

SIGNAL OR NOISE? TESTING HYPOTHESES

ABOUT FAUNAL TURNOVER

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PROBLEM

Patterns of faunal turnover have been interpreted as reflecting the impact of external stimuli such as climate change on mammalian evolution (e.g., Bobe *et al.*, 2002, Bobe & Behrensmeyer, 2004). Specifically, the origin of genus *Homo* has been linked to a pulse of turnover events caused by global climate change (Vrba, 1988, 1995).

However, stochastic speciation and extinction processes can also produce pulses of turnover events in the absence of external forcing. There is no agreed upon method for distinguishing signal from noise.

I use birth-death simulations with constant speciation probabilities to explore the factors influencing the detection of turnover pulses. Results can be used to calibrate hypothesis tests based on the fossil record in order to avoid false positives.

METHODS

Simulations: 200 fossil records were simulated (Figure 1A), using the paleotree R package (Bapst, 2012; R Core, 2015). Simulation parameters were drawn from uniform distributions with the ranges shown in Table 1. **Detecting pulses:** Fossil records were divided into time bins (Fig. 1B). Turnover rates for bins were computed using Foote's (2000) per-capita rates of origination. Rates were compared to other bins, excluding the first and last. A turnover pulse criterion (e.g., 1.5 times the interquartile range of origination rates) was applied (Fig. 1C), and any pulses noted (Fig. 1D).

Test of Simulation Parameters: Pulses were detected in each fossil record using time bins of 0.5 Ma and a pulse criterion of $1.5 \times \text{IQR}$. ANOVA was used to test for a relationship between simulation parameter values and the number of pulses detected. **Test of Analytical Parameters:** Pulses were detected in each fossil record with time bins of 0.5 Ma and a pulse criterion of 1.5. The proportion of fossil records showing at least one pulse was noted. Analytical parameters were incremented over the range in Table 1. At each increment, the pulse detection procedure was re-applied using the incremented analytical parameter. Loess regression was used to visualize the relationship between parameter values and the proportion of records showing a pulse.

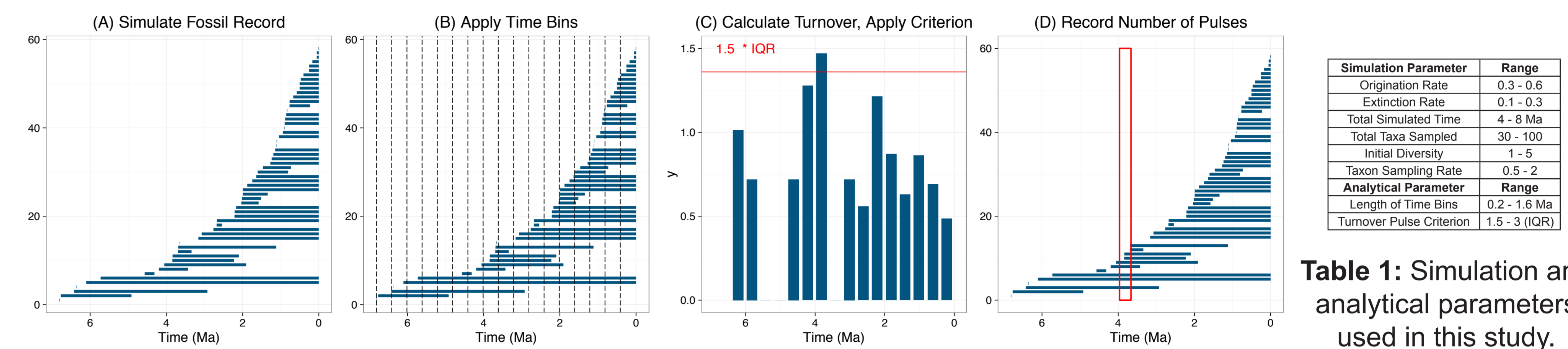


Figure 1 A-D: Schematic illustration of methods.

