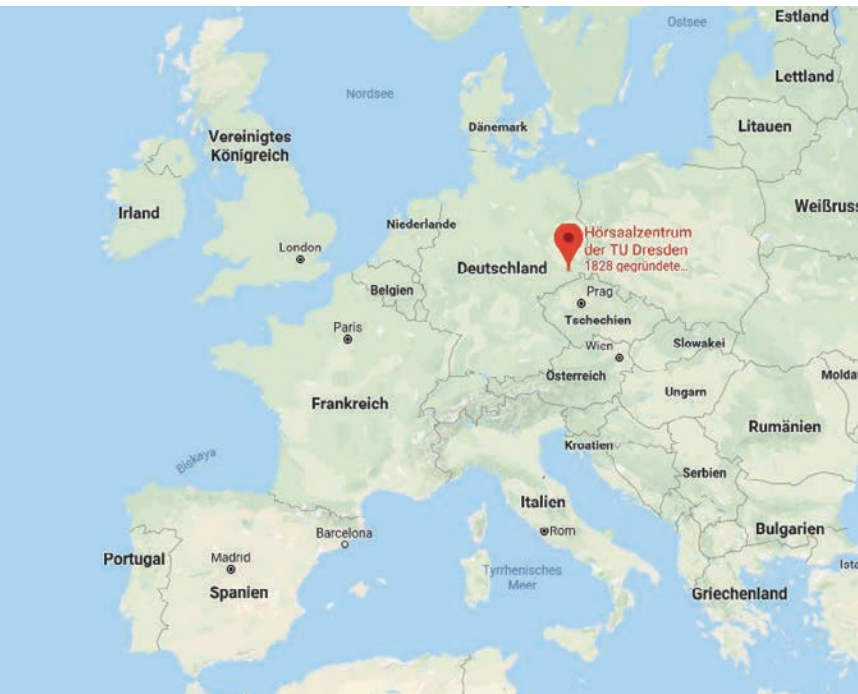


Is our world agent-based and can we model it?

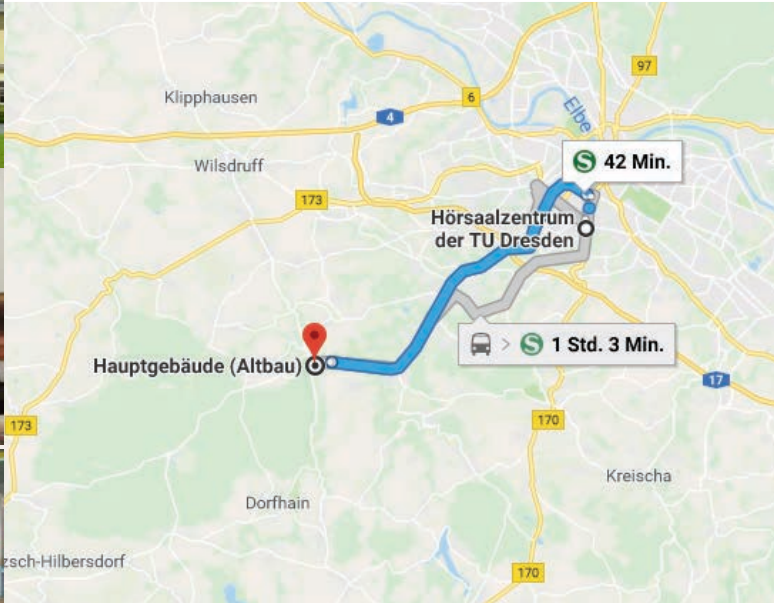
Uta Berger

- Faculty of Environmental Sciences
 - Department of Forest Sciences (Forest Faculty Tharandt)
 - Institute of Forest Growth and Computer Sciences
 - Professorship of Forest Biometry and Systems Analysis



Dresden

Tharandt



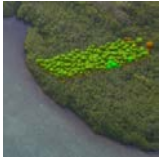
Forest Academy Tharandt

- 2nd oldest in the world (est. by Heinrich Cotta 2011)
- Term “sustainability” established



- ~ 700 students (~20.000 TU Dresden)
- Forest Growth And Computer Sciences, Silviculture and Forest Protection, Forest Botany and Forest Zoology, Soil Sciences, Forest Technics, Forest Economy, Plant Chemistry, ..





Professorship of Forest Biometry and Systems Analysis

Research interest

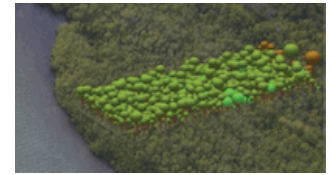
- link between **species adaptations**, **organismic interactions** and long term patterns in **biodiversity** and **systems dynamics**
- **restoration ecology** with a particular focus on **degraded forests**
- concepts of **conservation** under **changing environmental conditions**

Methodologies - Ecological modelling

- individual-based modelling (IBM)
- agent-based modelling (ABM)
- spatial statistics
- ...

Current systems we are working on

- Mangroves (Brazil, South Africa, Thailand, Vietnam)
- Renaturation of oak trees in pine dominated forests (Germany)
- Peat swamp forests (Indonesia)
- Bark beetle invasions (Germany)
- Pine forests (Syria)
- Farm systems in the Andean region (Peru)
- Antibiotica resistance of microbial communities (freshwater, Germany)
- Spiny lobster fishery (Ecuador)
- Krill (Antarctic Ocean)



Is our world agent-based and can we model it?

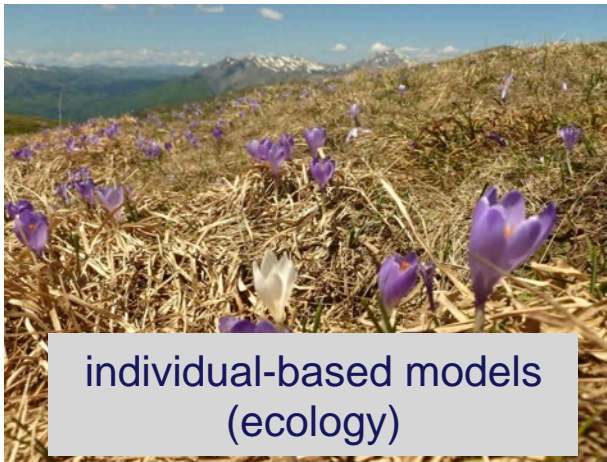
1. Basic principles of IBM / ABM (example models: Behavioural Ecology, Plasmid Biology)
2. IBM / ABM meets Machine Learning (example model: plant ecology)
3. IBM merges ABM towards agent-based sciences



agent-based models
(social sciences)



stand simulators
(forest sciences)



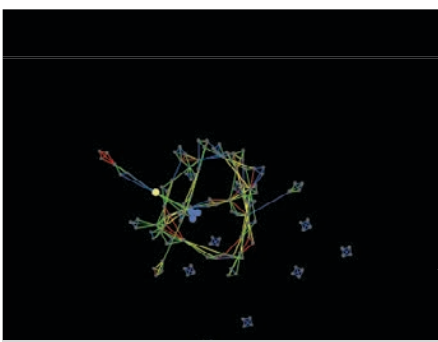
individual-based models
(ecology)

Same objectives

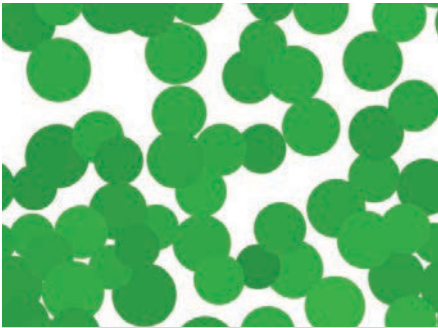
- understanding the functioning of complex systems
- multiple linkages of social-economic-ecological processes
- quantitative forecasts of effects of a changing environment (climate, extreme events, ...)

while considering

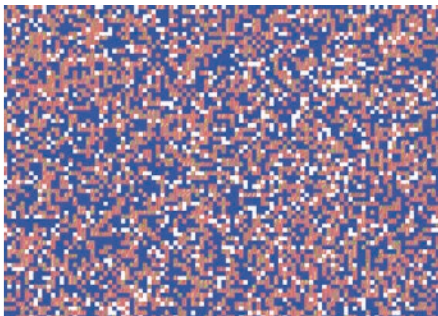
- variability of agents, organisms (plants animals), microbes, .. (entities)
- interactions
- behaviour
- adaptation
- decision making



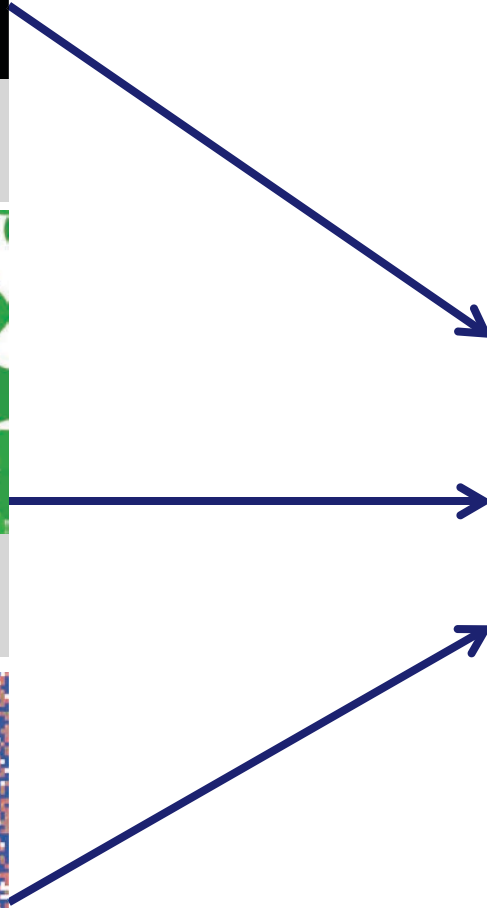
agent-based models
(social sciences)



stand simulators
(forest sciences)



indiv.-based models
(ecology)



