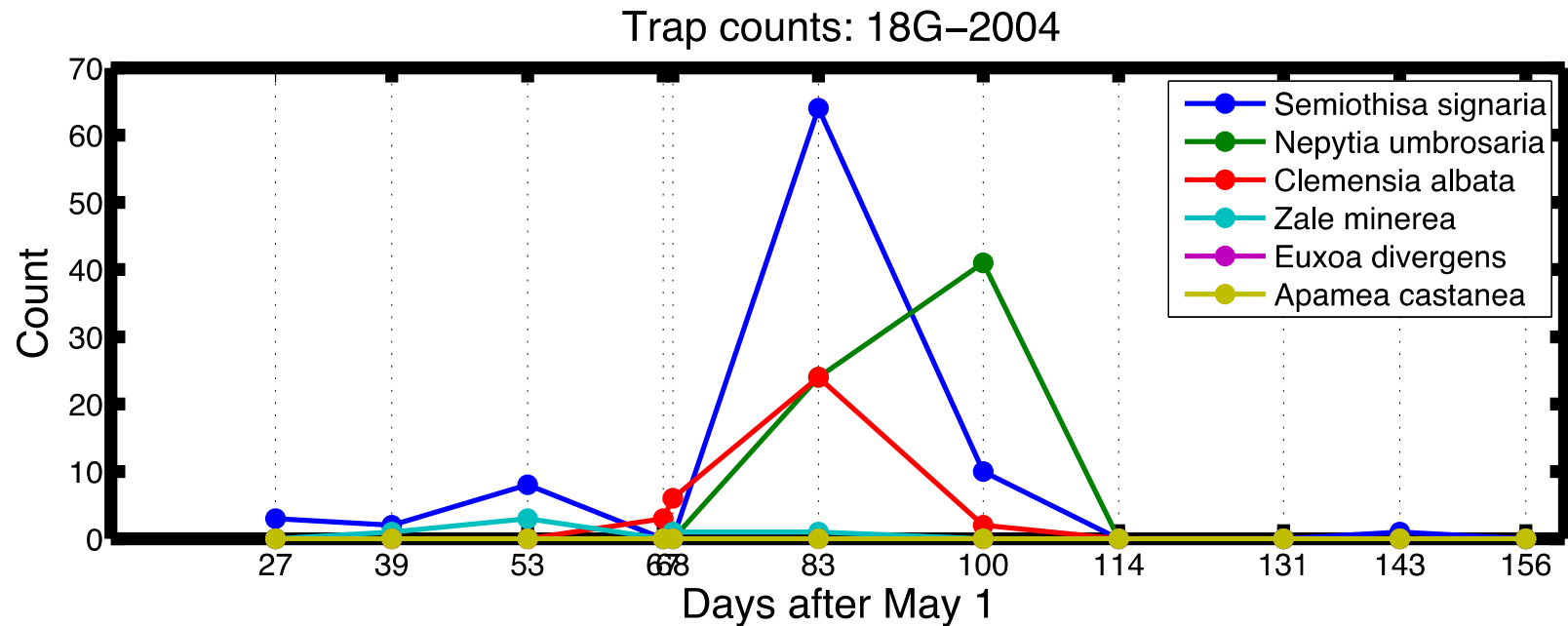


# A Common Problem in Ecology

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- ▶ ***What we have***: periodic observations of organism “activity”
  - ▶ Moth trap counts
  - ▶ Bird surveys
- ▶ ***What we want***: timing of life history events
  - ▶ When did adult moths emerge from cocoons?
  - ▶ When did migrating birds arrive?
- ▶ How to bridge the gap?

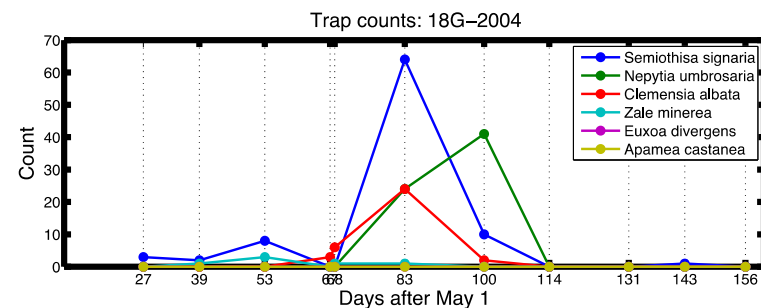
# Example: Moth Trap Counts



What was the flight period of *Nemytia umbrosaria* in 2004?