RESULTS

SIMULATION PARAMETERS

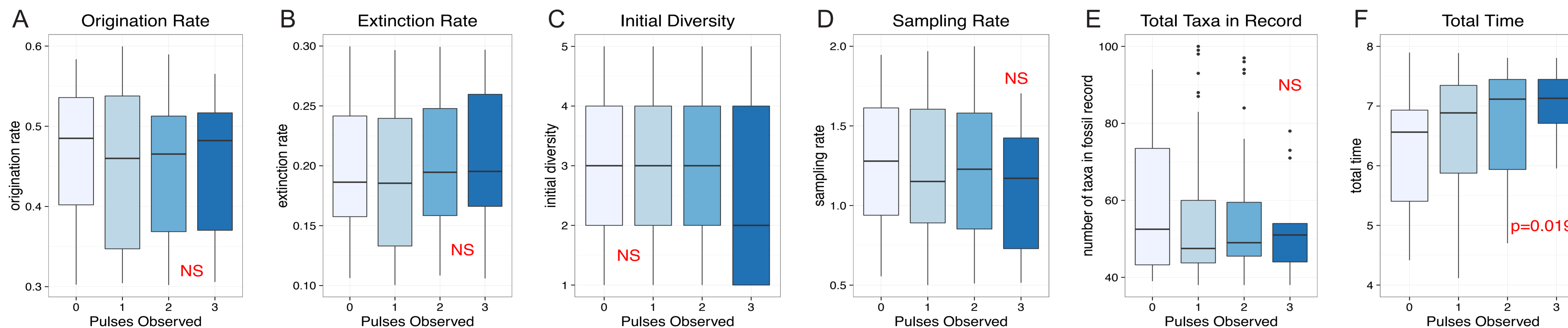


Figure 2 A - F: Boxplots showing relationship between simulation parameters and number of pulses detected. Significance test are from ANOVA.