12th Summer School in IBM/
ABM at the TU Dresden

1st Example

Behavioural Ecology

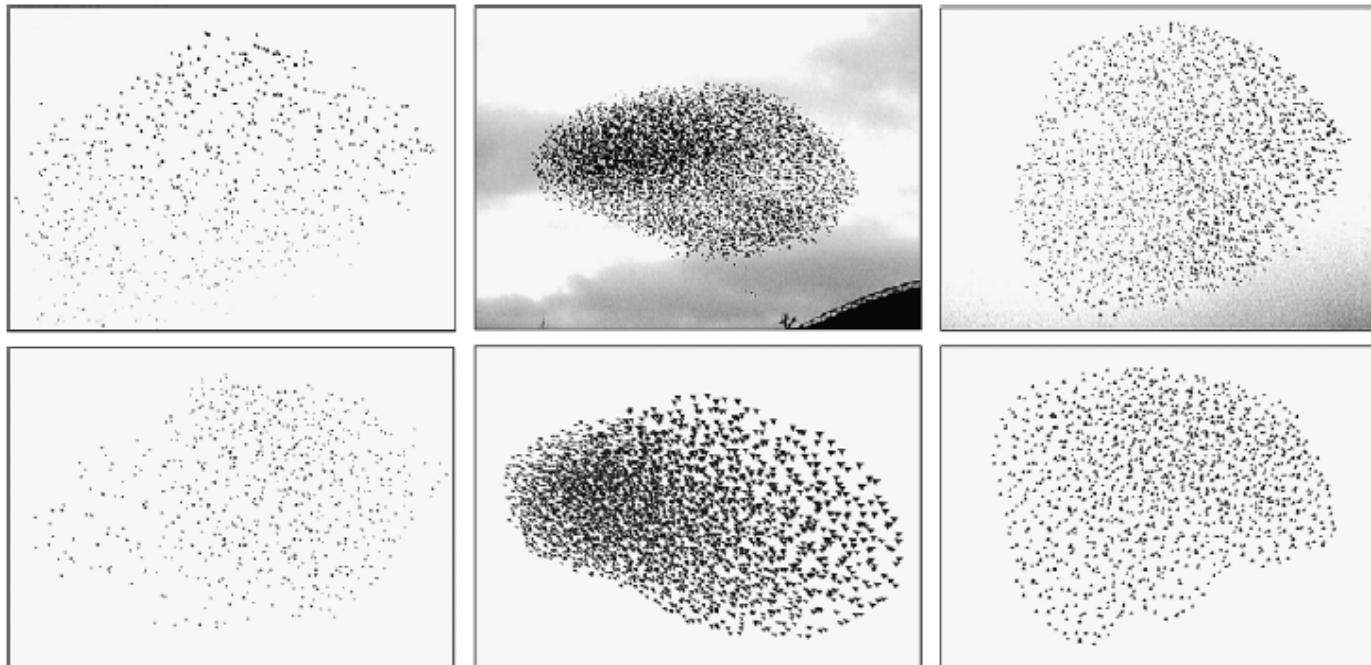
Basic principles of ABM: interactions

Self-organized aerial displays of thousands of starlings: a model

H. Hildenbrandt,^a C. Carere,^{b,c} and C.K. Hemelrijk^a

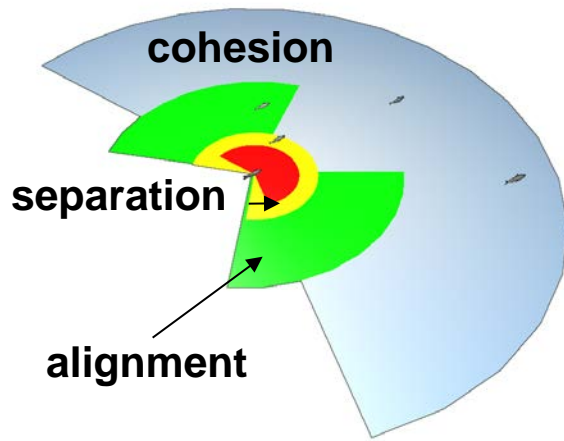
^aTheoretical biology; Behavioural Ecology and Self-organisation, Centre for Ecological and Evolutionary Studies, University of Groningen, PO Box 14, 9750 AA, Haren, The Netherlands, ^bCNR-INFM, Dipartimento di Fisica, Università di Roma La Sapienza, P.le A. Moro 2, 00185 Roma, Italy, and ^cDipartimento di Ecologia e Sviluppo Economico Sostenibile Università degli Studi della Tuscia, Viterbo, Italy

Through combining theoretical models and empirical data, complexity science has increased our understanding of social behavior of animals, in particular of social insects, primates, and fish. What are missing are studies of collective behavior of huge swarms of birds. Recently detailed empirical data have been collected of the swarming maneuvers of large flocks of thousands of starlings (*Sturnus vulgaris*) at their communal sleeping site (roost). Their flocking maneuvers are of dazzling

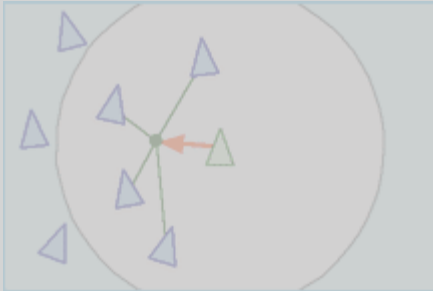




a starling flocks at the sky of Termini (Italy)

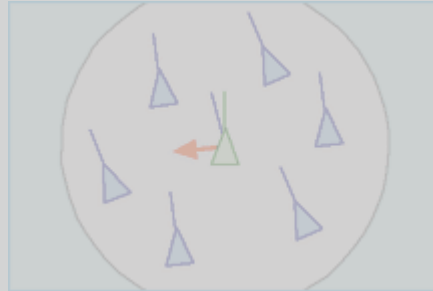


cohesion



$$\vec{F}_{cohesion} = \left(\frac{1}{n} \sum_n \vec{p}_n \right) - \vec{p}$$

alignment

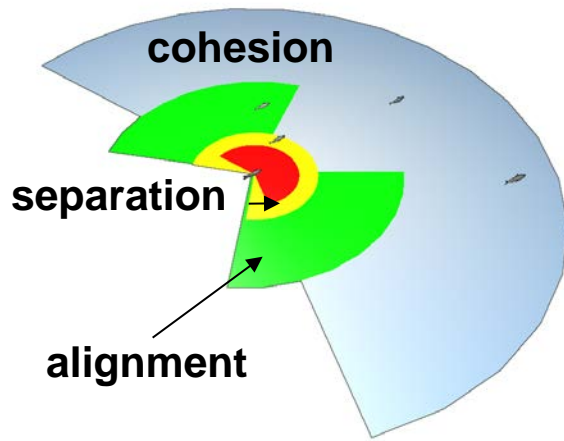


$$\vec{F}_{align} = \left(\frac{1}{n} \sum_n \vec{f}_n \right) - \vec{f}$$

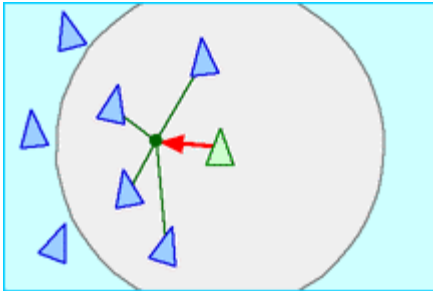
separation



$$\vec{F}_{separation} = \left(\frac{1}{n} \sum_n \frac{-(\vec{p}_n - \vec{p})}{\|\vec{p}_n - \vec{p}\|^2} \right)$$

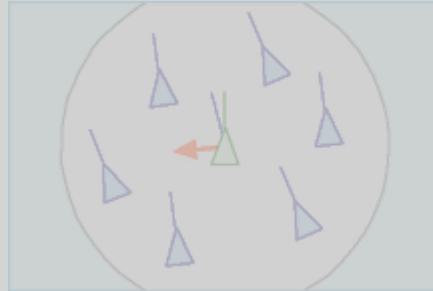


cohesion



$$\vec{F}_{cohesion} = \left(\frac{1}{n} \sum_n \vec{p}_n \right) - \vec{p}$$

alignment

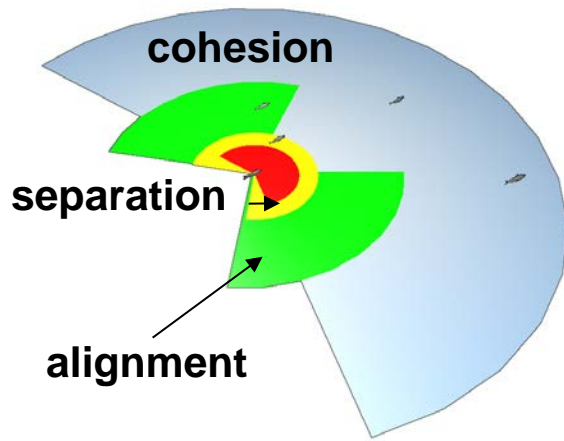


$$\vec{F}_{align} = \left(\frac{1}{n} \sum_n \vec{f}_n \right) - \vec{f}$$

separation



$$\vec{F}_{separation} = \left(\frac{1}{n} \sum_n \frac{-(\vec{p}_n - \vec{p})}{\|\vec{p}_n - \vec{p}\|^2} \right)$$



cohesion

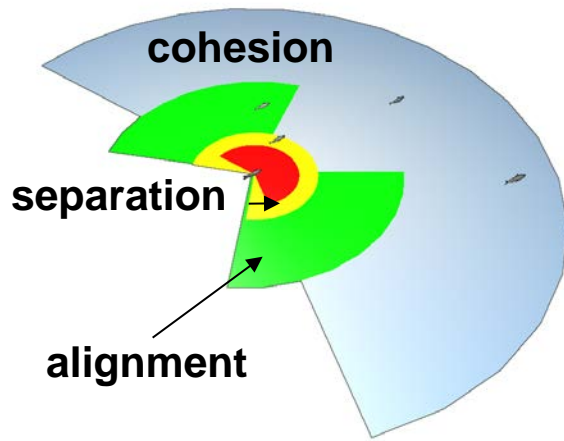
$$\vec{F}_{cohesion} = \left(\frac{1}{n} \sum_n \vec{p}_n \right) - \vec{p}$$

alignment

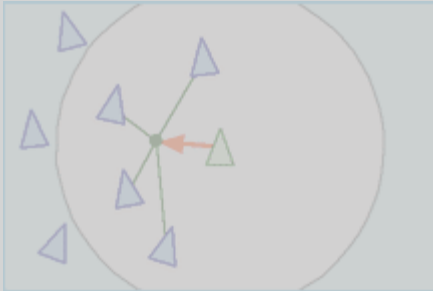
$$\vec{F}_{align} = \left(\frac{1}{n} \sum_n \vec{f}_n \right) - \vec{f}$$

separation

$$\vec{F}_{separation} = \left(\frac{1}{n} \sum_n \frac{-(\vec{p}_n - \vec{p})}{\|\vec{p}_n - \vec{p}\|^2} \right)$$

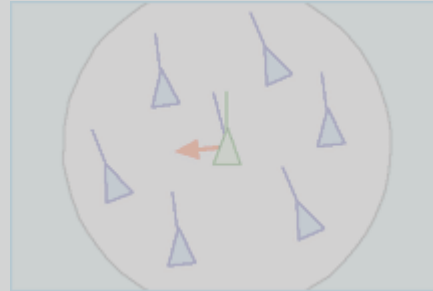


cohesion



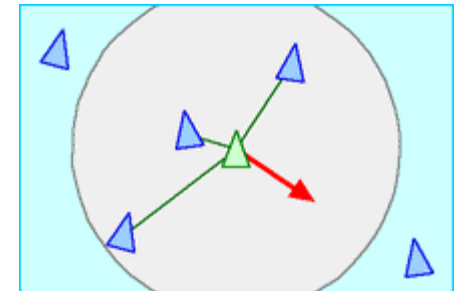
$$\vec{F}_{cohesion} = \left(\frac{1}{n} \sum_n \vec{p}_n \right) - \vec{p}$$

alignment

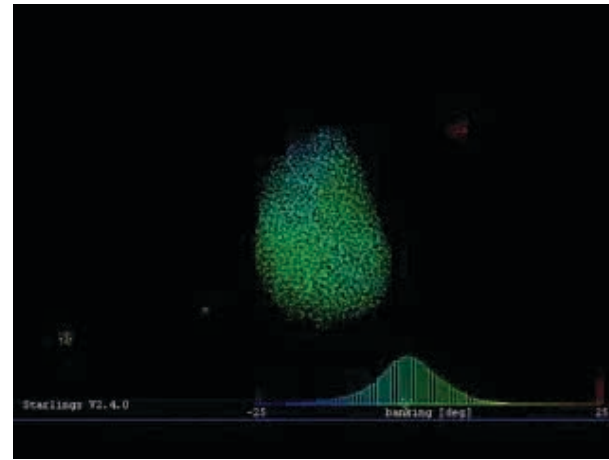
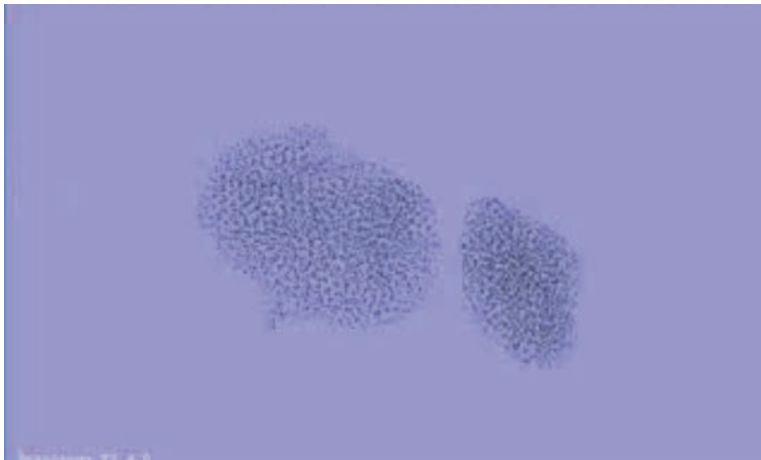


$$\vec{F}_{align} = \left(\frac{1}{n} \sum_n \vec{f}_n \right) - \vec{f}$$

separation



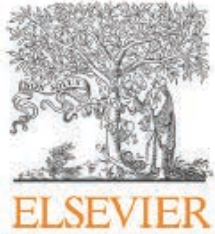
$$\vec{F}_{separation} = \left(\frac{1}{n} \sum_n \frac{-(\vec{p}_n - \vec{p})}{\|\vec{p}_n - \vec{p}\|^2} \right)$$



**screenshots of an ABM developed by Hanno Hildenbrandt
(Uni Groningen, Netherlands)**



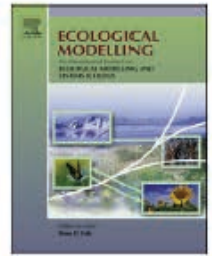
Locust Swarms - Subsahara



Contents lists available at [ScienceDirect](http://www.sciencedirect.com)

Ecological Modelling

journal homepage: www.elsevier.com/locate/ecolmodel



Self-organized spatial structures of locust groups emerging from local interaction



Jamila Dkhili^{a,b,c,*}, Uta Berger^d, Lalla Mina Idrissi Hassani^a, Saïd Ghaout^c, Ronny Peters^d, Cyril Piou^{a,b,c}

^a Faculté des sciences, Université Ibn Zohr, Agadir, Morocco

^b CIRAD, UMR CBGP, 34398 Montpellier Cedex 5, France

^c CNLAA, BP15, 86343 Inezgane, Morocco

^d TU Dresden, Faculty Of Environmental Sciences, Germany

J. Dkhili et al. / Ecological Modelling 361 (2017) 26–40

29



DeAngelis D and Zhang B:
F1000Prime Recommendation
of [Dkhili J et al., Ecol
Modell 2017, 361:26-40].