# Challenges

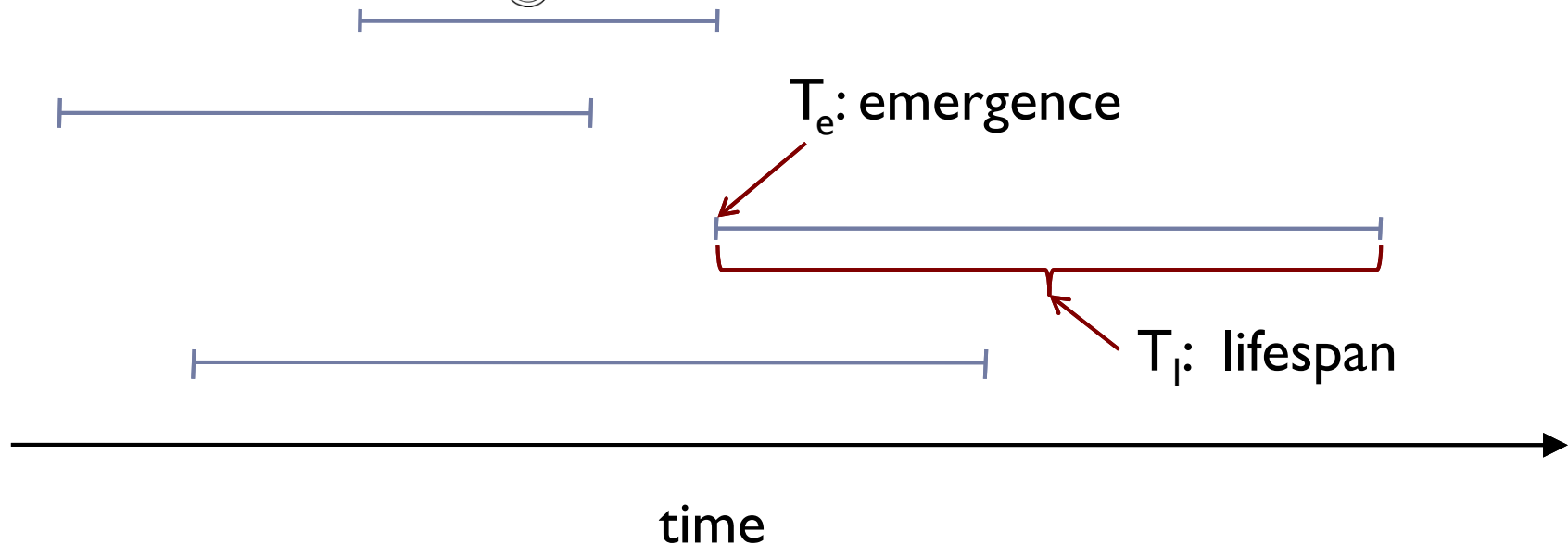
- ▶ We do not *directly* observe the events we are interested in
  - ▶ Moth emergence
  - ▶ Bird arrival
- ▶ Surveys are infrequent
  - ▶ May miss “peak” activity
- ▶ Naïve approaches don’t use all of the data
  - ▶ Date of first moth
  - ▶ Date of maximum abundance



# Probabilistic Modeling Approach

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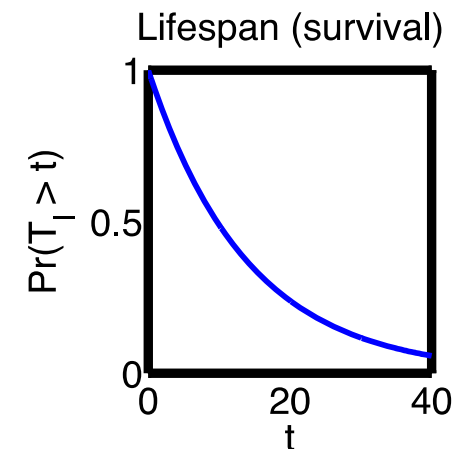
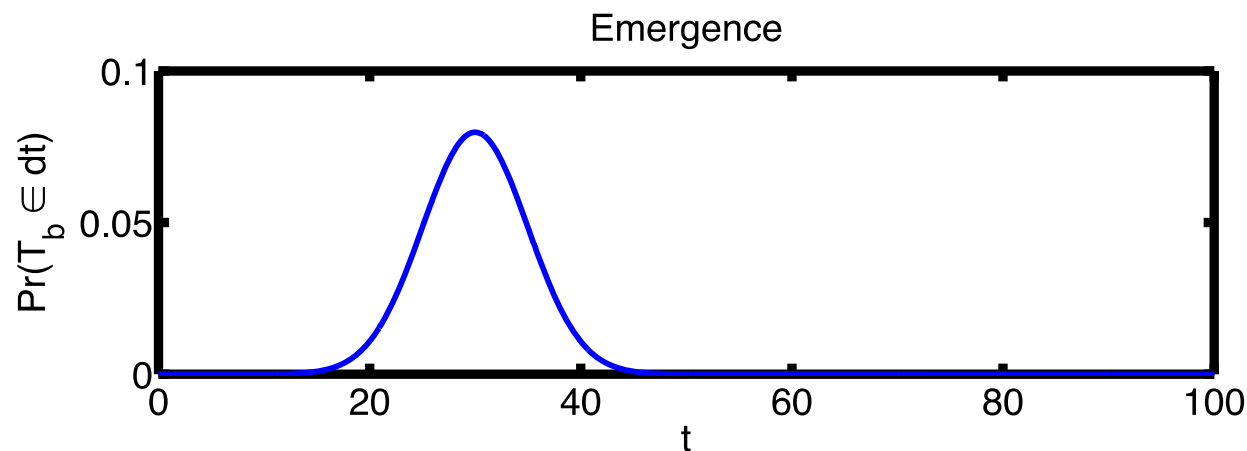
- ▶ Due to Zonneveld (1991), Manley (1974)
- ▶ Assume moths are independent and identical draws from a probability distribution



# Choose Lifetime Distributions

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- ▶ Emergence date is Normal( $\mu, \sigma^2$ )
- ▶ Lifespan is Exponential( $\lambda$ )

