A review of honeybee models and a short introduction to the new integrated colony model BEEHAVE

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Matthias Becher*, Pete Kennedy*, Jenny Swain, Judy Pell, Juliet Osborne*: Rothamsted Research, UK

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Grimm V, Becher MA, Kennedy PJ, Thorbek P, Osborne J. **Ecological modeling for pesticide risk assessment of honeybees and other pollinators**. In: Fischer D, Moriarty T. *Pesticide risk assessment for pollinators: A SETAC Pellston workshop*. SETAC Press, Pensacola, FL (in press)

Becher MA, Thorbek P, Kennedy PJ, Osborne J, Grimm V. **Towards a systems approach for understanding honeybee decline: a stock-taking and synthesis of existing models.** *Journal of Applied Ecology* (in press)

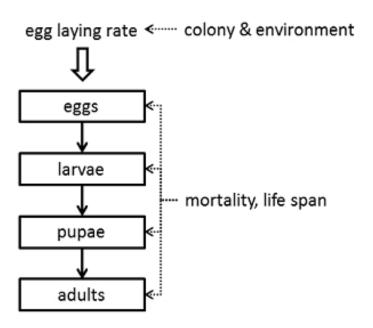
Becher MA, Thorbek P, Kennedy PJ, Grimm V, Osborne J. <**BEEHAVE: an integrated model of honeybees dynamics**>. To be submitted to *Journal of Applied Ecology* (within next 4 weeks)

Honeybee models: overview

Three categories of models:

- 1. Within-hive colony dynamics (8)
- 2. Varroa mite population dynamics within hives (11)
- 3. Foraging (12)

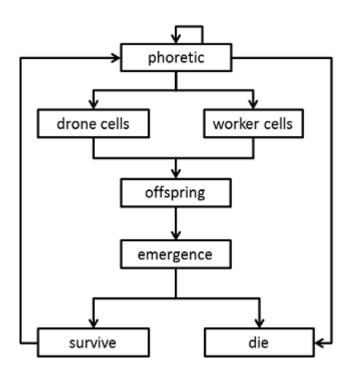
1. Within-hive colony dynamics



Most important models:

- BEEPOP (deGrandi-Hoffmann et al. 1989): beekeeping management
- HoPoMo (Schmickl and Crailsheim 2007): science (drivers, feedbacks)
- Khoury et al. (2011): Impact of forager mortality on colony development

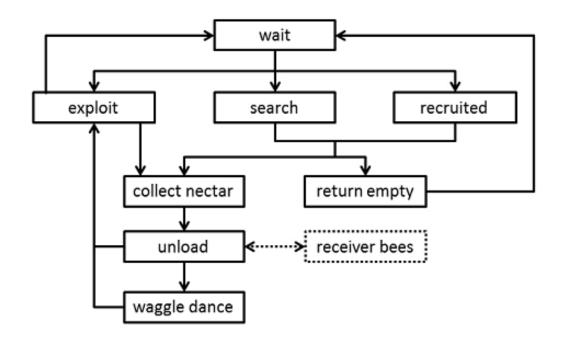
2. Varroa mite population dynamics



Most important models:

- Martin (1998): understand varroa effects on honeybees, beekeeping management
- Martin (2001): Martin (1998) combined with BEEPOP, virus transmission

3. Foraging



Most important models: hard to tell

- Most models use "energetic efficiency" as basis for foraging decisions
- None of the models linked to colony dynamics, explicit landscape structure and dynamics, or pollen collection