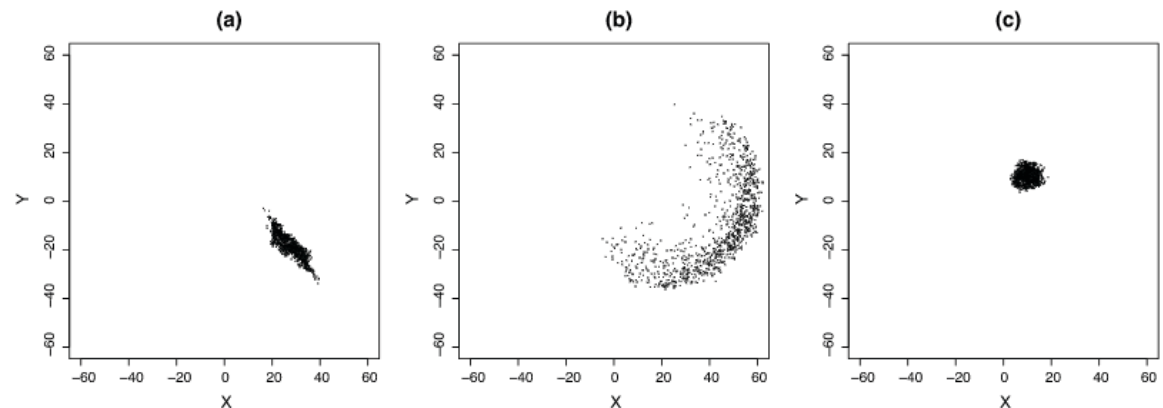


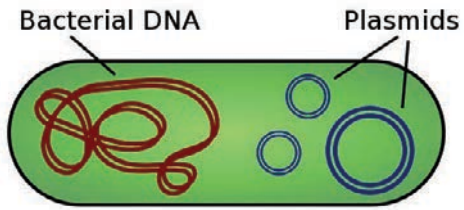
Fig. 2. Example of the three patterns generated by the model: (a) ribbons pattern; (b) band pattern; (c) spot pattern.

2nd Example

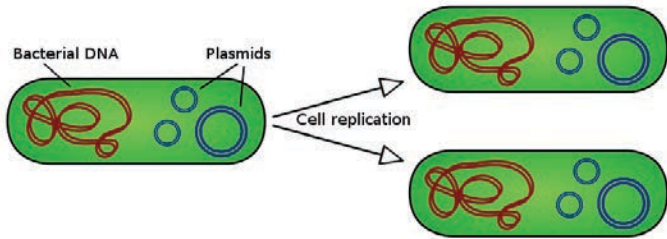
Antibiotica Resistance

(Plasmid Biology)

Basic principles of ABM: interaction & variability



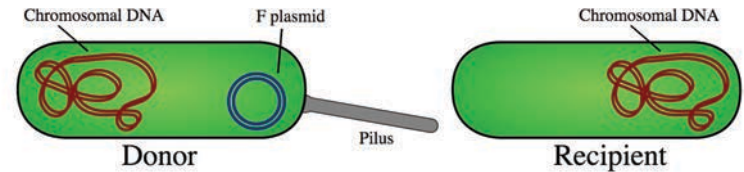
Vertical component (cell fission)
persistence from one generation
to the next



all figures: Wikimedia commons

non transmissible

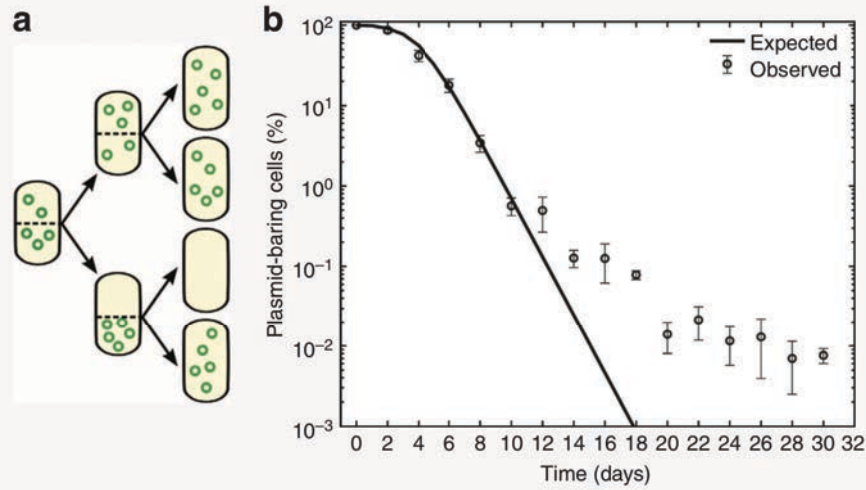
Horizontal component (conjugation)
persistence from one generation to
the next



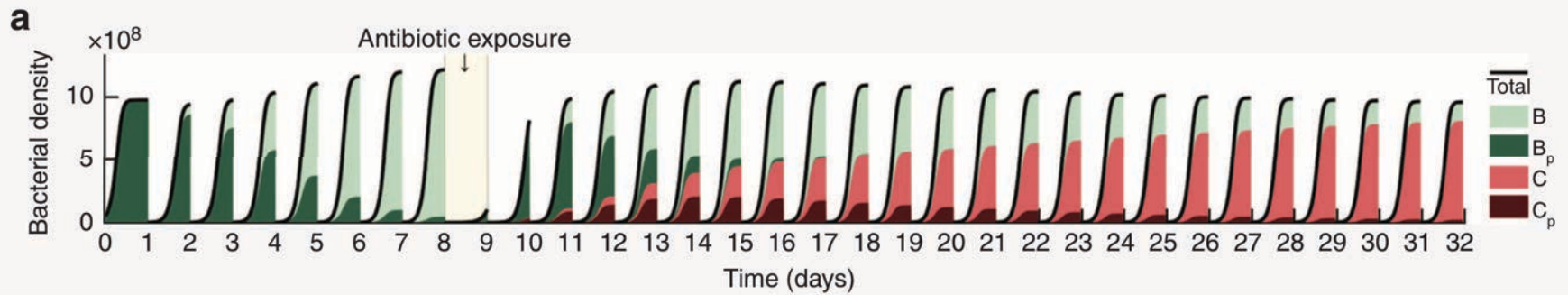
transmissible

- repressed
- derepressed

Theory: replication and plasmid burden lead to a loss without positive selection.

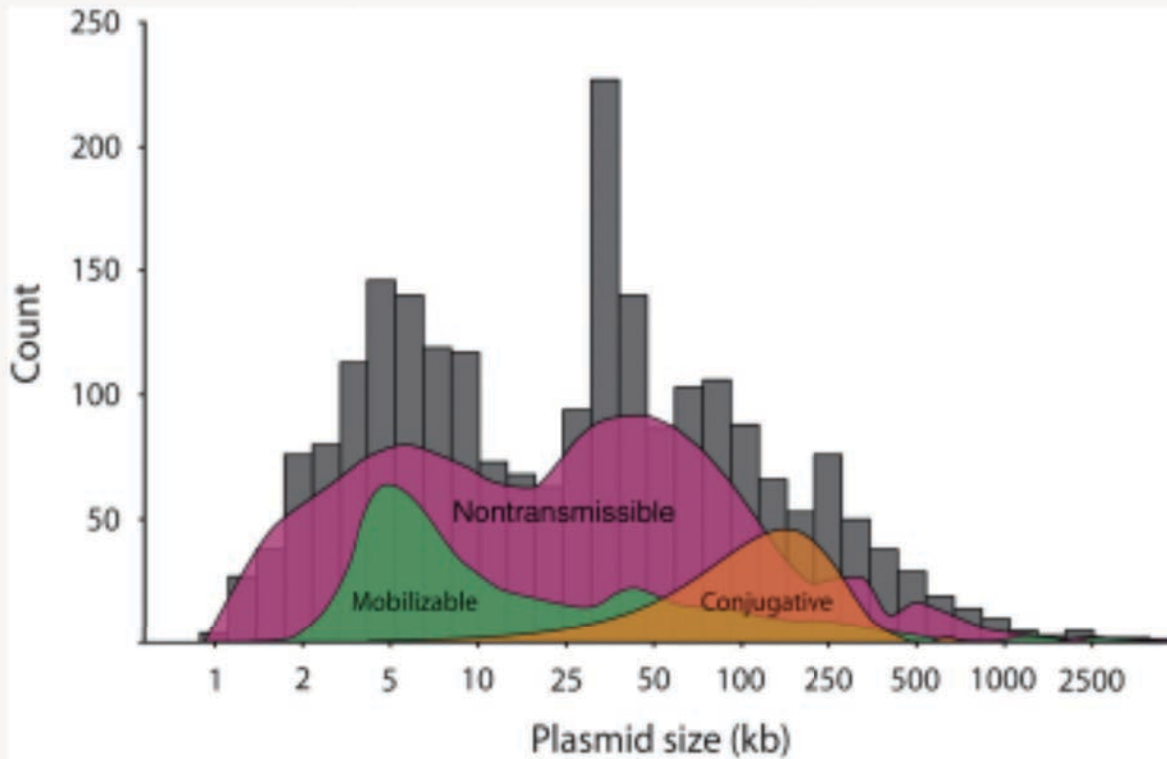


San Millan et al., Nat Commun, 2014



Reality

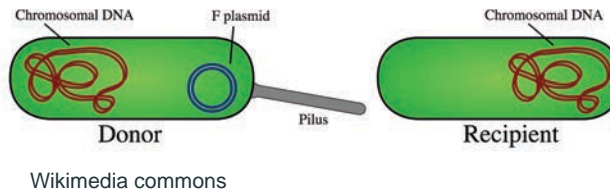
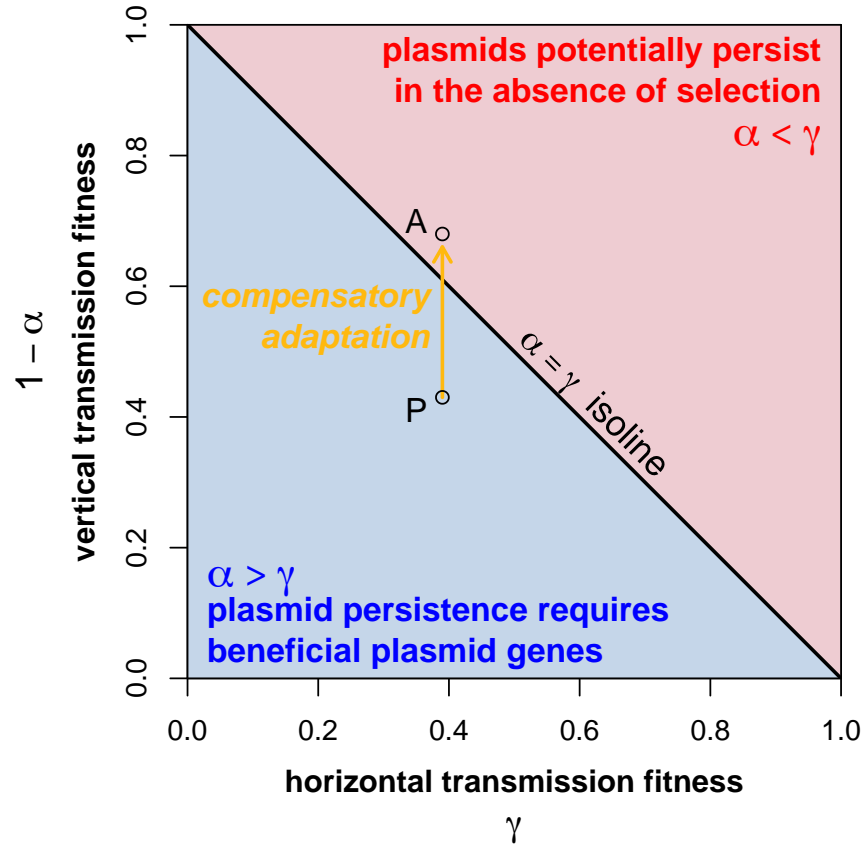
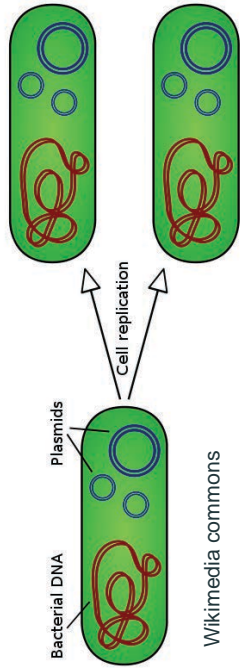
- ▶ 50% of the plasmids are non-transmissible