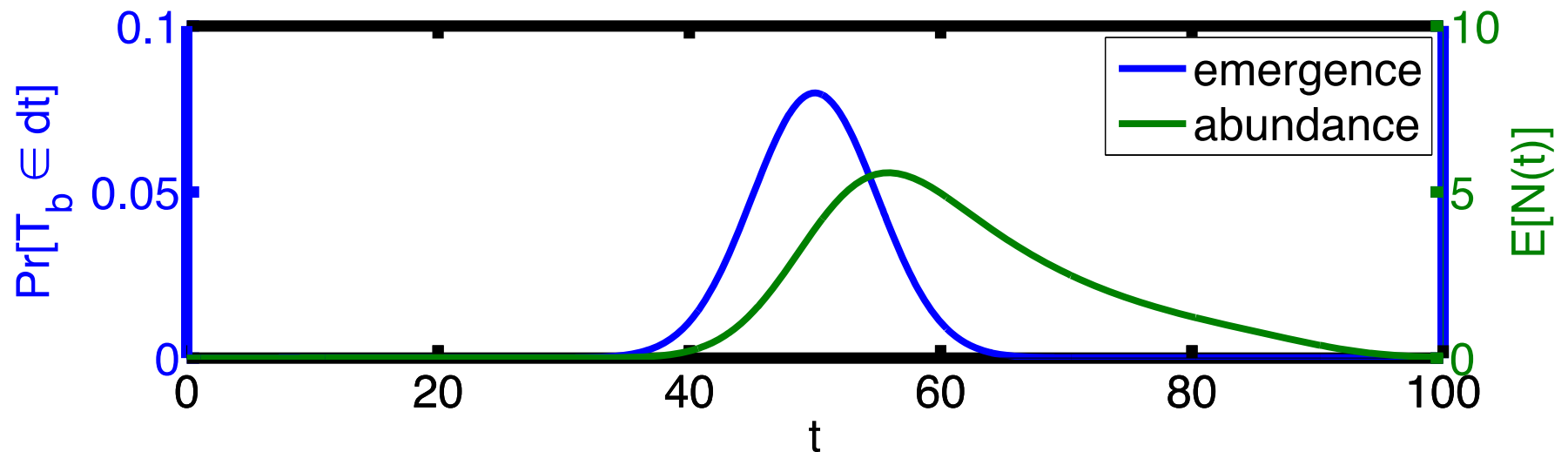


Note: *any* other parametric models can work  
(not restricted to Zonneveld's Logistic distribution)

# Abundance

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- ▶ Emergence and lifespan induce a model of abundance



- Goal: fit parameters  $(\mu, \sigma^2, \lambda)$

# Zonneveld (1991)

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- ▶ Compute expected number of moths  $X(t)$  flying at time  $t$  by solving

$$\frac{dx(t)}{dt} = N^* f(t; \mu, \sigma^2) - \lambda x(t)$$

- ▶ Assume Poisson distributed observation counts  
 $y(t) \sim \text{Poisson}(x(t))$

- ▶ Implemented in INCA ([www.urbanwildlands.org/INCA/](http://www.urbanwildlands.org/INCA/))
  - ▶ Bruggeman, Longcore & Zonneveld



# Differences between Transect Counts and Moth Trapping

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- ▶ **Traps are lethal**
  - ▶ requires change in the likelihood and survival functions
  - ▶ eliminates issues of double counting (either within a single night or across multiple nights)
- ▶ **Traps are very effective**
  - ▶ Less problem with detection rates
  - ▶ Nonetheless, we still include detection rate in our model

# Limitations of the Zonneveld Method

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- ▶ Differential equation approach makes it difficult to change the emergence distribution
- ▶ Likelihood requires an approximation (i.e., Poisson)

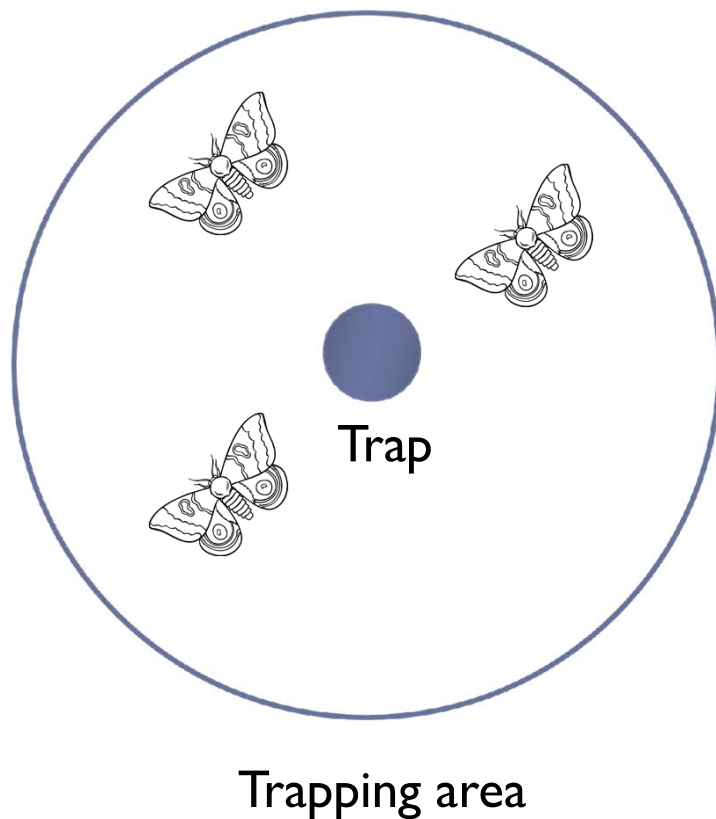
# Our Contributions

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- ▶ **General approach that can work with any emergence and mortality distributions**
  - ▶ Only requirement is the ability to evaluate a double integral over the convolution of the two distributions
    - ▶ Survival functions for emergence and mortality with weather covariates
    - ▶ Overlapping generations
  - ▶ Can model lethal and non-lethal trapping
- ▶ **Exact likelihood rather than an approximation**
  - ▶ Our likelihood converges to the Zonneveld likelihood in the limit where trapping probability goes to zero

