

# BeeNestABM: An open-source agent-based model of spatiotemporal distribution of bumblebees in nests

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## Software

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## Licence

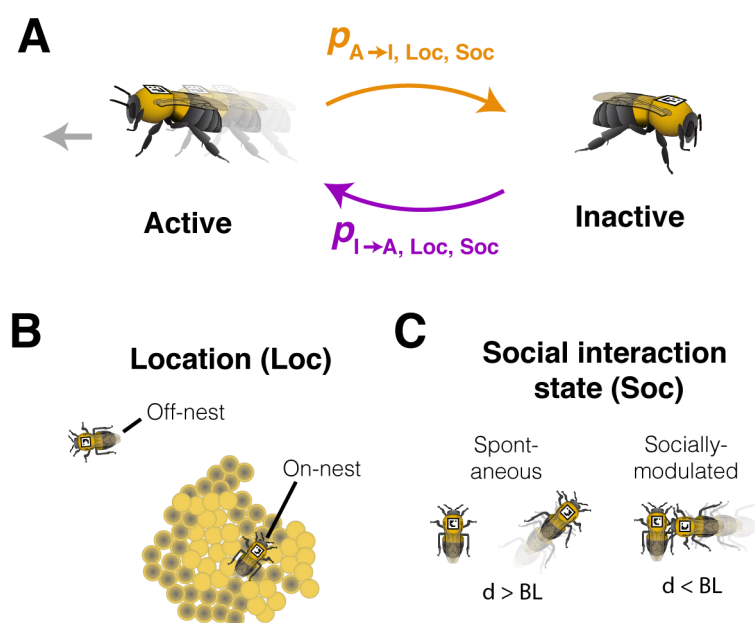
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## Summary

This software features the MATLAB source code for an interactive computational model that can be used to study the localized responses of bumblebees to sublethal exposures to a prevalent class of pesticides called neonicotinoids. The code involves an agent-based stochastic model for the interactions between and movements of individual bees within a nest and the nest-related disruptions that occur due to pesticide exposure. The dynamic states of the bees are stored in a matrix, the default data structure of MATLAB. Agent-based modeling allows for building understanding of the colony scale impacts of multiple interacting factors that affect numerous individuals in close proximity and how those change upon pesticide exposure. The scientific significance is that the model solved in the software focuses on the effects of pesticides that occur in bumblebees over short time scales (hours to days) inside a single nest, which includes a much smaller spatial region with finer resolution than that considered in the state-of-the-art (Thorbek, Campbell, and Thompson 2017). These temporal and spatial scales are appropriate for modeling the effects of neonicotinoid pesticides that account for colony size and interactions between exposed and unexposed individuals. The short and local scales allow the model to explicitly consider neighbor interactions and individual bee interactions with structures inside an isolated environment without confounding external factors. Novel research results using this software for scientific applications have been obtained (Crall, Switzer, et al. 2018, Crall, deBivort, et al. (2018)).

## References

- Crall, J. A., B. L. deBivort, B. Dey, and A. N. Ford Versypt. 2018. "Social buffering of pesticides in bumblebees: agent-based modeling of the effects of colony size and neonicotinoid exposure on nest behavior." *Submitted to Frontiers in Ecology and Evolution*.
- Crall, J. A., C. Switzer, R. Oppenheimer, S. Combes, N. Pierce, B. de Bivort, A. N. Ford Versypt, et al. 2018. "Chronic neonicotinoid exposure disrupts bumblebee nest behavior, social networks, and thermoregulation." *Submitted to Science*.
- Thorbek, P., P. J. Campbell, and H. M. Thompson. 2017. "Colony impacts of pesticide-induced sublethal effects on honeybee workers: A simulation study using beehave." *Environmental Toxicology and Chemistry* 36:831–40. <https://doi.org/10.1002/etc.3581>.



**Figure 1:** A) The BeeNestABM model tracks bumblebee activity and motility using empirically estimated probabilities for transitions between active (mobile) and inactive states. B) The location of a bee in relation to the structures such as brood and food pots within the nest influence the transition probabilities and the orientation of bee movement through a combination of random walk and attraction toward the nest structures. C) The transition probabilities contain a component that considers whether the transition is occurring spontaneously or due to social modulation upon collision with a neighboring bee.