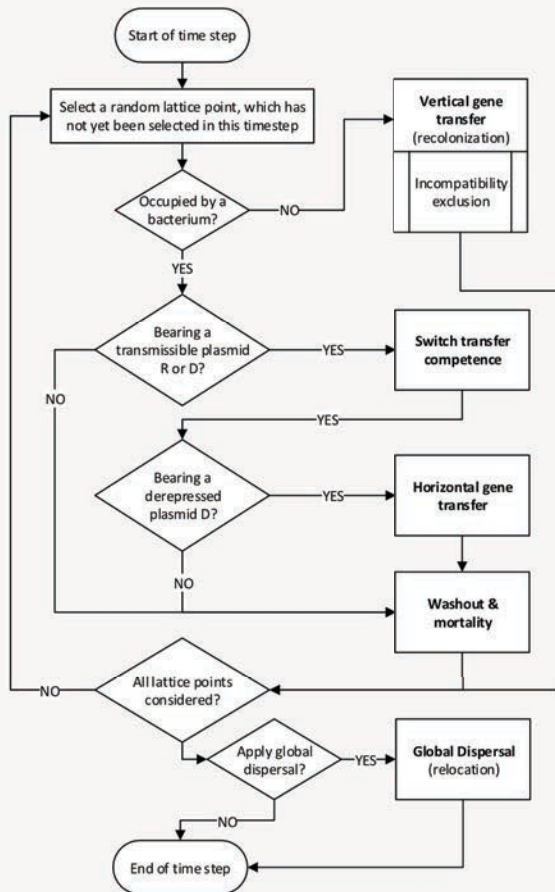


Garcillán-Barcia et al., FEMS Microbiol Rev, 2011

- ▶ What is the consequence of a co-occurrence of plasmids differing in mobility?

Concept for the persistence of plasmids without positive selection.





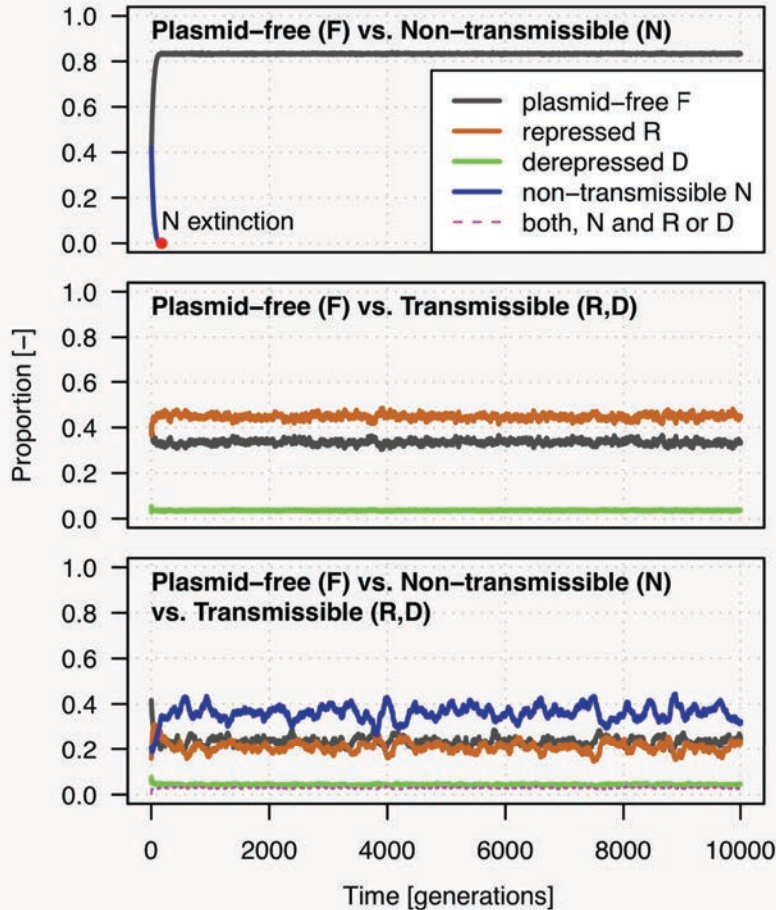
Werisch et al., Plasmid, 2017

- **ENTITIES:** BACTERIA and *PLASMIDS*

- *non-transmissible N*
- *conjugative (repressed R or derepressed D)*

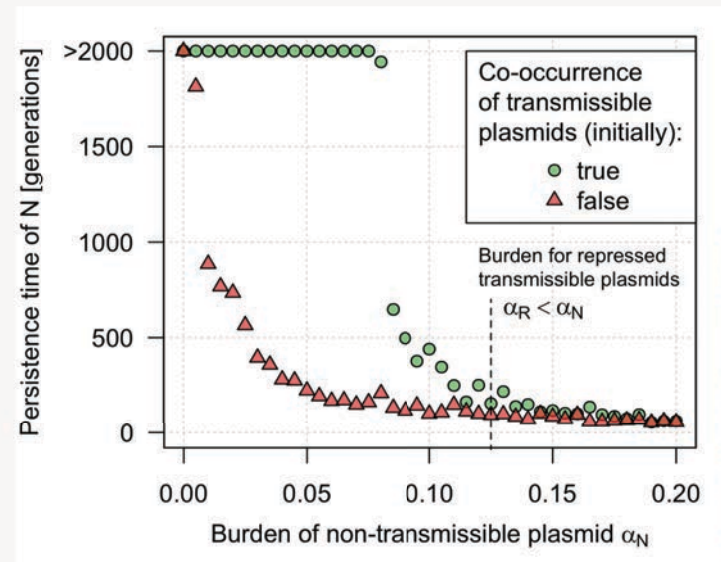
- **PARAMETERS:**

- Plasmid burden $\alpha_N, \alpha_R, \alpha_D$
- Incompatibility-exclusion probability $\kappa_N, \kappa_R D$
- Density dependency ρ_R, ρ_D for switch of transfer competence
- Transfer attempt success probability γ
- Death or washout probability ω



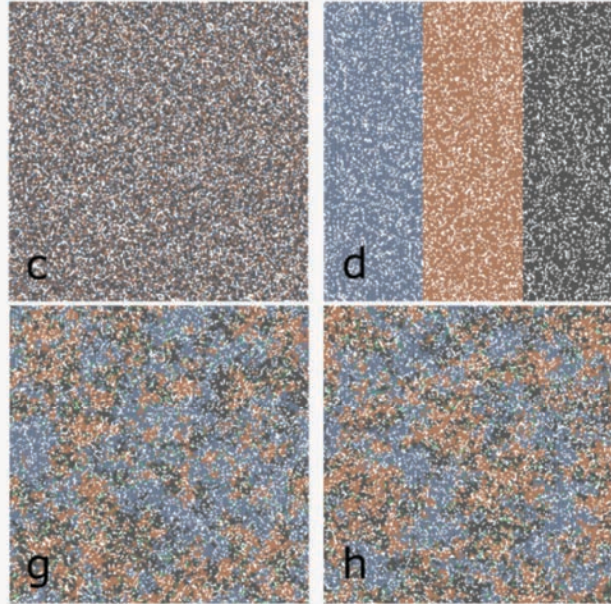
Werisch et al., Plasmid, 2017

► non-transmissible plasmids can be maintained indefinitely in the presence of an incompatible and more costly conjugative plasmid type



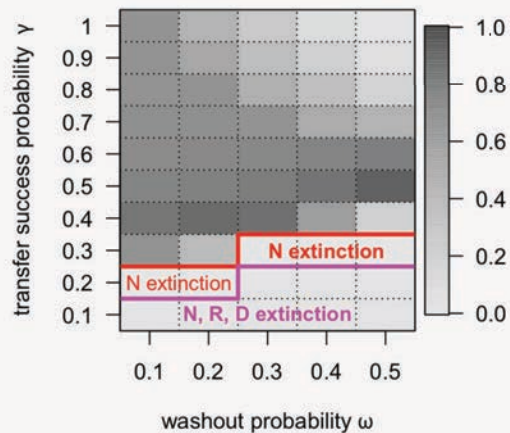
Werisch et al., Plasmid, 2017

► initial plasmid distribution

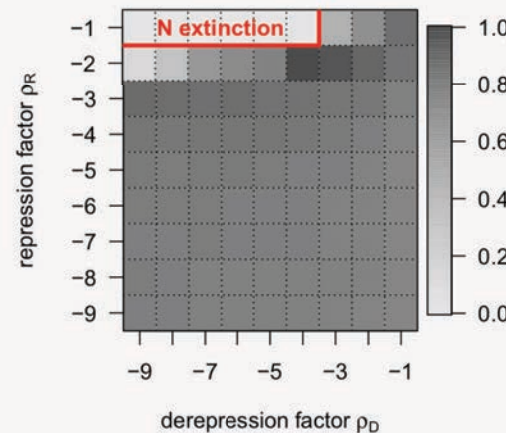


► model parameter settings

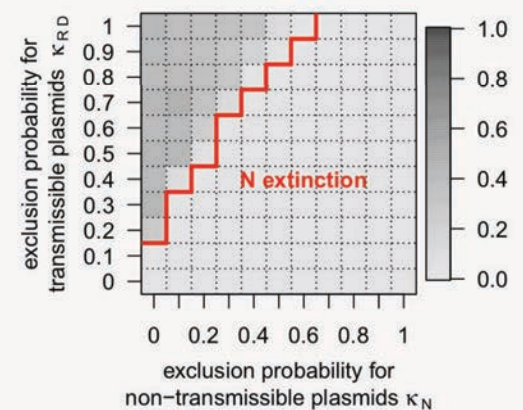
(a) Proportion of bacteria bearing non-transmissible plasmids



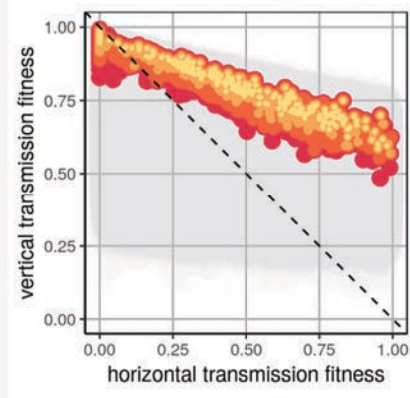
(b) Proportion of bacteria bearing non-transmissible plasmids



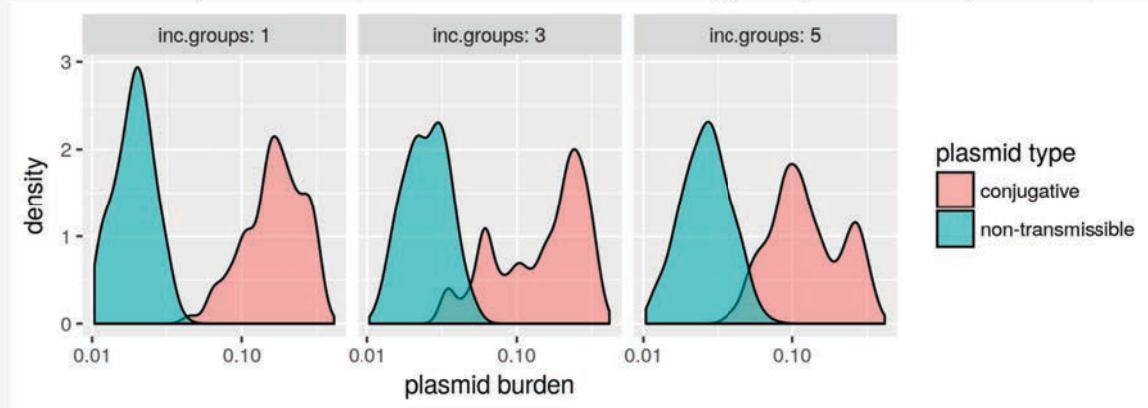
(c) Proportion of bacteria bearing non-transmissible plasmids



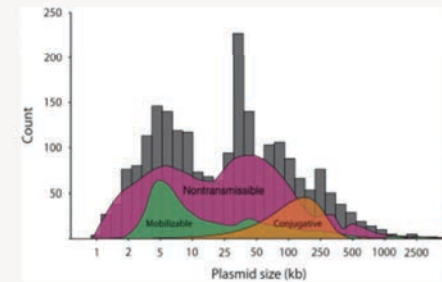
selected plasmid types in 100 simulations (inc. groups: 1)



distribution of plasmid burden of evolved community (100 simulations for each inc. group number)



► Inventory of artificial system fits basic features of natural systems (no selection for accessory-genes of plasmids considered!)