# Trapping model

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- ▶ Closed trapping area
- ▶  $N$  moths (unknown) emerge during season
- ▶ Trap dates  $t_1, \dots, t_k$
- ▶ Trapping probability  $\alpha$  for each moth alive on trap date

# Data Likelihood

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- Can categorize each moth by its possible fates

Outcome	Probability	Trap Count
Trapped $t_1$	$q_1$	$f_1$
Trapped $t_2$	$q_2$	$f_2$
...	...	...
Trapped $t_k$	$q_k$	$f_k$
Not trapped	$r = 1 - (q_1 + \dots + q_k)$	$U = N - (f_1 + \dots + f_k)$

- Thus, likelihood of observed trap counts  $f_1, \dots, f_k$  is Multinomial:

$$(f_1, \dots, f_k, U) \sim \text{Multinomial}(N; q_1, \dots, q_k, r)$$

# Likelihood Computation and Model Fitting

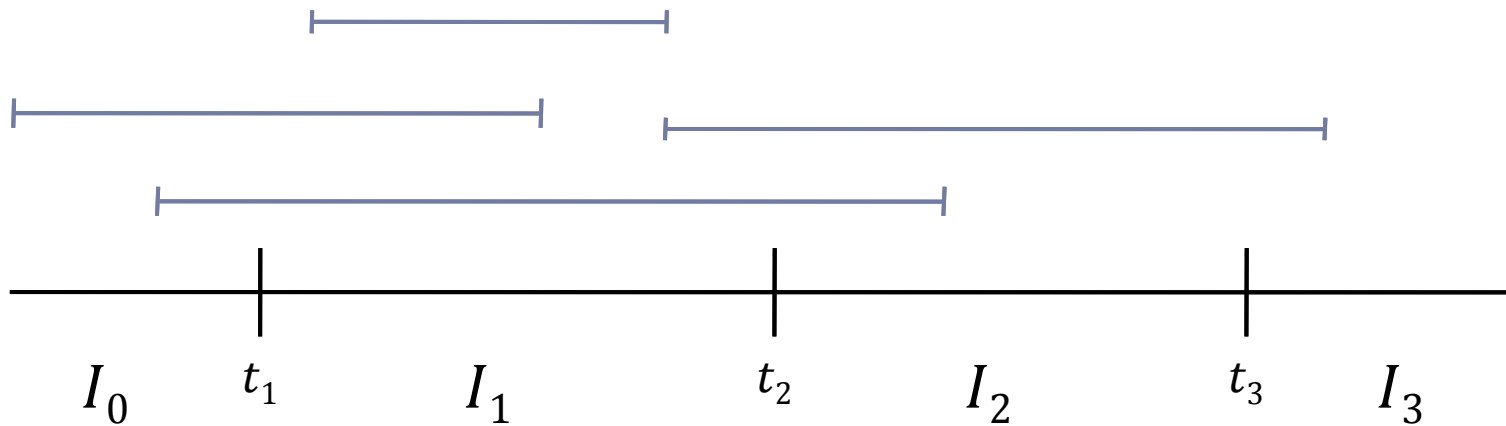
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- ▶ **To compute likelihood:**
  - ▶ Compute birth-death table  $P(i, j)$  from emergence and lifespan distributions (numerical integration)
  - ▶ Compute birth-trap table  $Q(i, j)$
  - ▶ Compute trapping probabilities  $q_1, \dots, q_k$
- ▶ Find  $(\mu, \sigma^2, \lambda)$  to maximize likelihood with numerical optimizer

# Intervals

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- ▶ Trapping times  $t_1, \dots, t_k$
- ▶ Intervals  $I_0, \dots, I_k$
- ▶ Trap counts  $f_1, \dots, f_k$