Representation of stressors

Factors																			
Factors	Omholt 1986	deGrandi-Hoffman et al. 1986	Martin 2001	AlGhamdi & Hoopingarner 2004	Thompson et al. 2005/2007	Schmickl & Crailsheim 2007	Becher et al. 2010	Khoury et al. 2011	Omholt & Crailsheim 1991	Calis et al. 1999a	Calis et al. 1999b	Boot et al. 1995	Wilkinson & Smith 2002	deGrandi-Hoffman & Curry 2004	Sumpter & Martin 2004	Vetharaniam & Barlow 2006	Vetharaniam 2012	All forager models	ВЕЕНАVЕ
Genetic diversity																			
Varroa mites			+	+				(+)	+	+	+	(+)	+	+	+	(+)	(+)		+
Viruses			+												+				+
Bacterial pathogens																			
Nosema spp.								(+)											+
Loss of forage quantity						+													+
Forage nutritional quality																			+
Beekeeping practice			(+)	(+)				(+)		(+)	(+)		(+)	+	(+)	(+)			+
Pesticides – inside hive					+														+
Pesticides – outside hive					+			(+)											+
Forager death unknown cause		(+)			(+)	+		+					(+)	(+)					+

Conclusions from review

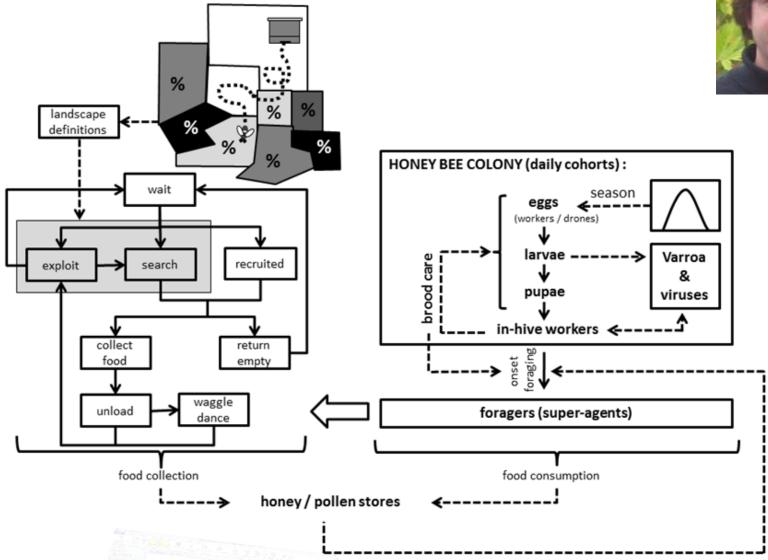
- Model testing, validation, and analysis of most models was very limited
 - No clear separation of imposed and emergent dynamics
 - No clear indication of how much calibration was involved
 - Limited or no sensitivity analysis
- For foraging models, a benchmark test exist: the Seeley et al. (1991) feeder experiment

Conclusions from review

- Well-tested building blocks exist in existing models
- A model that would allow integrating stressors within and outside the hive does not yet exist
- Colony structure and important feedback loops need to be included (e.g., "age of first foraging")
- Egg-laying rate, weather, colony structure, and availability of nectar and pollen should drive the dynamics

BEEHAVE: developed by Matthias Becher





BEEHAVE

Colony module (in-hive, daily time steps):

- Similar to BEEPOP
- Feedbacks: brood care, amount of honey and pollen
- Consumption of nectar and pollen

Varroa module (daily time steps):

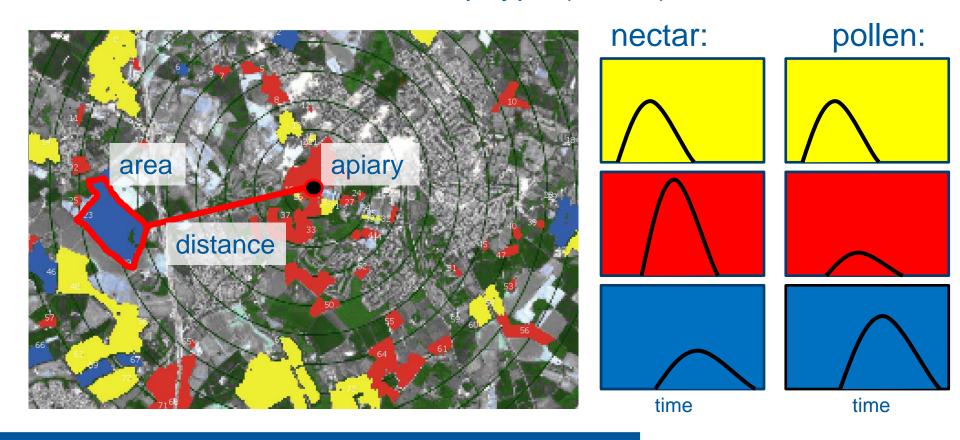
- Similar to Martin (2001)
- Transmits either deformed wing or acute paralysis virus

