



# Bayesian Population Analysis using WinBUGS: A hierarchical perspective

*Marc Kery, Michael Schaub*

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Bayesian statistics has exploded into biology and its sub-disciplines, such as ecology, over the past decade. The free software program WinBUGS, and its open-source sister OpenBugs, is currently the only flexible and general-purpose program available with which the average ecologist can conduct standard and non-standard Bayesian statistics.

- Comprehensive and richly commented examples illustrate a wide range of models that are most relevant to the research of a modern population ecologist
- All WinBUGS/OpenBUGS analyses are completely integrated in software R
- Includes complete documentation of *all* R and WinBUGS code required to conduct analyses and shows *all* the necessary steps from having the data in a text file out of Excel to interpreting and processing the output from WinBUGS in R

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