Method matters: pitfalls in analyzing phenology from occurrence records

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Abstract

Large occurrence datasets provide a sizable resource for ecological analyses, but have substantial limitations. Phenological analyses in Fric et al. (2020) were misleading due to inadequate curation and improper statistics. Our reanalysis of 22 univoltine species with sufficient data for independent analysis found substantive differences in macroscale phenological patterns.

Method matters: pitfalls in analyzing phenology from occurrence records

Running title: Caution needed in GBIF phenological analyses

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Data accessibility statement: No new occurrence data were used; only the data file provided in Fric et al. (2020) supplemental files (https://doi.org/10.1111/ele.13419). Voltinism data and sources are provided in Supplemental Table 1. All data and code used in this reanalysis are available on github at*https://github.com/RiesLabGU/Larsen-Shirey2020_ EcoLettersComment.*

Abstract:

Large occurrence datasets provide a sizable resource for ecological analyses, but have substantial limitations. Phenological analyses in Fric et al. (2020) were misleading due to inadequate curation and improper statistics. Our reanalysis of 22 univoltine species with sufficient data for independent analysis found substantive differences in macroscale phenological patterns.

Main Body:

The aggregation of large datasets from museum records and community science provides a valuable resource for macroscale ecological analyses. However, such data include spatiotemporal and taxonomic biases that must be addressed (Troudet *et al.* 2017; Ries *et al.* 2019). Given these biases, proper data curation and appropriate modelling strategies are necessary to ensure valid inferences.

Fric et al. 2020 used occurrence data for 100 species aggregated in the Global Biodiversity Information Facility (GBIF) in temperate regions of North America and Europe to track phenology across latitudes. Estimating phenology metrics and trends from large occurrence datasets is possible, but requires sufficient data density and appropriate statistical methods (Taylor and Guralnick 2017). The data from this study were frequently too sparse and insufficiently curated to estimate phenological patterns across latitudes. Further, using regression of residuals resulted in spurious patterns; after correcting for altitude and year, onset and termination phenology appeared the same at low and high latitudes for most species, contrary to previous findings (Karlsson 2014, Matechou et al. 2014). We show that applying appropriate data curation and methods, most species demonstrated later onset and shorter flight periods at higher latitudes.

Many species analyzed (in 105 datasets separated by continent) in Fric et al. (2020) had insufficient data for independent analysis. Data were analyzed with as few as 15 occurrence records across >20 degrees latitude and >100 years. Phenological "onset" and "termination" of flight periods were extracted simply as the first and last day-of-year (DOY) values within latitudinal bands, pooled across all years and altitudes. Pooling data increased spatiotemporal bias, lowered the resulting power to detect patterns, and resulted in only one observation date being used as both "onset" and "termination" of flight periods (resulting in one-day flight periods and "peak flight") in an average of 20% of latitudinal bands per species (Figure 1).

Fric et al.'s data curation was inadequate regarding spatial precision and outlier detection. Altitudes were extracted using imprecise GIS coordinates, sometimes representing sea floor or vague place names (eg, "Mt Shasta") and skewed left, giving high altitude observations outsized leverage in regressions. Temporal outliers were problematic; one species' onset at 68° N was in January, when the next occurrence across all latitudes was in June. No sources were cited for species traits, and we found evidence documenting additional generations in portions of their range for 22 species identified as obligate univoltine (Supplemental Table 1).

Finally, the analytical approach in Fric et al. (2020) produced biased results. Beyond regressing individual species' phenometrics against latitude, altitude, and year separately, regression of residuals was used for corrected regressions. This resulted in biased parameter estimates due to collinearity among explanatory variables and reduced statistical significance (Freckleton 2002). Results suggested most species' onset (67 datasets) and termination (71 datasets) were similar across latitudes (Figure 2). These results were surprising, considering well-documented delayed and/or shortened flight periods at high latitudes (Karlsson 2014).

We sought to validate those results with a more robust analysis, applying stricter data standards and curation. For 72 species (76 datasets) we confirmed as univoltine, we filtered data for altitude (0-500m) and timing (March-November). We calculated phenometrics for year-latitude combinations with at least 10 observations. Only 22 datasets, all European, met these requirements in at least three latitudinal bands (Supplemental Table 1). For these, onset and termination were estimated from a Weibull distribution using R package phest (Pearse 2017) and bounded by days (60,330). To estimate unbiased parameters, we modeled each species phenometric using multiple regression (DOY ~ latitude + year) using R version 3.6.2 (R Core Team 2019). We compared our results to those from Fric et al. Supplemental Table 2.