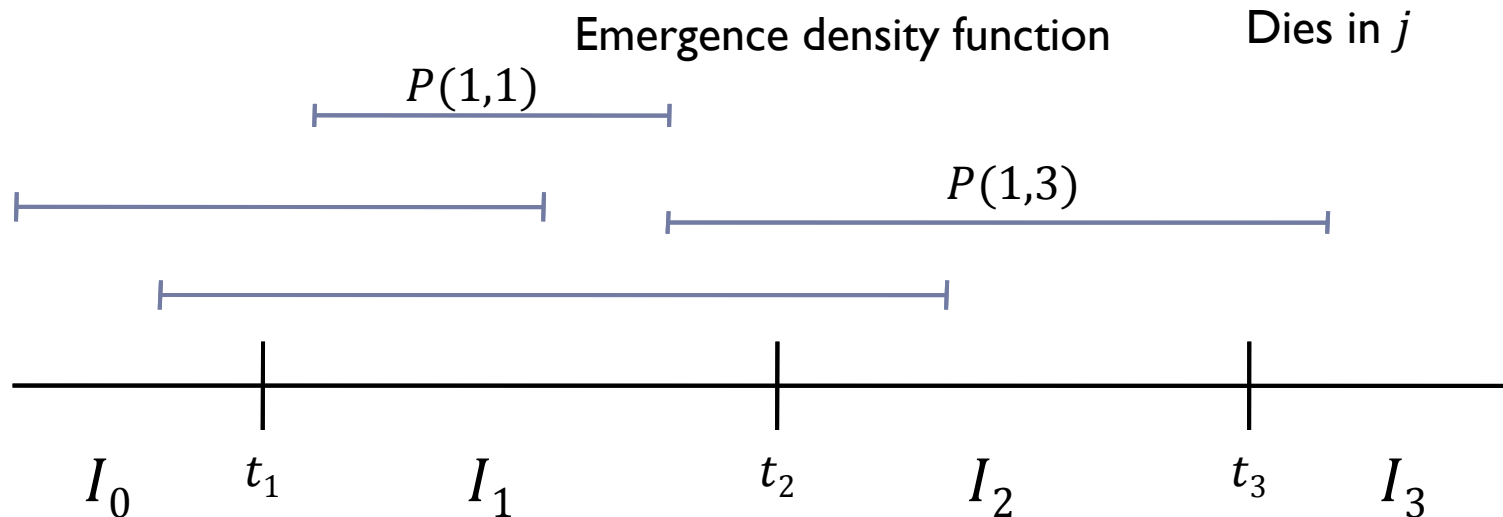


# Birth-Death Table

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- ▶ Let  $P(i, j)$  be probability a moth is born in  $I_i$  and dies in  $I_j$

$$P(i, j) = \int_{t_i}^{t_{i+1}} f(s; \mu, \sigma^2) \Pr[t_j \leq s + T_l \leq t_{j+1}] ds$$



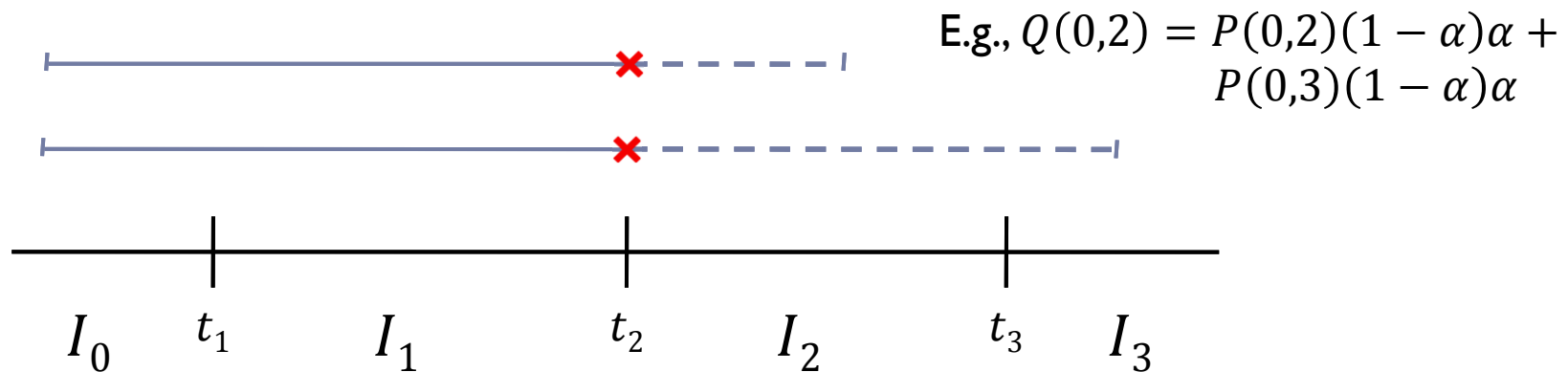


# Birth-Trap Table

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- ▶ Let  $Q(i, j)$  be probability a moth is born in  $I_i$  and trapped at  $t_j$

$$Q(i, j) = \sum_{k \geq j} P(i, k) (1 - \alpha)^{j-i-1} \alpha$$



# Trap Probabilities

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- ▶ Overall probability of being trapped at  $t_j$