Foraging module (minutes):

- Driven by energetic efficiency
- Can be linked to hetegerogeneous and dynamics landscape
- Includes pollen collection

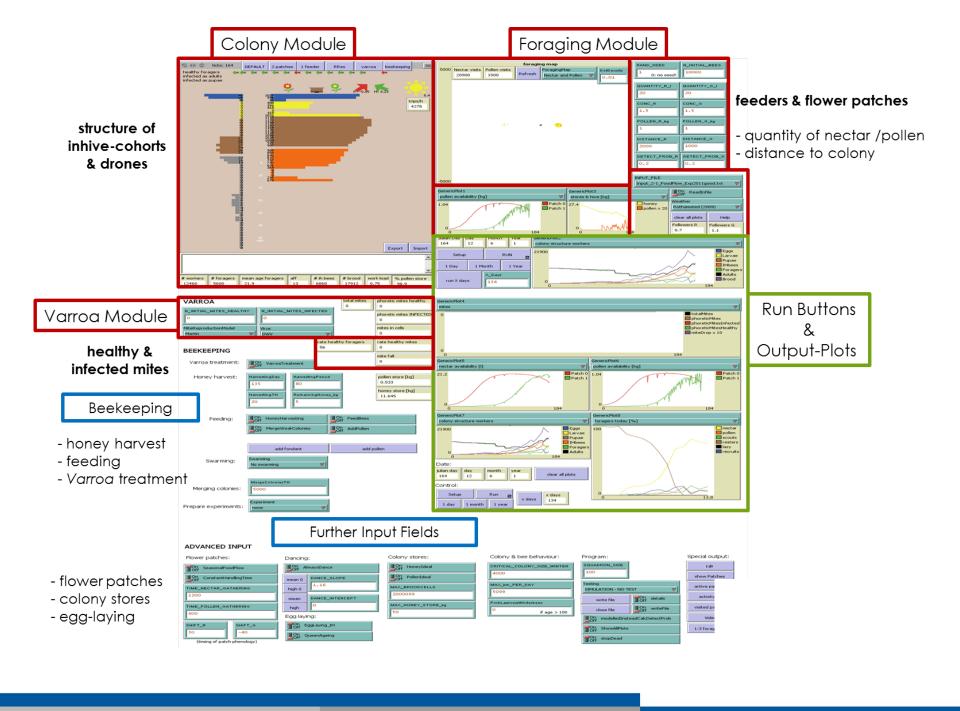
Automated calculation of: - number of patches

- distance to apiary
- area of patch
- chance to find the patch
- crop type (colour)



Documentation, testing, validation

- Implemented in NetLogo (free software platform)
- Documented in ODD format (ca. 40 pages)
- User manual and guided tour exist (ca. 60 pages)
- Extensive testing (debug code, consistency tests, visual output)
- Validation:
 - Age of first foraging, lifespan
 - Number of reproductive cycles of varroa in a year
 - Seeley's feeder experiment



1st Example Scenario

NO VARROA

N initial bees: 10000

2 patches:

Distance: 1500 m 500 m

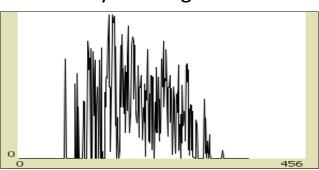
Nectar concentration: 1.5 mol/l 1.5 mol/l