Garcillán-Barcía et al., 2011

“WE ARE NOW
SEEING A REAL
CONVERGENCE
OF WHAT
EXPERIMENTALISTS
WANT AND WHAT
THEORISTS
CAN DELIVER.”

Genetic Inspiration: Human Genome Project (MIT, early 2000)

Computer Science: Bioinformatics, Data Mining, Simulation Experiments (last decade)

Material Science: currently

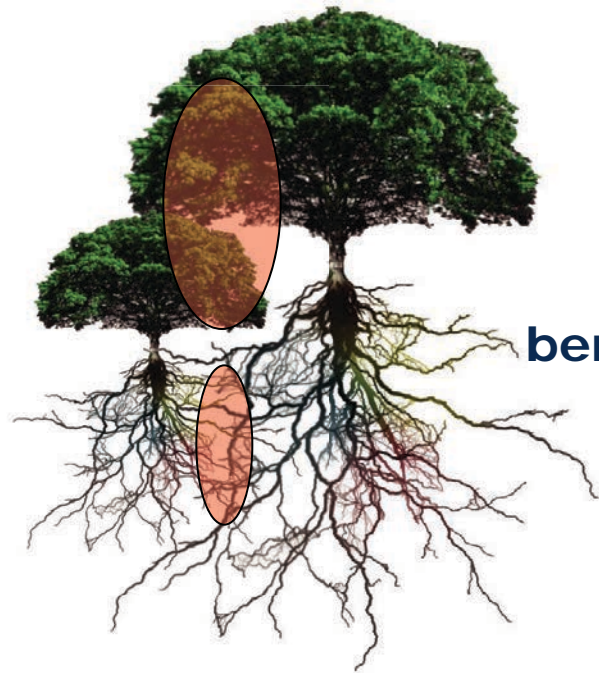
3rd Example

Plant Ecology

ABM meets Machine Learning

Competition

Asymmetric



benign \longleftrightarrow stress gradient \longleftrightarrow harsh

Symmetric

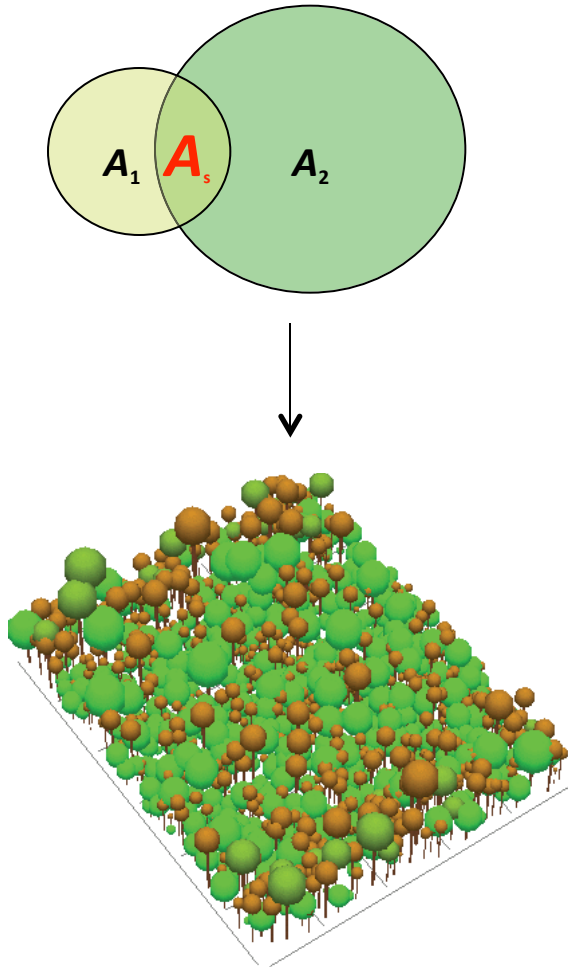
Facilitation

Asymmetric



Symmetric

Plant-Interaction (PI) Modell



$$dm/dt = f_p f_q c A [1 - (m / M)^{1/4}]$$

m .. Current biomass

M .. Maximum biomass

C .. Initial growth rate

A .. Area within the plant exploits resources

f_p .. competition among neighbouring plants

$$f_p = (A_{no} + \sum_{k=1}^{n_o} \frac{v_i m_i^p}{\sum_{j=1}^{n_j} v_j m_j^p} A_{o,k}) / A$$

f_q ... facilitation among neighbouring plants (in case of stress)

$$f_q = 1 - \frac{S}{A_f + 1} = 1 - \frac{S}{\sum_{k=1}^{n_o} (1 - \frac{w_i m_i^q}{\sum_{j=1}^{n_j} w_j m_j^q}) A_{o,k} + 1}$$

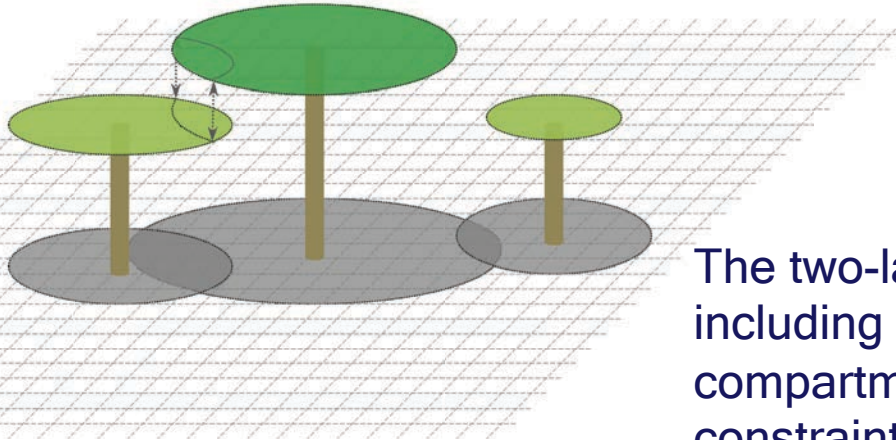
A_{no} area of plant i without overlap

$A_{o,k}$ overlapping area with neighbouring plant j

p, q parameter of symmetry – asymmetry

v, w species-specific weight

S stress level



The two-layer version of the PI model including both above- and belowground compartments connected via metabolic constraints.

$$\frac{\Delta m}{\Delta t} = \begin{cases} 2 \times \Delta AGR = 2 \times f_{R,a} f_{p,a} c_a A_a [1 - (m / M_a)^{1/4}], & \Delta AGR < \Delta BGR \\ 2 \times \Delta BGR = 2 \times f_{R,b} f_{p,b} c_b A_b [1 - (m / M_b)^{1/4}], & \Delta AGR > \Delta BGR \\ \Delta AGR + \Delta BGR, & \Delta AGR = \Delta BGR \end{cases}$$

$\Delta AGR, \Delta BGR$.. above- and belowground growth rate
 a, b .. above- and belowground compartment

Plant-Interaction (PI) Modell

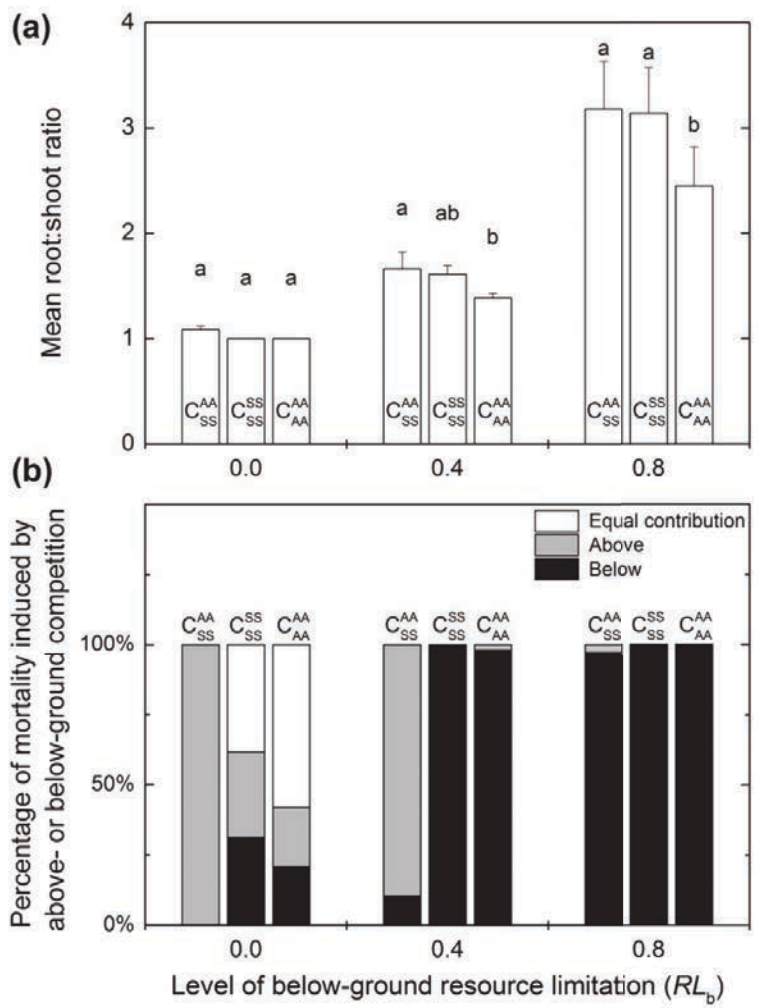
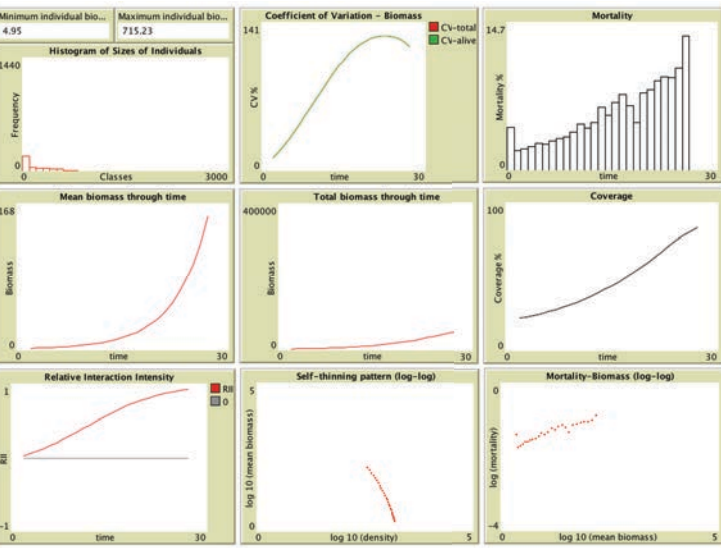
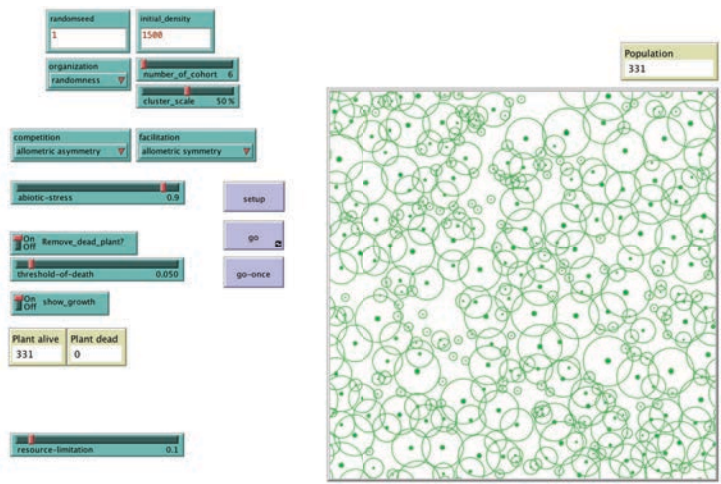
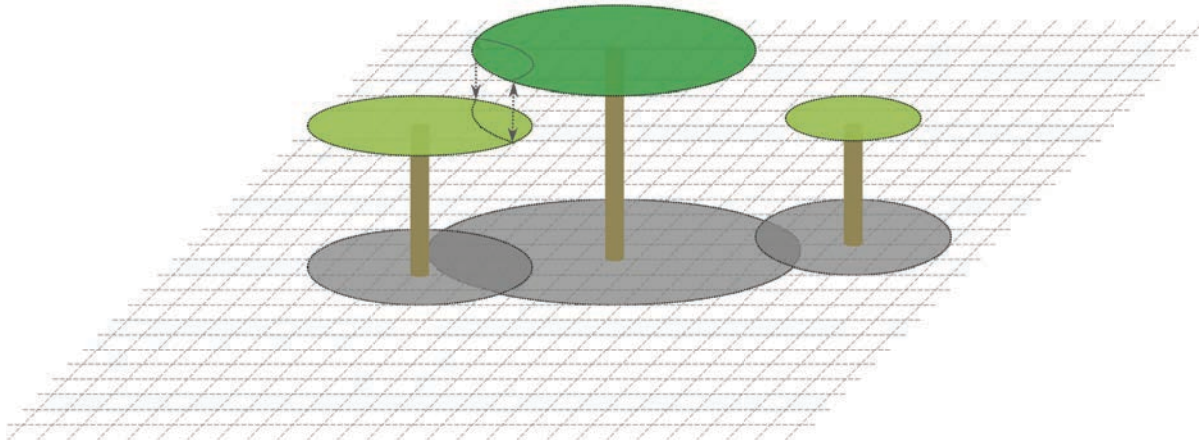


