



The generalised abundance index approach

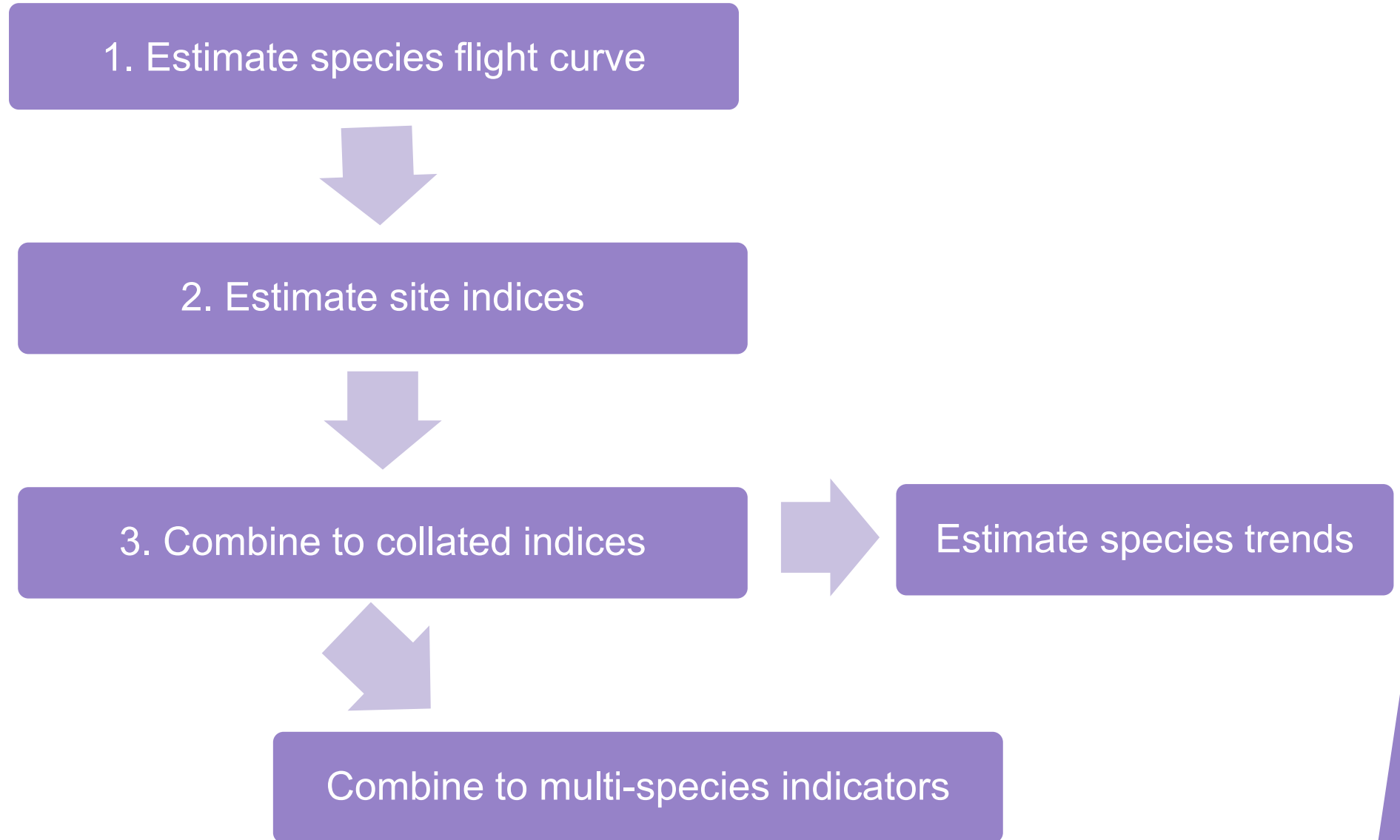


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Overview of modelling steps





Biological Conservation

Volume 12, Issue 2, September 1977, Pages 115-134



A method for assessing changes in the abundance of butterflies

[https://doi.org/10.1016/0006-3207\(77\)90065-9](https://doi.org/10.1016/0006-3207(77)90065-9)

