DIALECTICS IN NONAUTONOMOUS MATRIX POPULATION MODELS: ACCURACY OF CALIBRATION VERSUS CERTAIN PREDICTION

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A great advantage of the matrix model for a discrete-structured population, $\mathbf{x}(t) \in \mathbb{R}^n$, is the possibility to calibrate the "projection" matrix $\mathbf{L}(t)$ form the data of only two consecutive counts (at time monts t and t + 1) and to calculate $\lambda_1(\mathbf{L}(t))$, the adaptation measure of he local population under study [1]. This is the power of matrix models as a tool for comparative demography, but here also a methodological problem arise when we have a time series of data and it is necessary to summarise the outcome of the entire observation period. The nonautonomous matrix model represents a finie set of *one-step* matrices, $\mathbf{L}(t)$, each yielding its own set of quantitative characteristics of the population, which may even be contradictory in the forecast of its fate.

The contradictions are eliminated by averaging the set M of nonnegative matrices that forms the basic model equation

 $\mathbf{x}(t+1) = \mathbf{L}(t)\mathbf{x}(t), t = 0, 1, ..., M-1,$ (1) and model logic leads to the problem of *geometric* average [2]. Defined by the life cycle graph for the individuals of a given species, the fixed *pattern* of these matrices deprives this problem of *exact* solution, so that the approximate *pattern-geometric* mean becomes the correct mode of averaging [3] – a novel concept in the theory and practice of modelling biological populations.

For the case where the data bear "reproductive uncertainty" [2] and the calibration yields the whole family of matrices, $\{L(t)\} = T(t) + \{F(t)\}$, at each time step, a heuristic method of averaging is proposed, namely, *TF-averaging*. It enables calculating uniquely the pattern-geometric mean of the transition matrices, T(t), hence gaining certain age-specific traits from the stage-structured model and, in particular, answering the question how specifically short a *short-lived* perennial lives.

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References

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