Figure 1. The (a) root:shoot ratio (mean \pm 1 SD) and (b) percentage of mortality induced by above- or belowground competition in simulated plant populations at different levels of belowground resource limitation, RL_b , ranging from 0 (no resource limitation) to 1 (maximum resource limitation). C with superscript and subscript indicated the mode of competition for above- and belowground part correspondingly (AA: allometric asymmetry; SS: size symmetry). Bars in the same group that share the same letter do not differ significantly (p > 0.05, Holm-Sidak test).

”I have a dream ...”

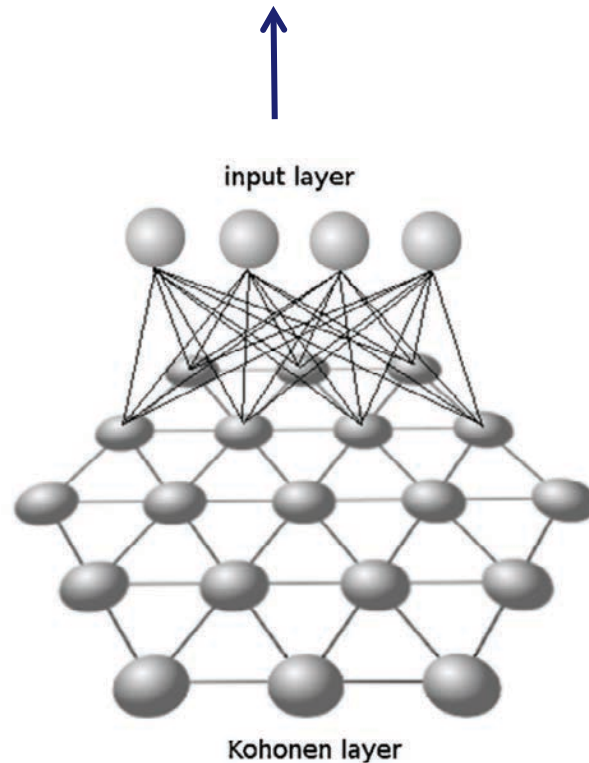
Observations above-ground Dynamic (Mortality, Spatial distribution ...)



.. can be used to reveal mechanisms below-ground (competition mode, resource limitation ..)

Dynamic observations at stand level:

- Mortality
- Spatial distribution (Clark Evans Index)



The self-organizing map (SOM = Kohonen layer) links the factors without defining dependencies.

Mechanisms to be detected:

- Below-ground competition mode
- Resource limitation

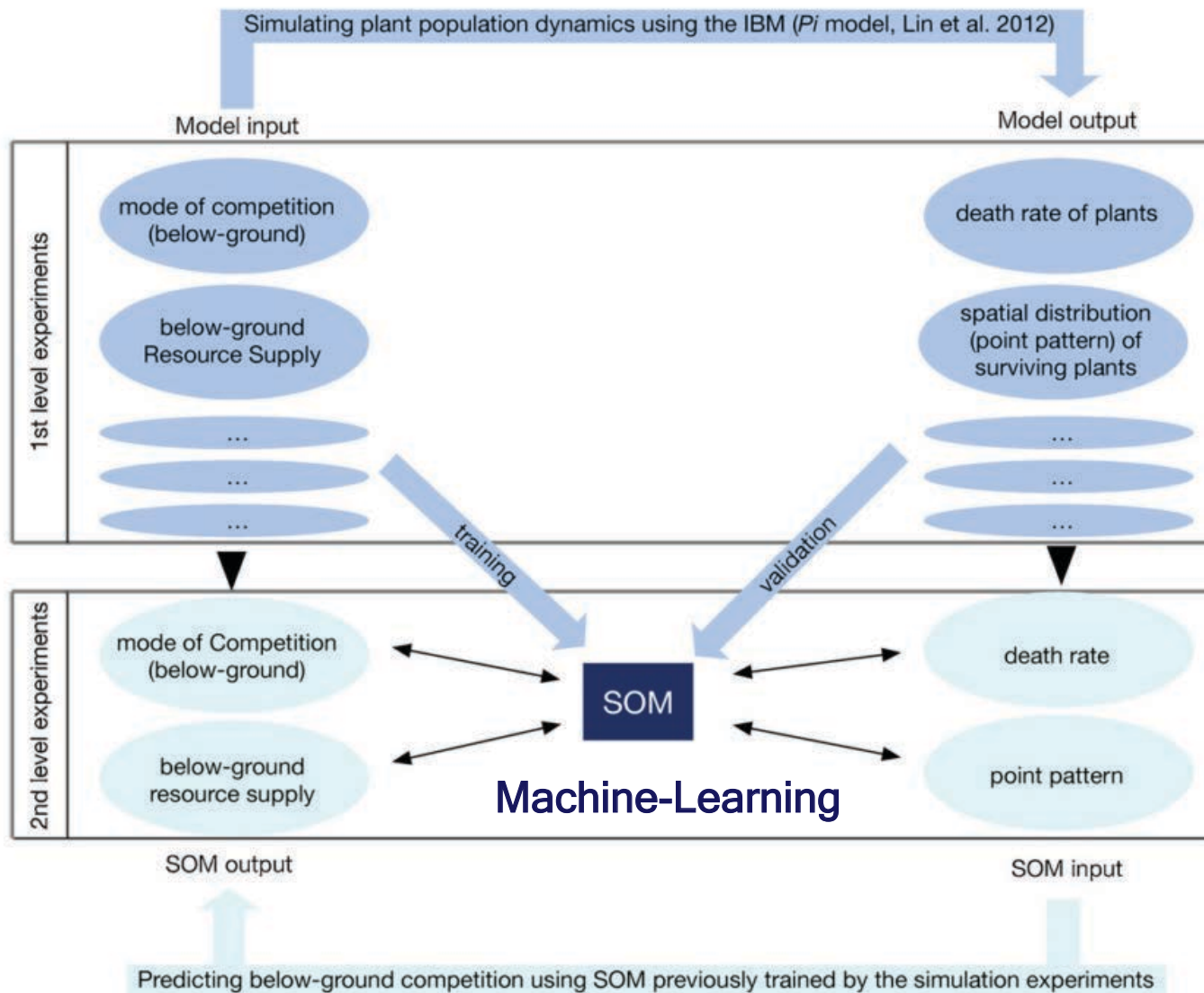


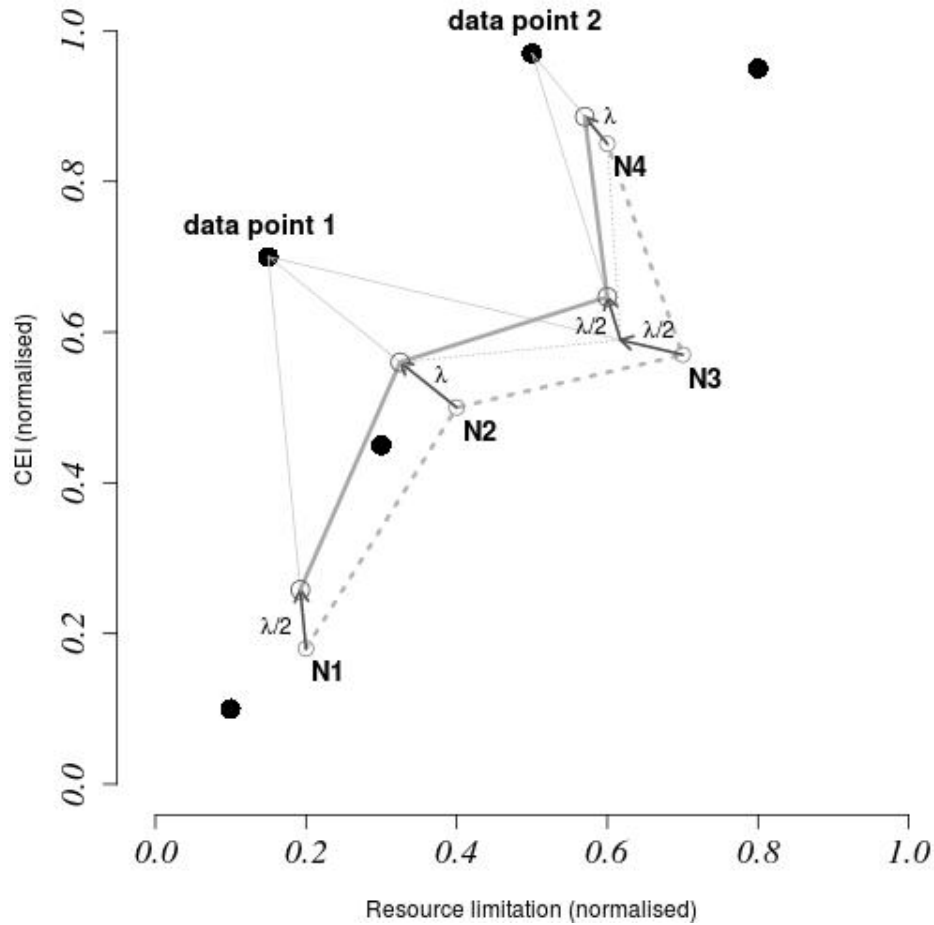
Fig. 1. Strategy of the IBM-ANN approach applied in this study. The figure shows the relationship between the input and output variables of the individual-based simulation model (*Pi* model, above) and their further use by self-organising maps (SOM).

Table 1

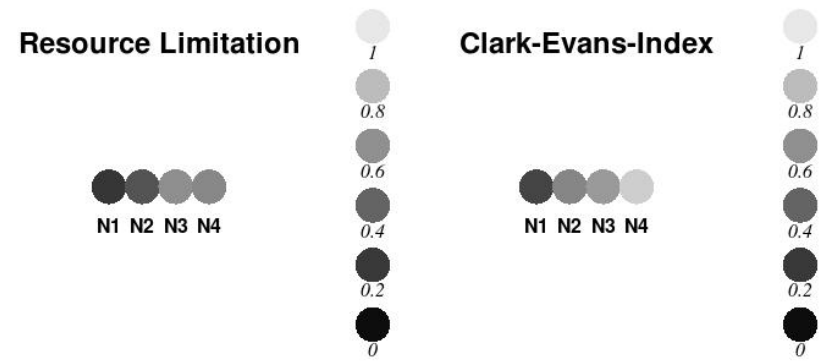
The parameter settings used for data generation by means of the individual-based *Pi* model.