Pollen (max): 1 kg/day 1 kg/day

Nectar flow:

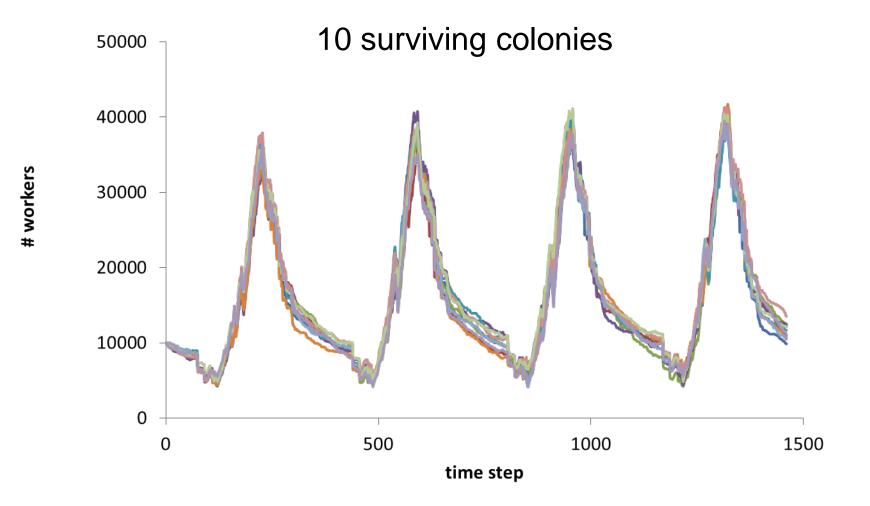
Jan May Aug Dec

Daily foraging period based on real weather:



1st Example Scenario

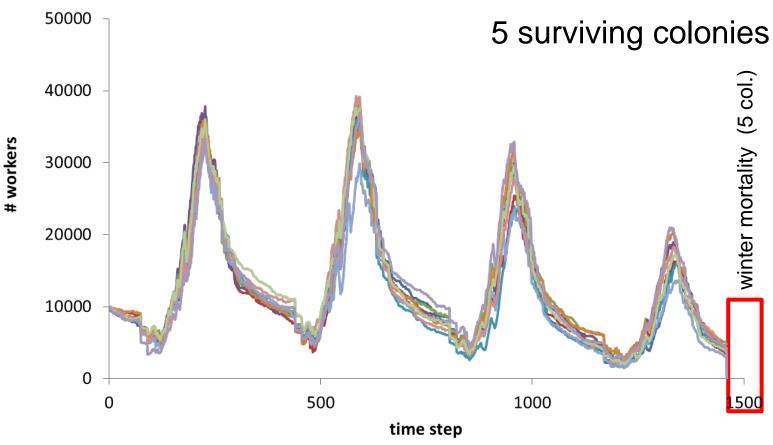
N adult workers (10 colonies) - no varroa - flower patches at 500 & 1500 m