We were unable to validate most patterns reported in the original study. Our results varied substantially in

both onset and termination across species (Figure 2). In contrast to Fric et al. (2020), we found significantly later and/or shorter flight periods at higher latitudes for most species. These new results were consistent with the latitudinal gradient in climate and growing season length (Kobayashi et al. 2016). This evidence of inaccurate phenological patterns also discredits Fric et al.'s downstream trait analyses.

Despite this critique, we recognize that occurrence data have great potential to address many ecological questions. New aggregations of large datasets provide valuable inputs for macroscale ecological research, and the sheer amount of data accumulated across time and space may provide statistical power. However, "with great power must come great responsibility" (Lee 1962); robust scientific inference requires careful data curation and robust analytical models. Other phenology metrics are less confounded by abundance and effort (Belitz et al. 2020); integrated community models with random species effects or informed priors better suit community phenological analyses (Ellwood et al. 2012). We enthusiastically support continued digitization and use of collection data in ecological analysis, but urge researchers to exercise caution when using these data.

Acknowledgements

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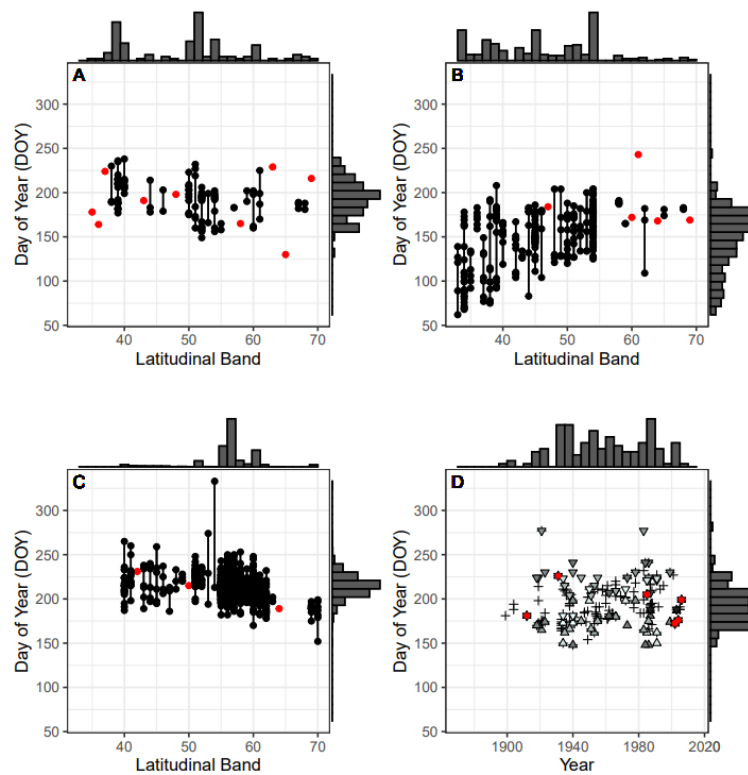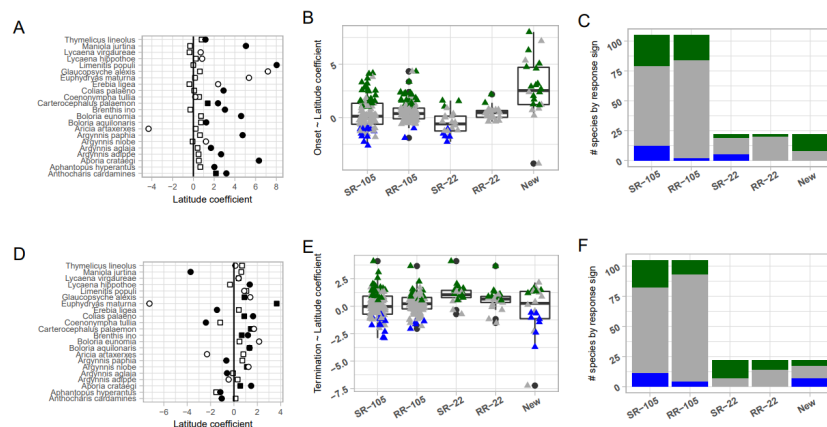
Figure 1:

Figure 2:

Figure 1 Caption:

Raw occurrence data used in the Fric et al. (2020) analysis and flight period lengths by latitude for 3 datasets (A: *Agriades glandon* in N. America, B: *Glaucopsyche lygdamus* in N. America, C: *Hesperia comma* in Europe) and year for 1 dataset (D: *Parnassius smintheus* in N. America). Results for these species were presented in Fric et al. (2020) Figure 1 to demonstrate varied phenological patterns. Red points represent observations used as both onset and termination. Marginal histograms show data density across latitudes (above the plot) and time of year (to the right of the plot). Because onset and termination analyses used only one day of year (DOY, calculated as SuccDay) per latitudinal band, panel D identifies onset $()$ and termination (V) observations $(fill colored by latitude where darker colors represent higher latitudes)$ in addition to all observ
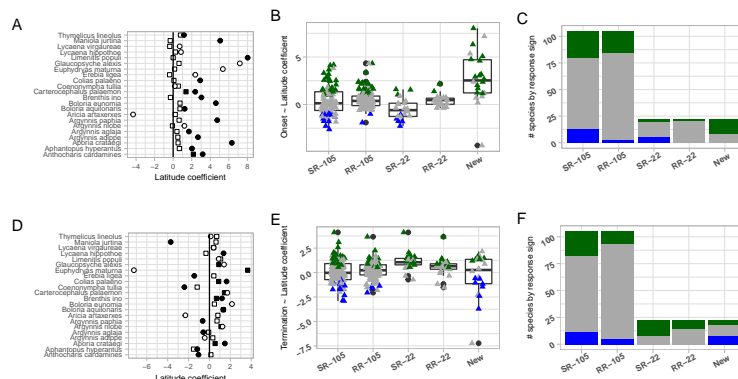
Figure 2 Caption:

Comparison of model parameters for onset (panels A, B, C) and termination (panels D, E, F) dates as a function of latitude. Panels A and D show latitude coefficients for 22 species onset and termination phenology. Values extracted from Fric et al. (2020) Supplemental Table 2 using regression of residuals against latitude, corrected for year and altitude, are shown as squares. Circles are parameters from the reanalysis. Filled symbols indicate significant slopes with latitude. Panels B and E show values and boxplots of the model coefficients for DOY ~ latitude, colored by response: positive (green), non-significant (gray), or negative (blue) correlations. Panels C and F are stacked barplots of the response signs representing the number of datasets demonstrating positive (green), non-significant (gray), or negative (blue) correlations with latitude. Panels B, C, E, and F show four sets of model results extracted from Fric et al. Supplemental Table 2, and one set of reanalysis results. For Fric et al. (2020) results, "SR" represents single regressions for DOY ~ latitude, while "RR" represents regression of residuals against latitude, corrected for altitude and year. "SR-105" and "RR-105" show results for all 105 datasets in the original analysis, while "SR-22" and "RR-22" only include results for the 22 species-region datasets used in our reanalysis. "New" shows results from our curation and reanalysis of those 22 species-region datasets. See Supplemental Figure 1 for a comparison of species-specific data and results.

Supplemental Figure 1 Caption:

These panels display the data and regression results for onset and termination across models. Each row of 3 panels represents a species in the re-analysis. In the left column, all raw occurrence data are shown as "+" while onset and termination are shown as ?; red points indicate individual observations used as both onset and termination. Fric et al.'s single regression results of DOY ~ latitude for onset and termination are overlaid on the data. In the middle

column, the residuals used in the Fric et al. (2020) regression of residuals are shown for onset
$()$ and termination (V), respectively, with red indicating records of observations used as both onset and termination. Because each
significant (black, dotted), or negative (blue, dashed) correlations with latitude in the corresponding models. While the Fric et al.
0.05 for the following single regression analyses : E. maturna onset and L. virgaurea termination.

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`LarsenShirey_SuppFig1.pdf` available at https://authorea.com/users/309075/articles/502821-method-matters-pitfalls-in-analyzing-phenology-from-occurrence-records