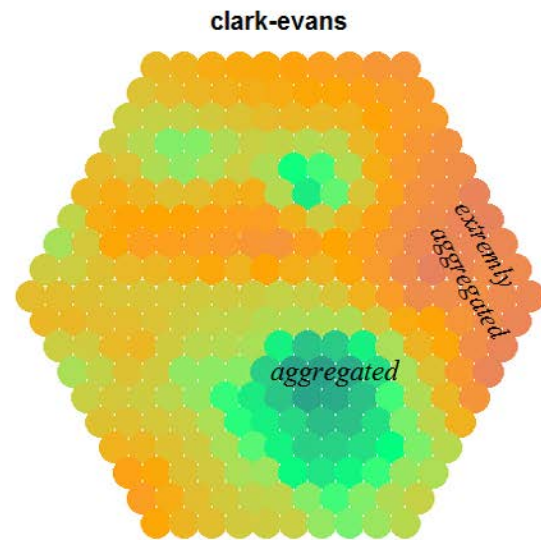
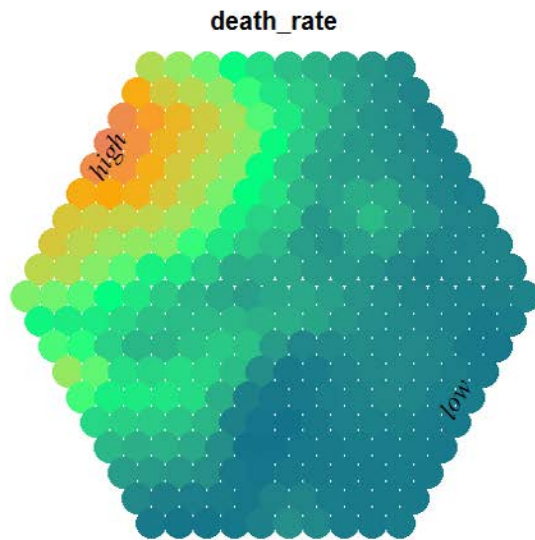
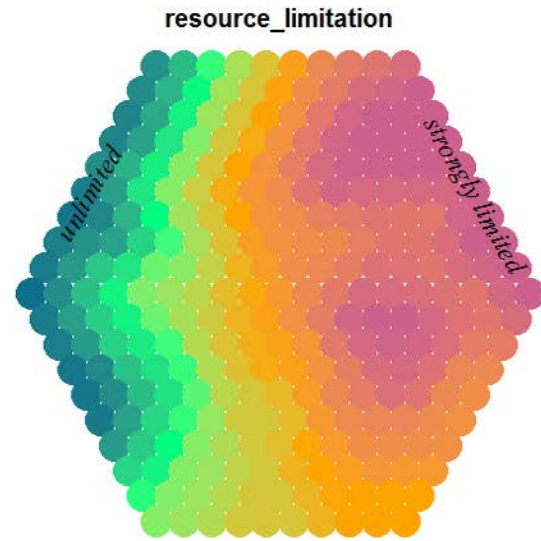
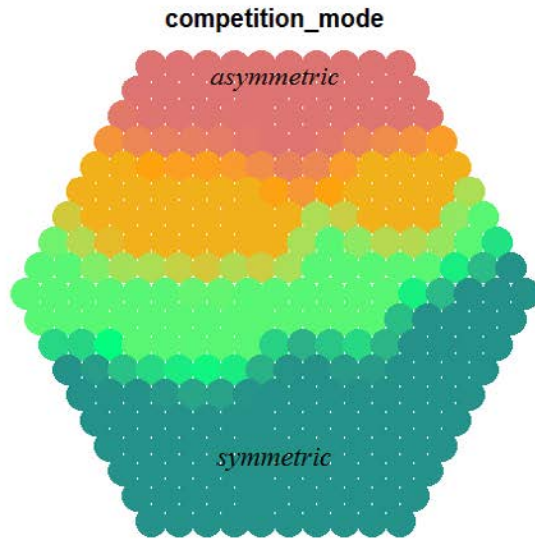
<i>Pi</i> model parameter setting	Training data	Validation data
Above-ground competition mode	Complete asymmetry	Complete asymmetry
Below-ground competition mode	Complete asymmetry	Complete asymmetry
	Allometric asymmetry	Allometric asymmetry
	Size symmetry	Size symmetry
	Complete symmetry	Complete symmetry
Initial individual number	7000	7000
	8000	8000
	9000	9000
Resource limitation	0.0 0.1 ... 0.9	0.05 0.15 ... 0.85
Total number of model runs	120	108

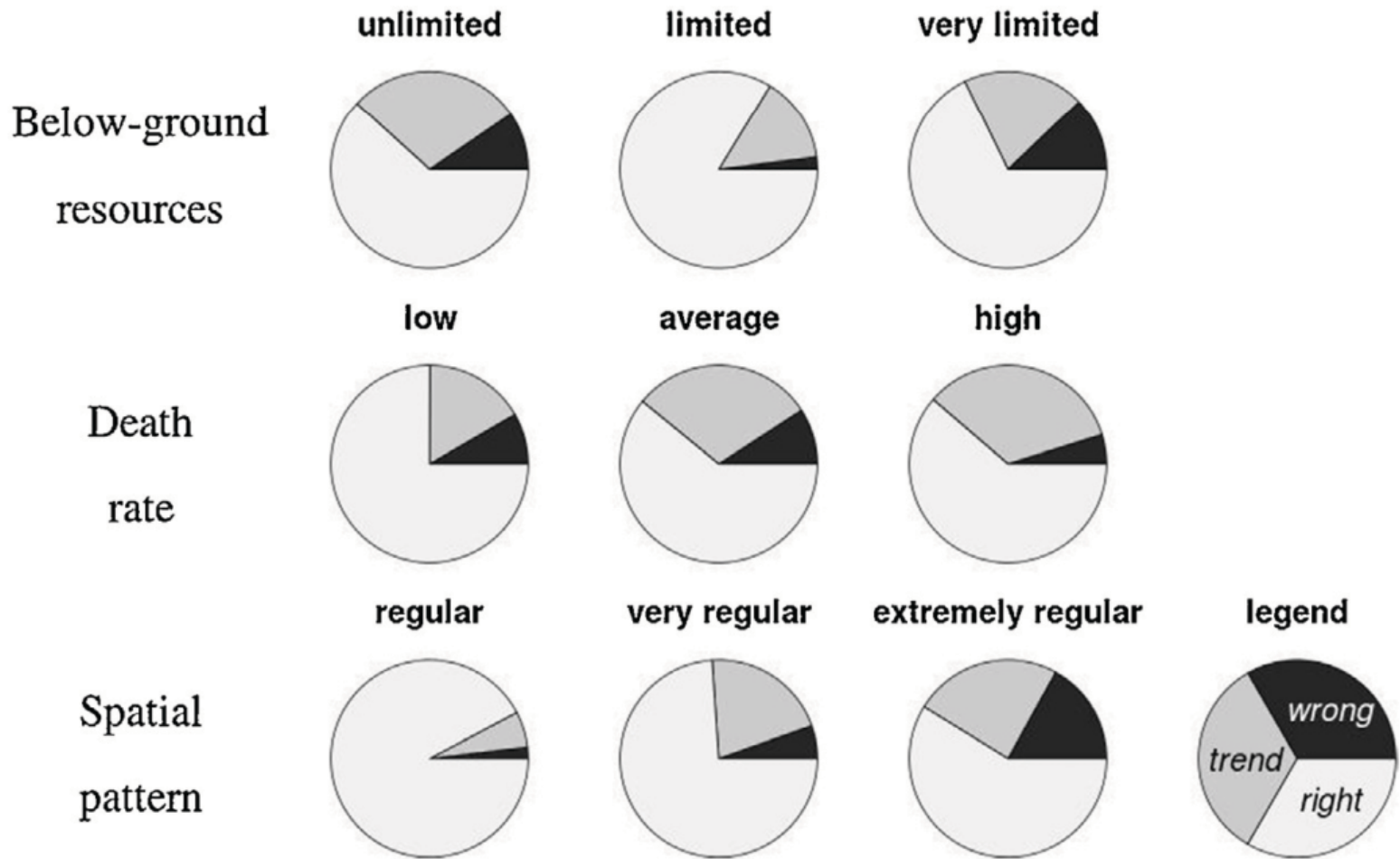
SOM training



SOM layer after training

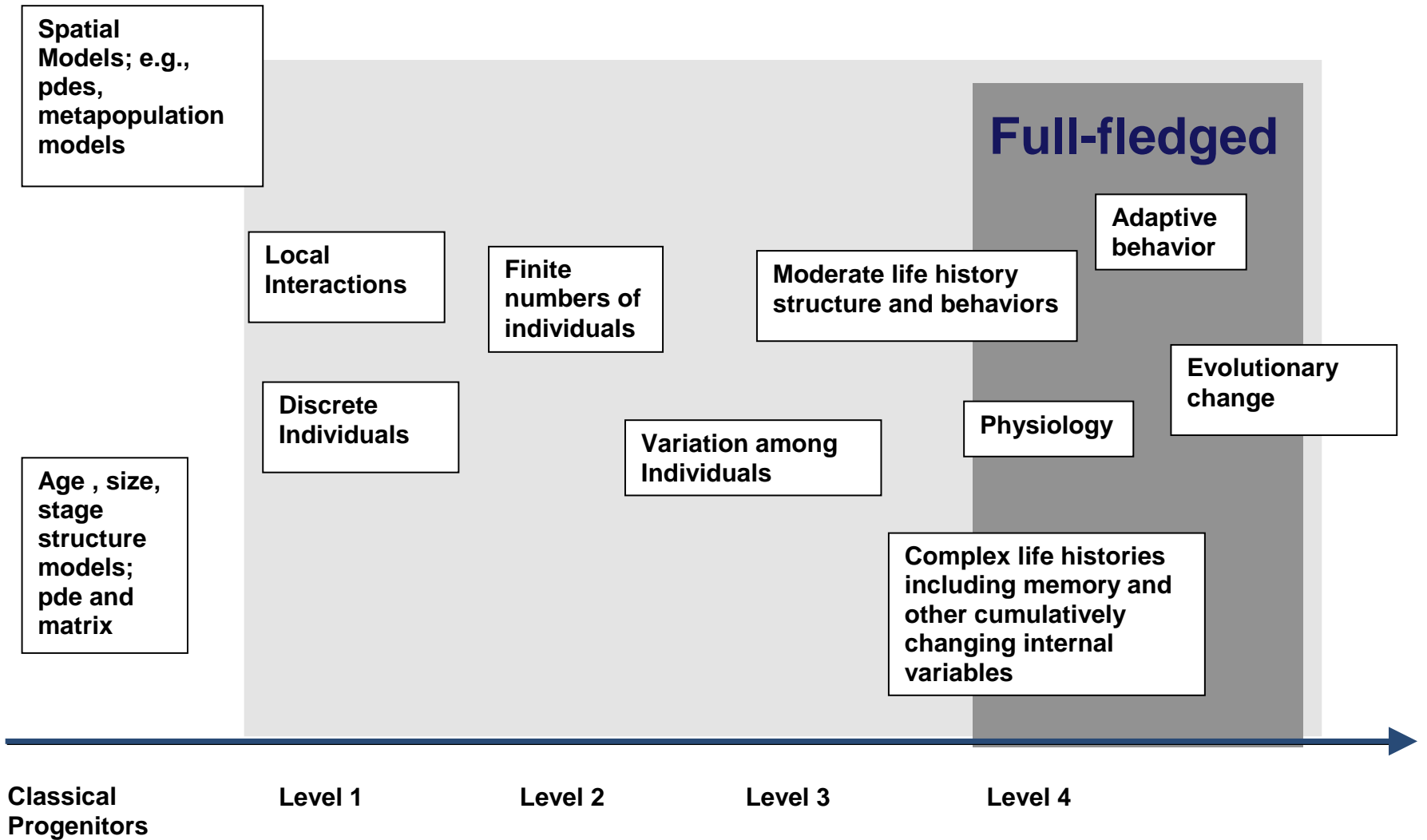


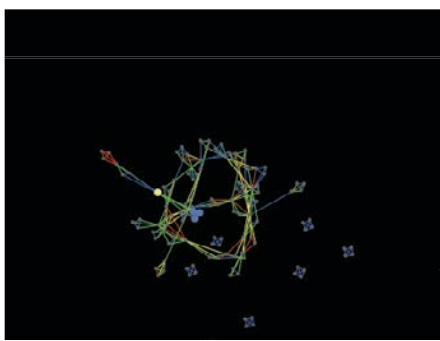




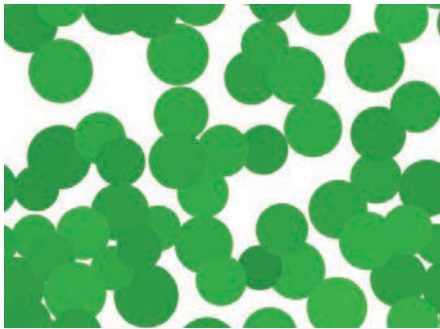
IBM / ABM

towards agent-based science?