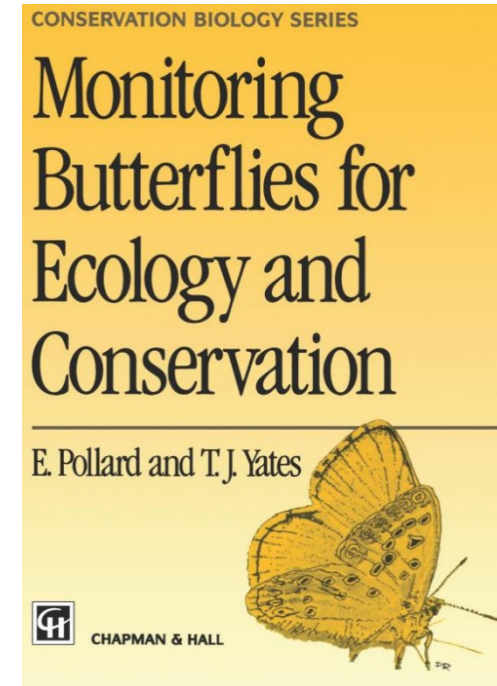
E. Pollard



Calculation of collated indices of abundance of butterflies based on monitored sites

D. MOSS, E. POLLARD

First published: February 1993 | <https://doi.org/10.1111/j.1365-2311.1993.tb01083.x>



Original Articles

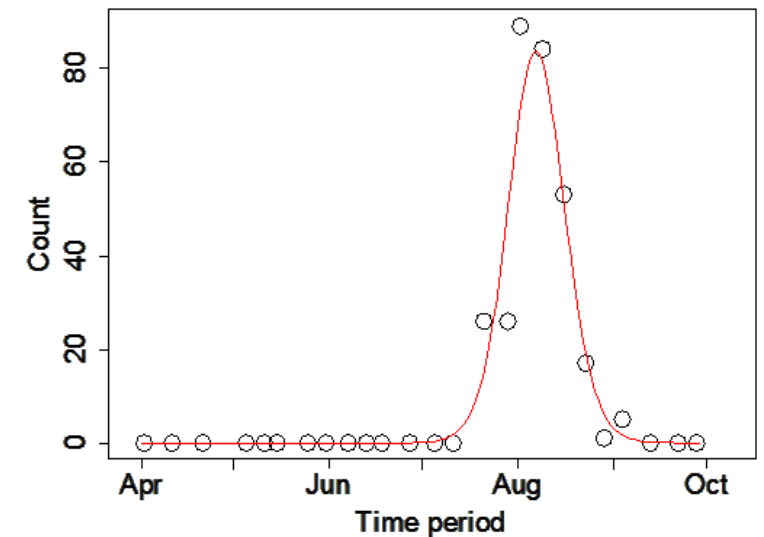
Application of generalized additive models to butterfly transect count data

Peter Rothery & David B. Roy

Pages 897-909 | Published online: 02 Aug 2010

<https://doi.org/10.1080/02664760120074979>

- A generalized additive model (GAM) is fitted to each site and year individually
- Excludes data where peak flight period is unrecorded or more than 30% data requires estimation
- For the UK, nearly 40% of monitored 10km grid squares were excluded