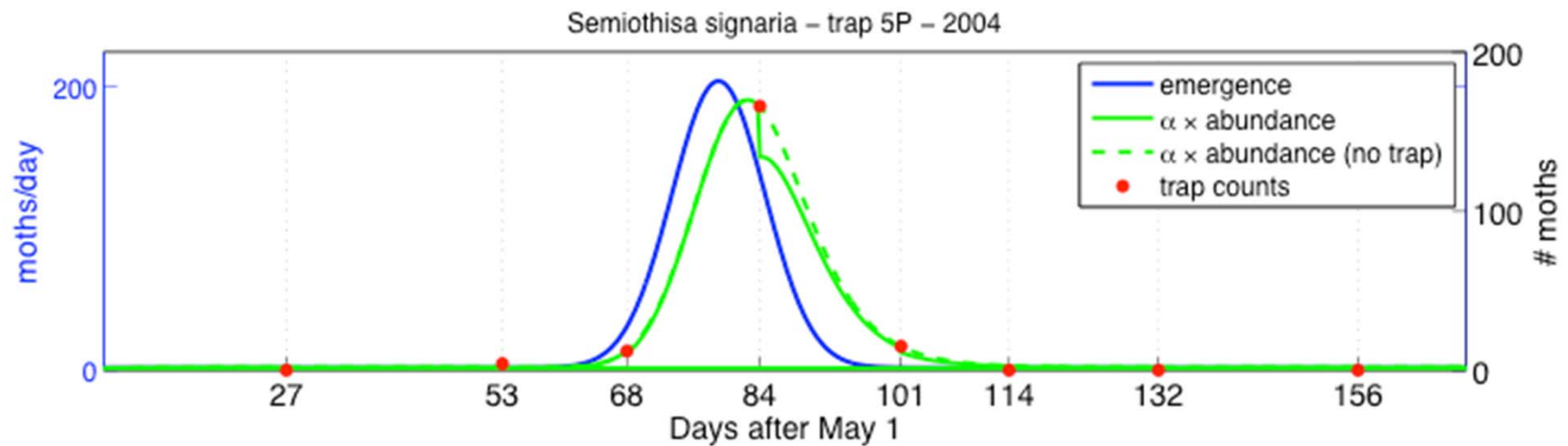
$$q_j = \sum_{i < j} Q(i, j)$$

- ▶ Probability not trapped at all

$$r = 1 - \sum_j q_j$$

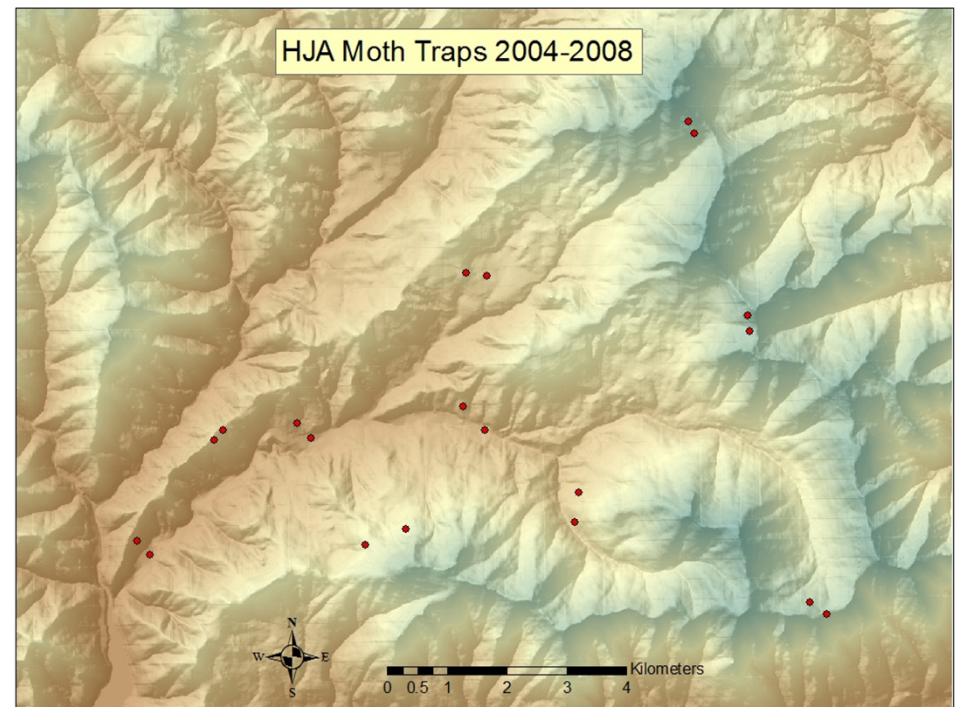
# Results

## ▶ Example of fitted model



# Data

- Moth trapping at HJA, 2004-2008
- 20 paired sites
- 10 trapping dates/year
- Approx. 2 week intervals
- > 500 species