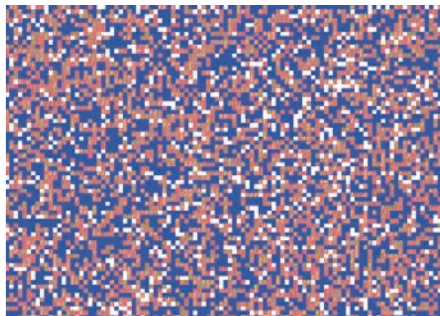




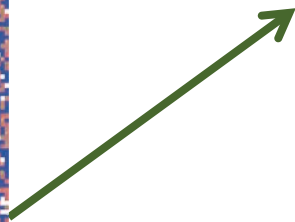
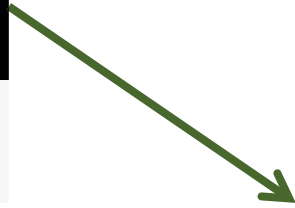
agent-based models
(social sciences)



stand simulators
(forest sciences)



indiv.-based models
(ecology)



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Research



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How new concepts become universal scientific approaches: insights from citation network analysis of agent-based complex systems science

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Progress in understanding and managing complex systems comprised of decision-making agents, such as cells, organisms, ecosystems or societies, is—like many scientific endeavours—limited by disciplinary boundaries. These boundaries, however, are porous and can be gradually eroded.

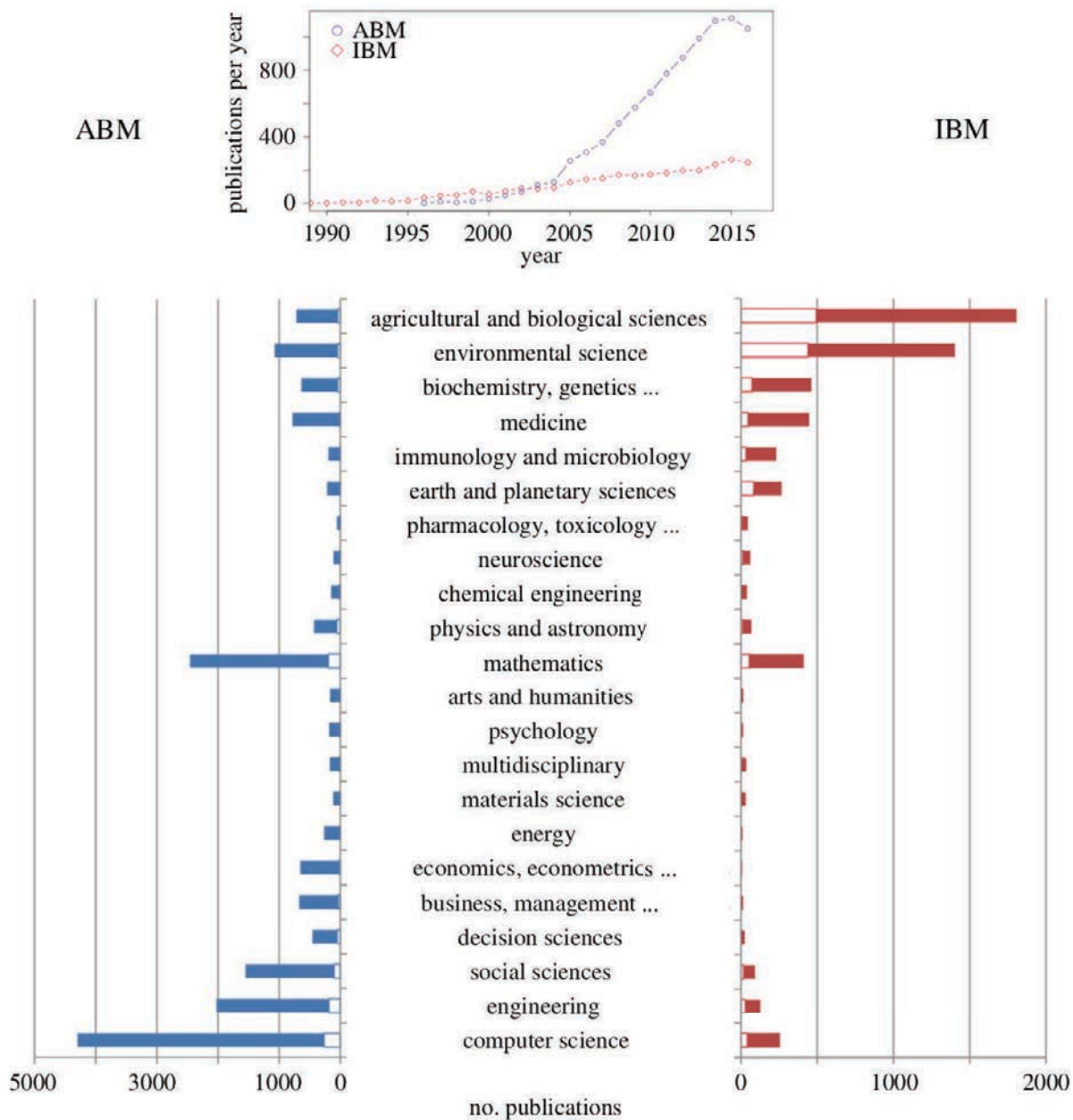


Figure 1. Sciences in which the terms ABM (left) and IBM (right) were dominant until 2005 (empty bars) and post-2005 (solid bars) measured in number of publications referenced on Scopus ($n = 9515$ and 3168 , respectively). Publications can belong to more than one scientific domain. The top inset shows the number of publications per year related to ABM and IBM from 1989 to 2016. Publications belonging to both corpora were counted twice. (Online version in colour.)

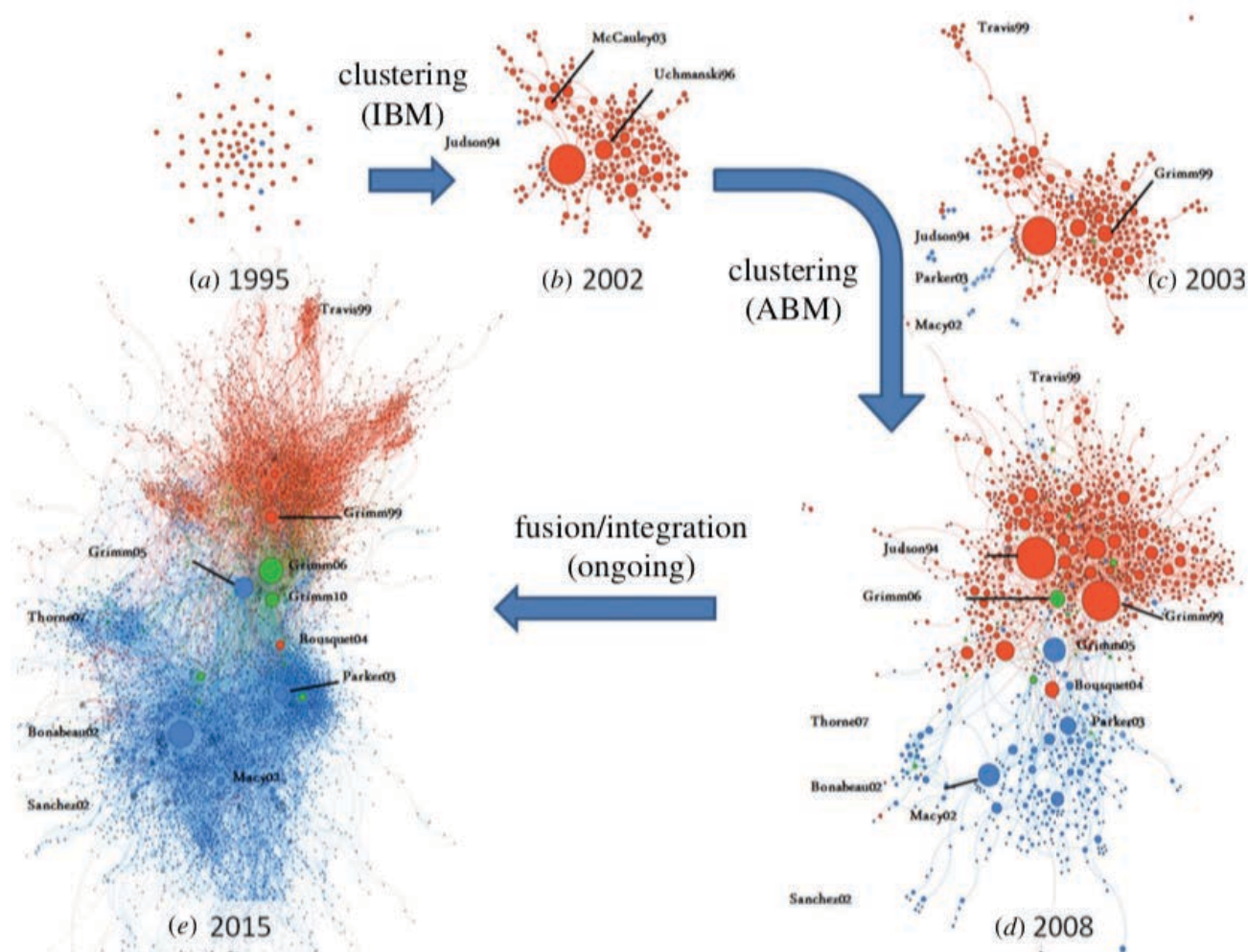


Figure 4. Evolution of the citation graph of ABM/IBM literature over time. Publications are represented by vertices (blue for ABM, red for IBM and green for both), which are attracted to each other proportionally to the number of edges (i.e. citations) linking them. Important publications are labelled with first author name and publication year. For full citations, see references and the electronic supplementary material, table S3. Vertex size is proportional to the number of citations within each time-dependent subgraph, but scaling varies between subgraphs. (a) IBM papers appeared first, but had weak interconnections. (b) A clustering then took place after the appearance of influential papers. (c) ABM papers followed the same pattern with a lag of a decade. (d) Clustering of the ABM corpus occurred around central papers. (e) Fusion of ABM and IBM emerged under the impulse of key papers. The formation of sub-clusters is also visible; for example, publications centred around Thorne *et al.* [22] focused on cell-scale agent-based models. (Online version in colour.)

Conclusions

- tendency to full-fledged models (mechanistic, physiology based)
- IBM / ABM toolbox (standardization, ODD, TRACE)
- merging with formerly separated modelling techniques
- ready to go towards agent-based sciences

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- Ronny Peters (TU Dresden, Germany)
- Hanno Hildenbrandt (University of Groningen, The Netherlands)
- Christian Vincenot (Kyoto University, Japan)



Research Projects

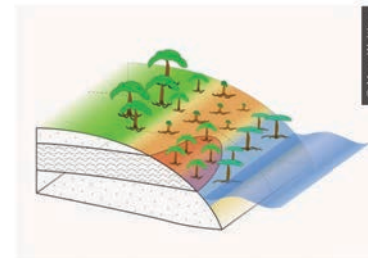
Our chair executes a variety of research projects. Topics, contact persons and funding agencies are listed below.

Index

[>current projects](#) | [>completed projects](#)

CURRENT PROJECTS

MARZIPAN - Adopting mangrove vegetation zonation patterns to gain information on subsurface aquifer structures and advance below-ground plant competition concepts in individual-based modelling



- Since 2018
- Funded by German Research Foundation (DFG)
- Contact person: [>Ronny Peters](#)

RESCuE - Monitoring and REstoration for Sustainable Coastal Ecosystems



- Since 2018
- Funded by SOUTHEAST ASIA-EUROPE JOINT FUNDING SCHEME FOR RESEARCH AND INNOVATION
- Contact person: [>Uta Berger](#)