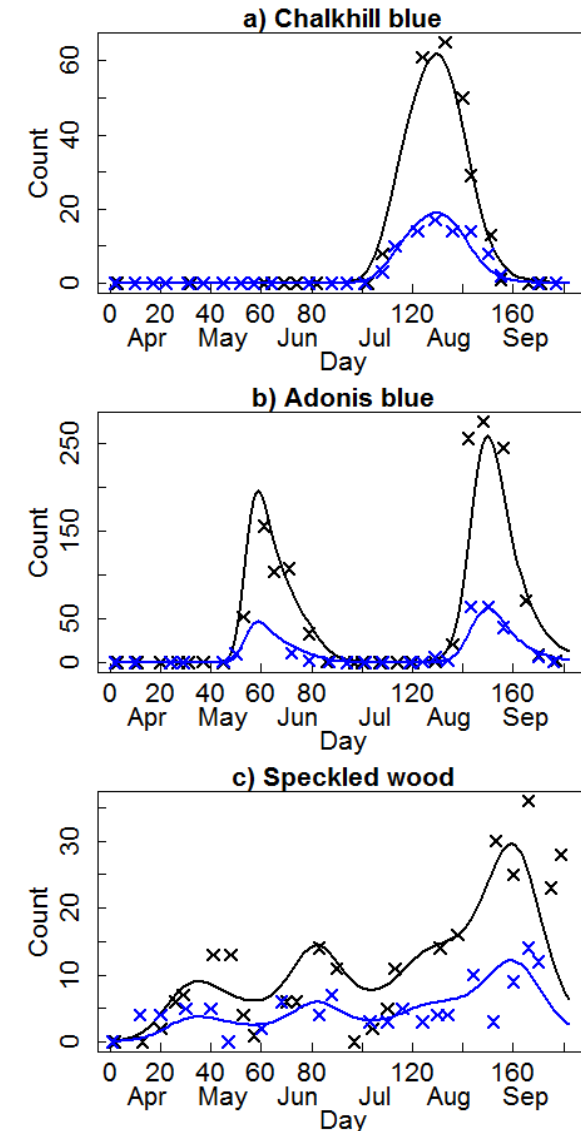
Standard Paper | [Free Access](#)

Indexing butterfly abundance whilst accounting for missing counts and variability in seasonal pattern

Emily B. Dennis , Stephen N. Freeman, Tom Brereton, David B. Roy

First published: 26 March 2013 | <https://doi.org/10.1111/2041-210X.12053> (open)

- A GAM is used to estimate a **common** flight period **across sites** for each year.
- More robust indices and trends than single-site GAM or linear interpolation
- Greater use of data
- Disadvantage – can be slow when there are lots of data



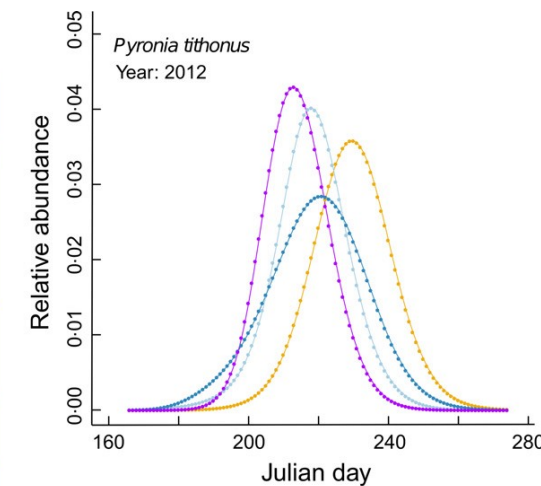
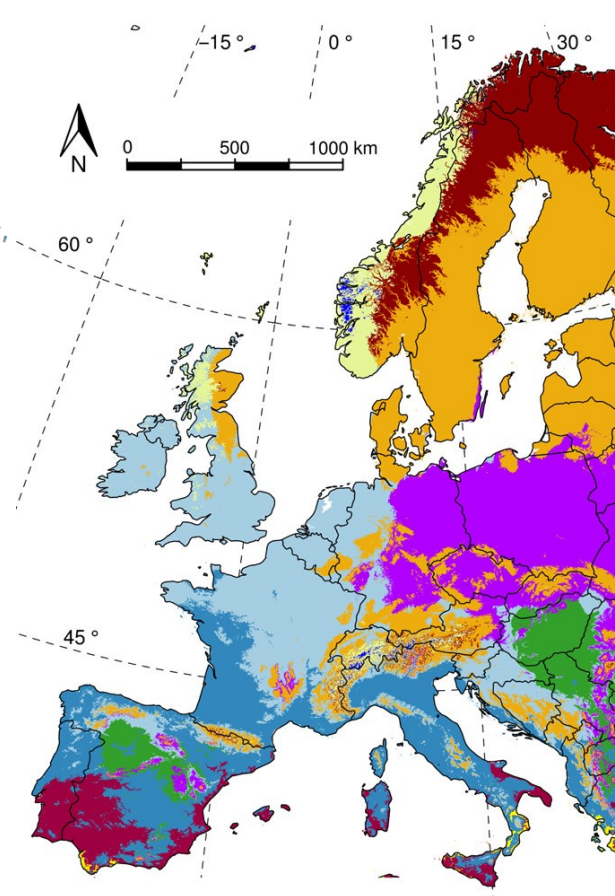
Standard Paper | [Open Access](#) | [CC](#) [i](#)