2nd Example Scenario

N adult workers (10 colonies) - with varroa and DWV

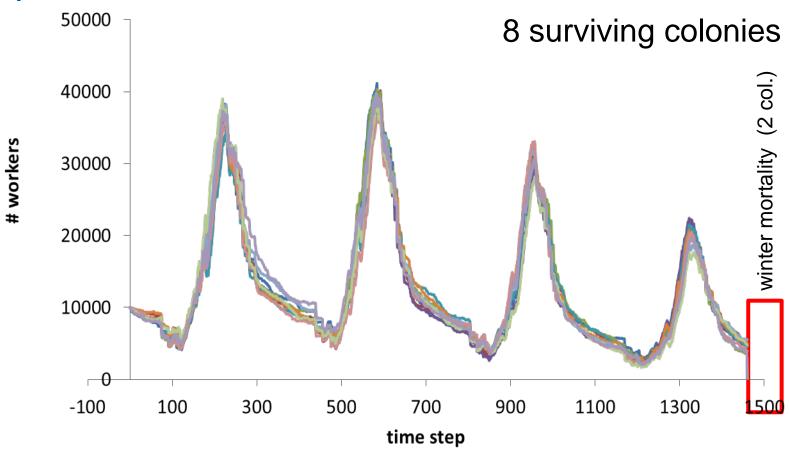
- flower patches at 500 & 1500 m



3rd Example Scenario

N adult workers (10 colonies) - with varroa and DWV

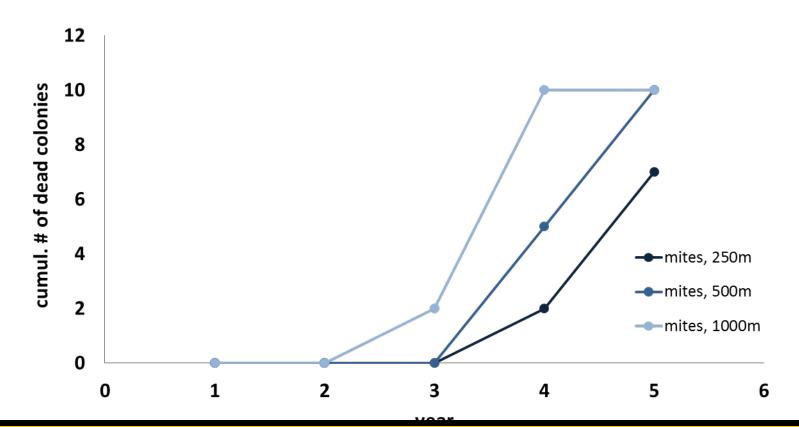
- flower patches at 250 & 1500 m



3rd Example Scenario

N adult workers (10 colonies) - with varroa and DWV

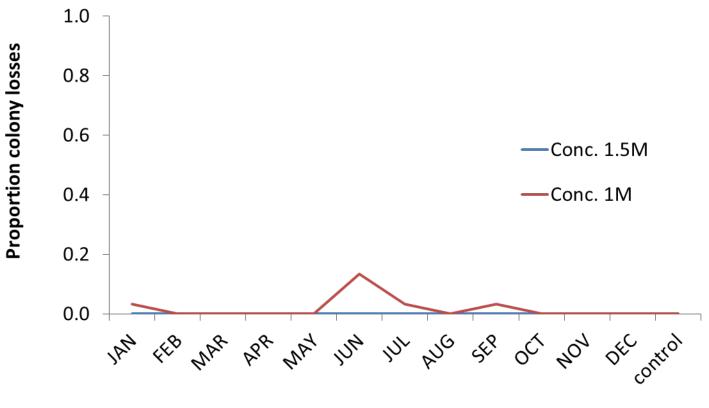
- close flower patch: 250, 500 & 1000m



First combination of foraging AND Varroa model:

Improved food availability can compensate loss of diseased bees

No varroa – nectar feeder at 1000 m - sugar concentration 1.0/1.5 mol/l – double mortality per foraging trip for 30 days (equivalent to increase in forager mortality from ca. 15 to 30%) (similar to **Khoury/Henry** scenario)



Month of treatment

BEEHAVE: to do list

Get it published as soon as possible

- Designed so that others can test and use it
- Offer training courses, workshops

2 PhD students currently working on

- Multiple stressors, landscape structure and dynamics (Juliane Horn, UFZ)
- Specific pesticide module (Jack Rumkee, Univ. Exeter/Syngenta)

From one to many colonies

Summary

- Three types of models (within-hive, varroa, foraging)
- Well-tested modules exist, but no integrated model
- BEEHAVE: first attempt to link within-hive dynamics to foraging in heterogeneous and dynamic landscape
- BEEHAVE (or refinements) would be suitable for regulatory risk assessment

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