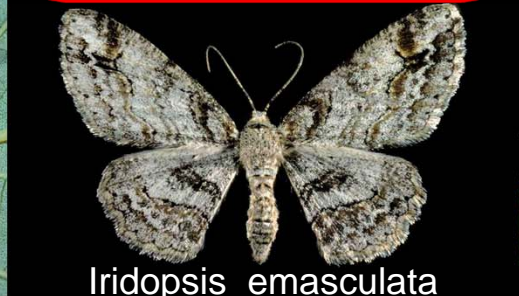




Clemensia\_albata



Nepytia\_umbrosaria



Iridopsis\_emasculata



Panthia\_portlandia



Melanolopia\_imitata



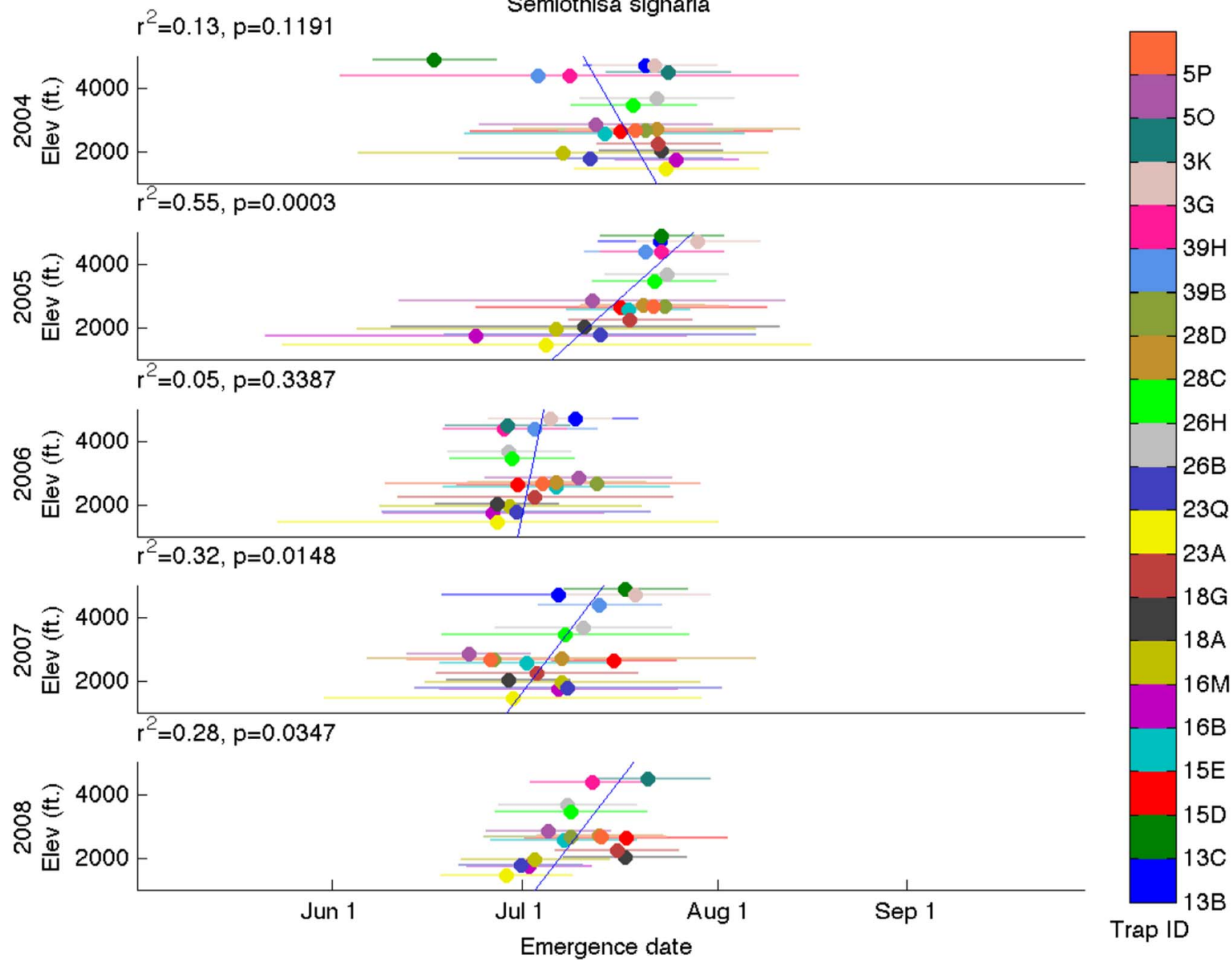
Pero\_mizon



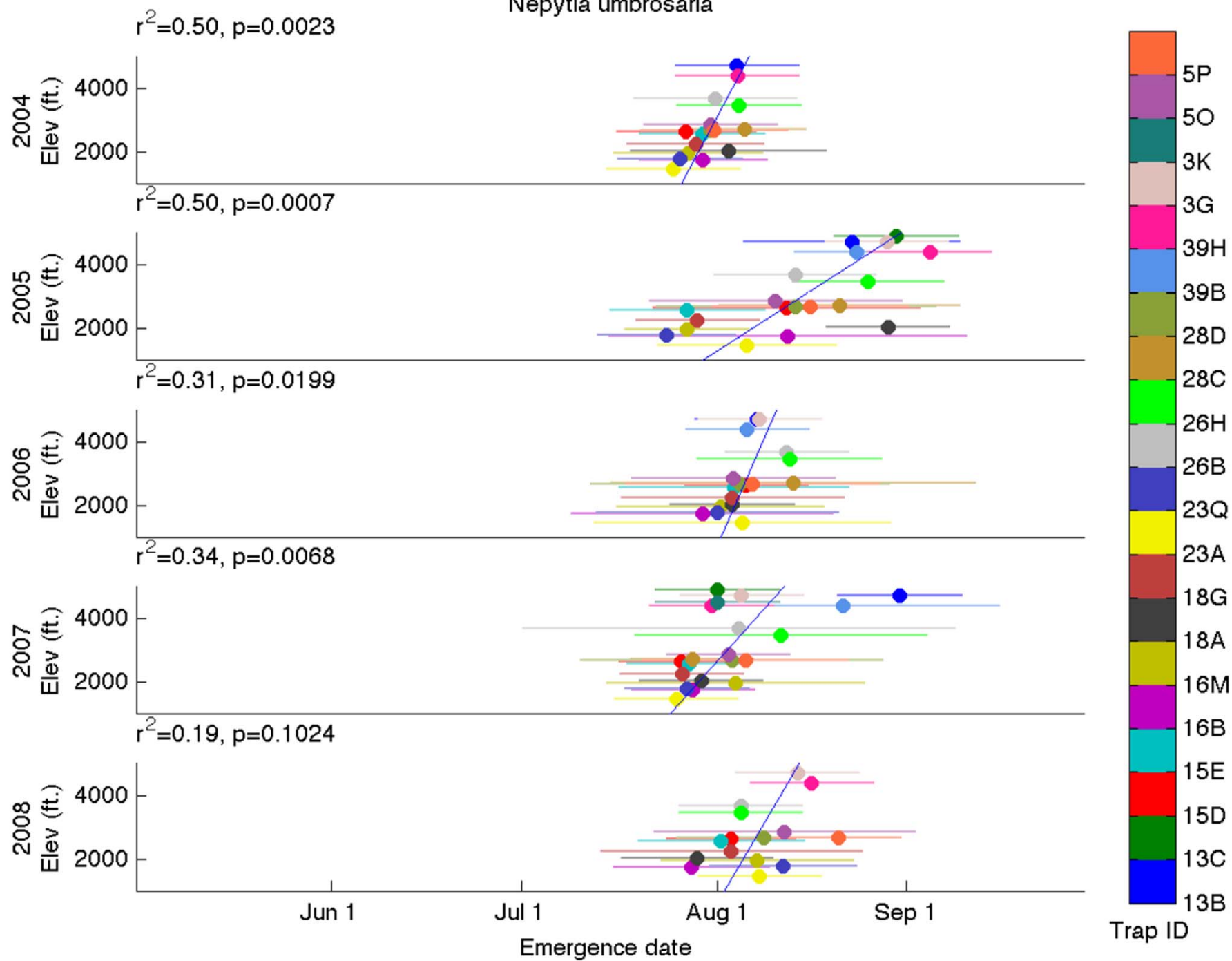
Semiothisa\_signaria

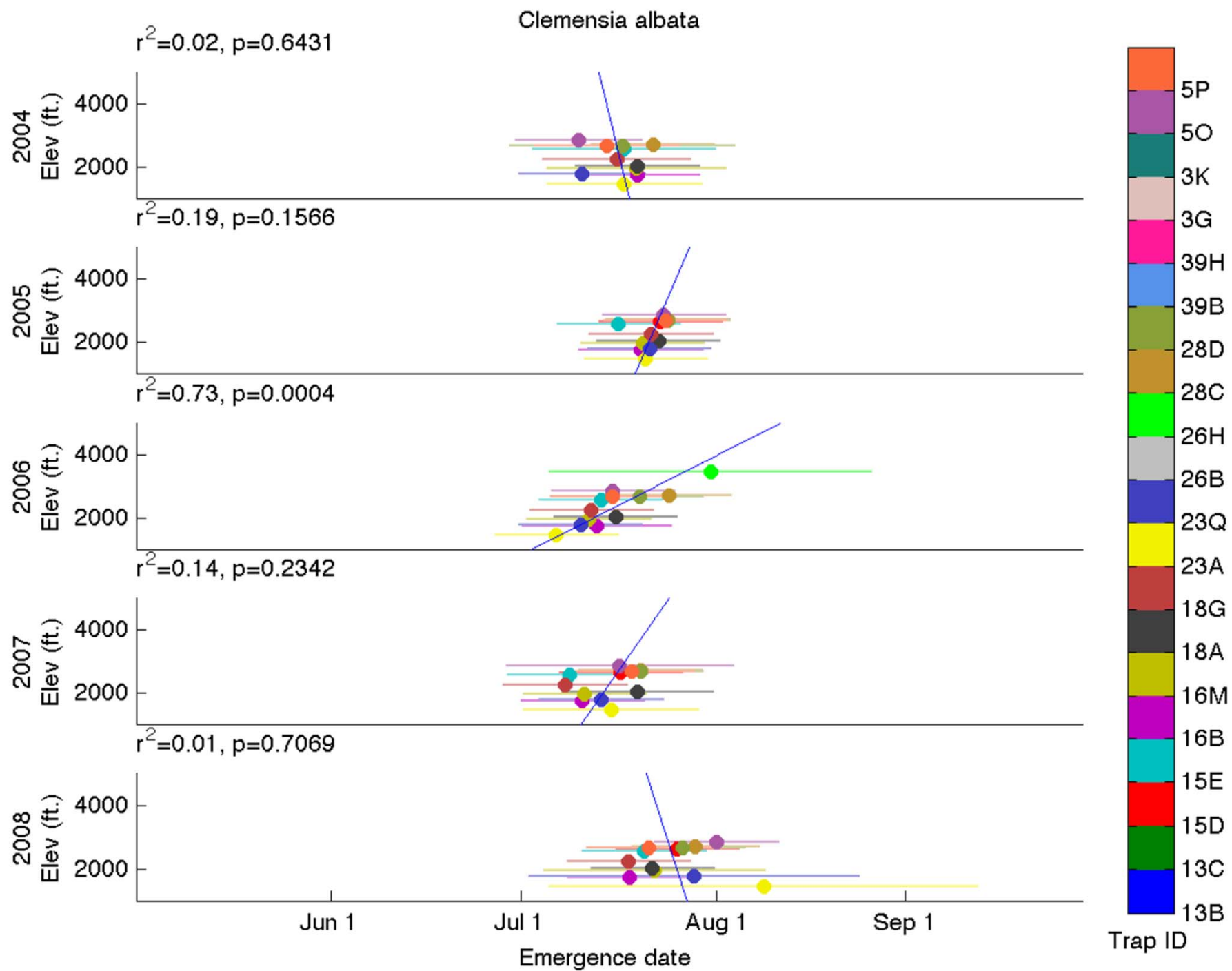
Seven of the eight most common moths in the HJ Andrews (photos by Jeff Miller)

*Semiothisa signaria*



*Nepytia umbrosaria*







# Summary

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- ▶ **Generalization of the Zonneveld model**
  - ▶ Works with any parametric birth (arrival) and death (departure) processes
  - ▶ Provides exact likelihood
  
- ▶ At HJA, often significant correlation between elevation and moth emergence, but also unexplained variability in this pattern
  
- ▶ **Future work:**
  - ▶ Incorporate environmental covariates into model (e.g., degree days)
  - ▶ Explore model limitations via simulation study
  - ▶ Obtain confidence intervals on parameters
  - ▶ Extend to other phenology questions: Bird Migration

Thank-you to Nick Haddad for introducing  
us to the Zonneveld Model

NSF LTER; NSF Ecoinformatics IGERT (Jones); NSF  
Computational Sustainability Expedition (Dietterich); NSF  
Ecoinformatics Summer Institute; NSF BIO Postdoc  
(Sheldon)