A regionally informed abundance index for supporting integrative analyses across butterfly monitoring schemes

Reto Schmucki [✉](#), Guy Pe'er, David B. Roy, Constanti Stefanescu, Chris A.M. Van Swaay, Tom H. Oliver, Mikko Kuussaari, Arco J. Van Strien, Leslie Ries, Josef Settele, Martin Musche ... [See all authors](#) [v](#)

First published: 28 October 2015 | <https://doi.org/10.1111/1365-2664.12561> | Citations: 14

<https://doi.org/10.1111/1365-2664.12561> (open)



- Extremely cold & wet
- Extremely cold & mesic
- Cold & wet
- Cold & mesic
- Cold temperate & moist
- Cold temperate & dry
- Cool temperate & xeric
- Warm temperate & mesic
- Warm temperate & xeric
- Hot & dry

A Generalized Abundance Index for Seasonal Invertebrates

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SUMMARY. At a time of climate change and major loss of biodiversity, it is important to have efficient tools for monitoring populations. In this context, animal abundance indices play an important rôle. In producing indices for invertebrates, it is important to account for variation in counts within seasons. Two new methods for describing seasonal variation in invertebrate counts have recently been proposed; one is nonparametric, using generalized additive models, and the other is parametric, based on stopover models. We present a novel generalized abundance index which encompasses both parametric and nonparametric approaches. It is extremely efficient to compute this index due to the use of concentrated likelihood techniques. This has particular relevance for the analysis of data from long-term extensive monitoring schemes with records for many species and sites, for which existing modeling techniques can be prohibitively time consuming. Performance of the index is demonstrated by several applications to UK Butterfly Monitoring Scheme data. We demonstrate the potential for new insights into both phenology and spatial variation in seasonal patterns from parametric modeling and the incorporation of covariate dependence, which is relevant for both monitoring and conservation. Associated R code is available on the journal website.

