

*Ninth Workshop Dynamical Systems Applied
to Biology and Natural Sciences DSABNS 2018
Turin, Italy, February 7-9, 2018*

STOCHASTIC MODELING OF BIOLOGICAL POPULATIONS THROUGH BRANCHING MODELS. APPLICATION TO BLACK VULTURE COLONIES

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We investigate appropriate stochastic models to describe the population dynamics of some biological species. We develop a discrete-time branching model which is indexed by the time instead of the generation, as usual in branching process literature [2]. In particular, by considering approximate Bayesian computation methods, we estimate some relevant biological parameters. As illustration, we apply such a model to describe the probabilistic evolution of Black Vulture colonies in Extremadura (Spain) [1]

References

- [1] J.C. Del Moral, J. De la Puente. (2017). *Buitre negro- Aegyptius Monachus*. *Enciclopedia virtual de los vertebrados*, Museo Nacional de Ciencias Naturales, Madrid, 2–41.
- [2] P. Jagers. (1975). *Branching Processes with Biological Applications*, John Wiley and Sons, London/New York.