ANALYTICAL PARAMETERS

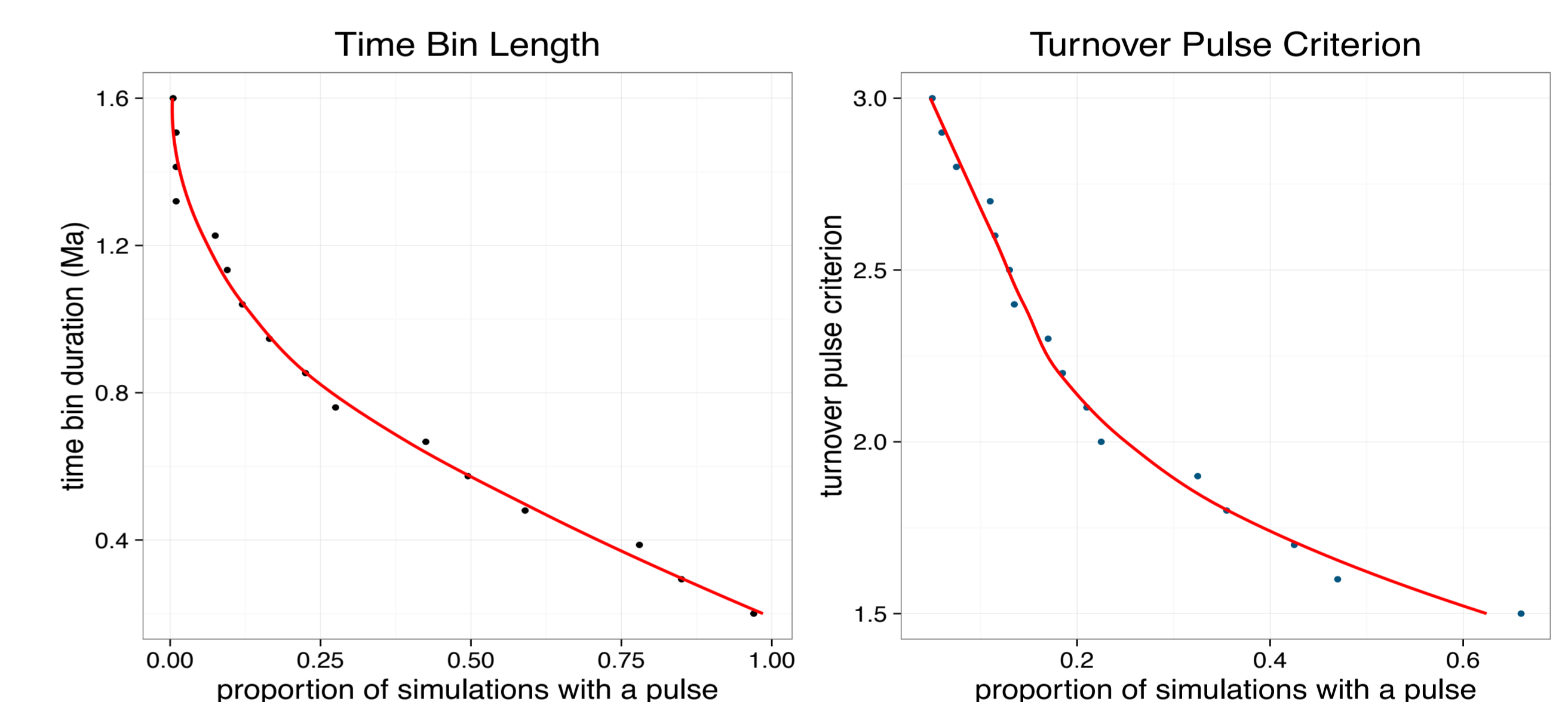


Figure 3 A and B: Regressions showing relationship between analytical parameters, and proportion of records showing a turnover pulse.

DISCUSSION

Most of the simulation parameters do not significantly predict the number of turnover pulses detected (Fig 2). The only significant predictor of the number of pulses was the total time of the simulation (Fig 2F). Other things being equal, longer simulations tend to have more turnover pulses. This is likely because longer simulations reduce the average number of taxa per bin, which results in a noisier dataset.

While the total number of taxa is not a significant predictor of the number of pulses, high numbers of taxa are rarely associated with high numbers of pulses.

Extinction and origination rates do not have a consistent impact on number of pulses observed. This is likely because the simulations are constrained in the number of taxa allowed. If the number of taxa were unconstrained, higher origination rates and lower extinction rates would increase the total number of taxa, and the number of taxa per time interval. This would likely reduce the number of pulses.

The analytical parameters (bin length and the pulse criterion) have very clear relationships to the number of pulses detected (Fig 3). More stringent criteria and longer time bins rapidly reduce the probability of detecting a pulse.

CONCLUSIONS

Analytical parameters have a profound impact on probability of detecting turnover pulses. Time bin length and turnover pulse criterion must be chosen carefully to ensure reasonable Type I error rates.

There are potentially high Type I error rates in simulations with parameters similar to estimates from the African bovid fossil record. This underscores the recent conclusion of Bibi & Kiessling (2015) that bovid turnover is best viewed as relatively continuous.

REFERENCES

Bibi F, and Kiessling W. 2015. Continuous evolutionary change in Plio-Pleistocene mammals of eastern Africa. *PNAS* 112:10623–10628.
Bapst DW. 2012. paleotree: an R package for paleontological and phylogenetic analyses of evolution. *Methods in Ecology and Evolution* 3:803–807.
Bobe R, and Behrensmeyer AK. 2004. The expansion of grassland ecosystems in Africa in relation to mammalian evolution and the origin of the genus *Homo*. *Palaeogeography, Palaeoclimatology, Palaeoecology* 207:399–420.
Bobe R, Behrensmeyer A, and Chapman R. 2002. Faunal change, environmental variability and late Pliocene hominin evolution. *Journal of Human Evolution* 42:475–497.
Foote M. 2000. Origination and extinction components of taxonomic diversity: general problems. *Paleobiology* 26:74–102.
R Development Core Team. 2015. R: A language and environment for statistical computing. Vienna, Austria: R Foundation for Statistical Computing.
Vrba ES. 1988. Late Pliocene climatic events and hominid evolution. In: Grine FE, editor. *Evolutionary History of the "Robust" Australopithecines*. New York: Aldine, p 405–426.
Vrba ES. 1995. The Fossil Record of African Antelopes (Mammalia, Bovidae) in Relation to Human Evolution and Paleoclimate. In: Vrba ES, Denton GH, Partridge TC, Burckle LH, editors. *Paleoclimate and Evolution with Emphasis on Human Origins*. New Haven: Yale University Press, p 383–424.

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