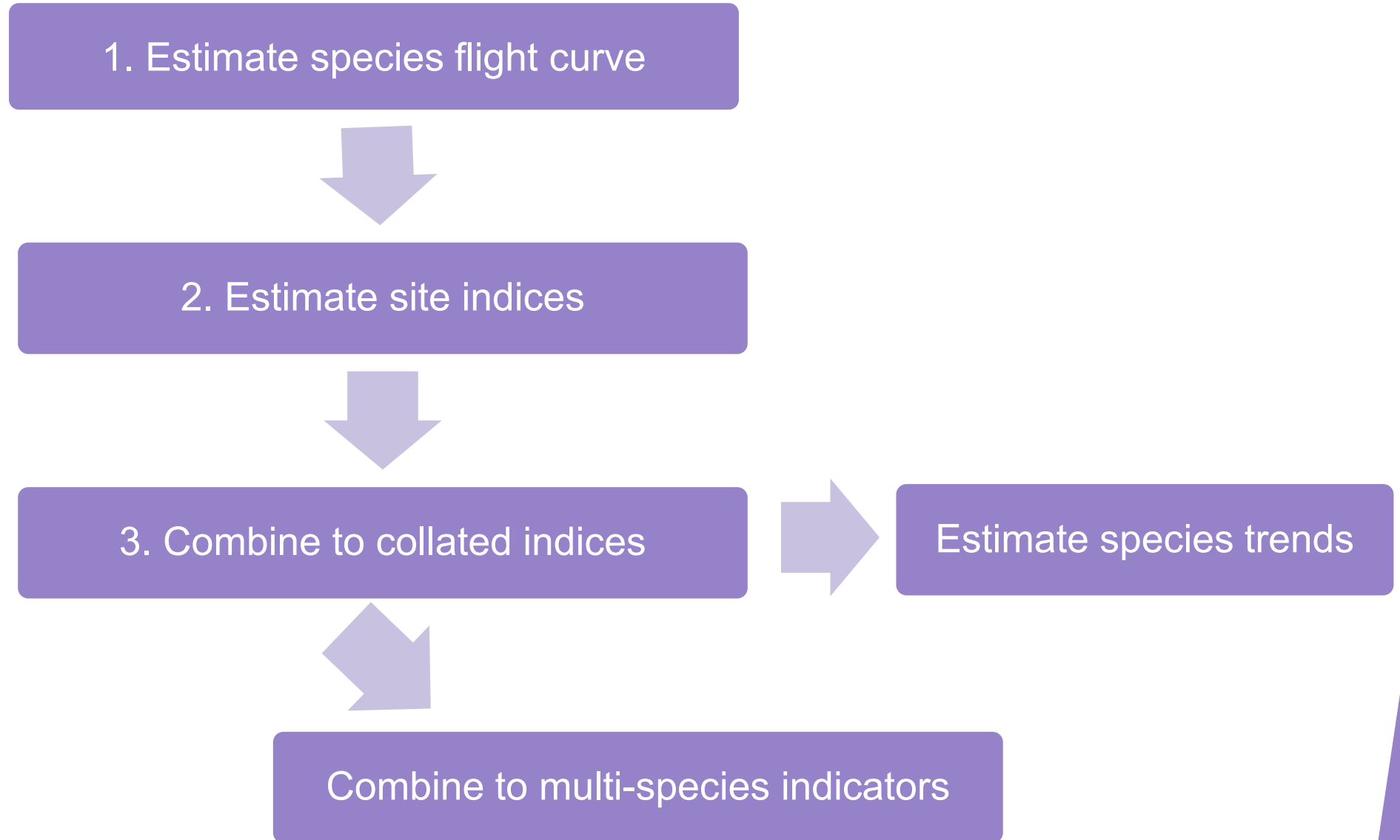
KEY WORDS: Butterflies; Citizen science; Concentrated likelihood; Normal mixtures; Phenology; UKBMS.

<https://doi.org/10.1111/biom.12506> (open)

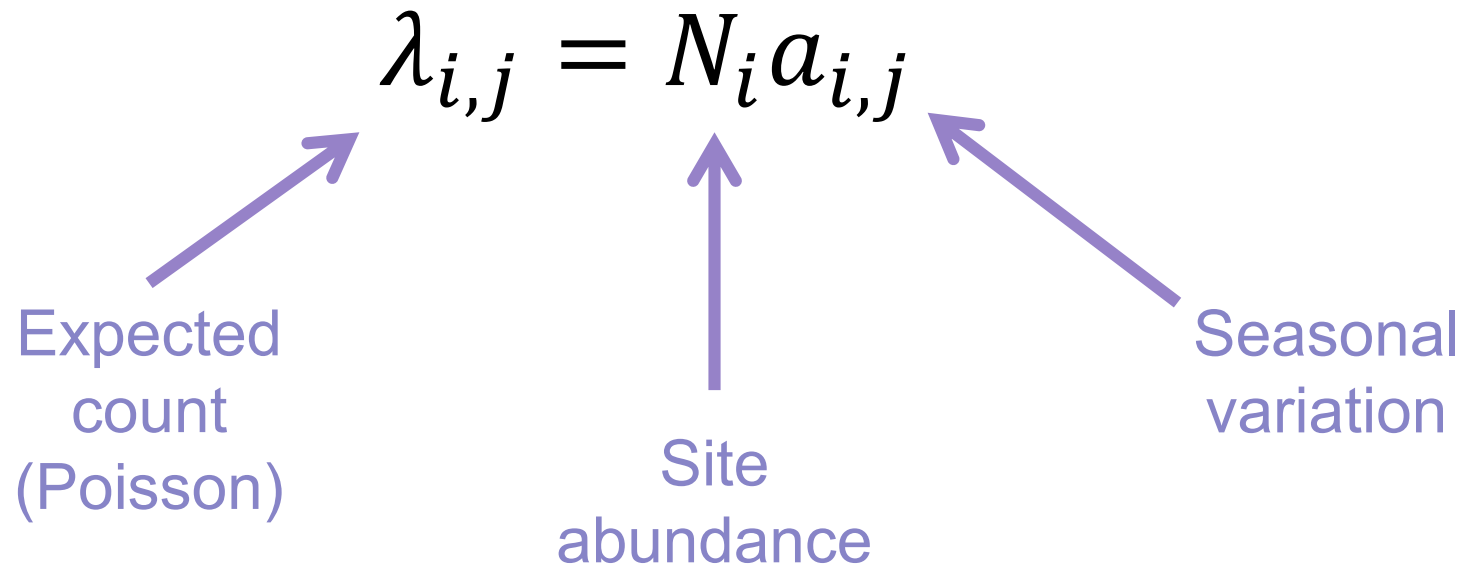
GAI approaches offers three options for describing seasonal variation:

- Flexible spline/GAM across sites each year
- Parametric description for each brood using Normal distributions
 - Phenology estimation, using covariates
- “Stopover model”
 - Mechanistic description with certain assumptions – estimates adult lifespan
 - Relevant paper: <https://doi.org/10.1111/1365-2664.12208> (open)

Overview of modelling steps

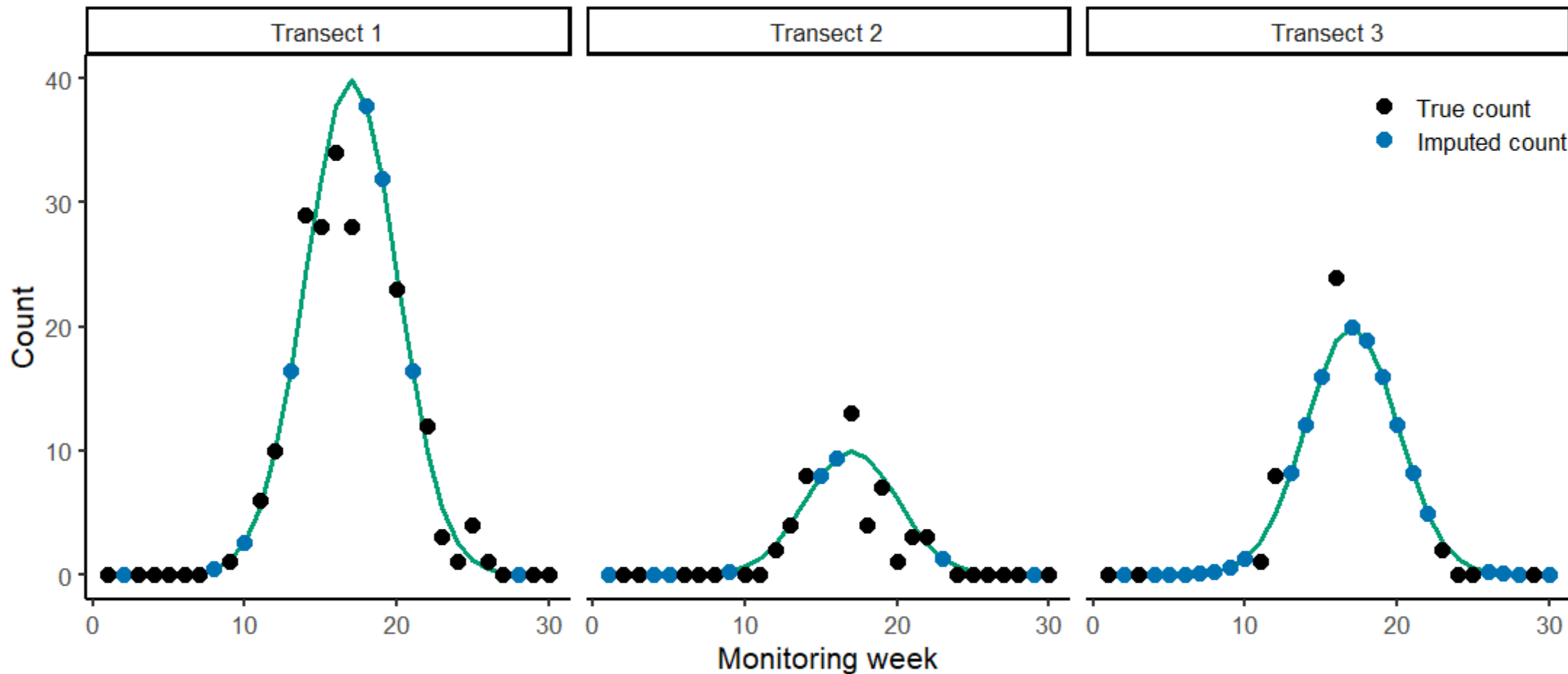


1. Estimate species flight curve



- We use a flexible curve to describe the species flight curve across sites
- The “height” of the curve is reflected by the site effect
- Efficient modelling of the site effects N_i - concentrated likelihood – up to 75 times faster

1. Estimate species flight curve



2. Combine to site indices

For each year:

$$N_i = \frac{\sum_j y_{i,j}}{\sum_j a_{i,j}}$$

Real counts, summed
across visits

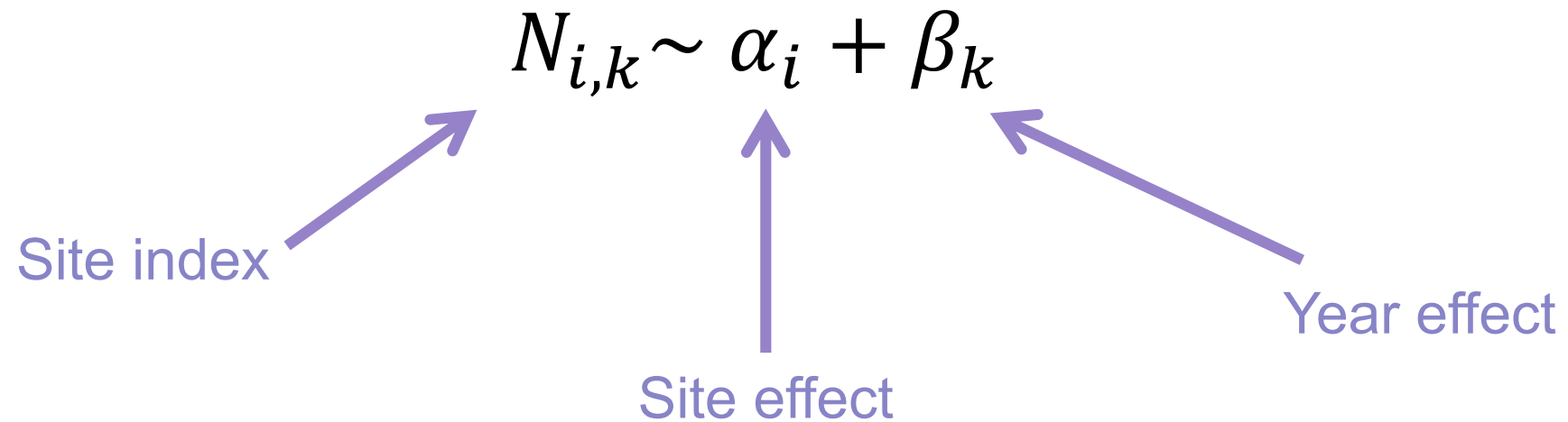
Site index

Estimated season pattern,
summed across visits

Site indices for each site and year for a given species

3. Combine to collated indices

We fit a Poisson generalised linear model (GLM):

$$N_{i,k} \sim \alpha_i + \beta_k$$


The diagram illustrates the components of the Poisson GLM equation $N_{i,k} \sim \alpha_i + \beta_k$. Three purple arrows point from labels below to the terms in the equation: an arrow from 'Site index' points to α_i , an arrow from 'Site effect' points to α_i , and an arrow from 'Year effect' points to β_k .

- Also weight by the proportion of the flight curve sampled
 - Well sampled sites contribute more
- We can then obtain expected total butterfly counts
- And convert to collated indices to consider changes in relative abundance over time

Other points to add/for later

- Trends
- Bootstrapping
- Indicator methodology