



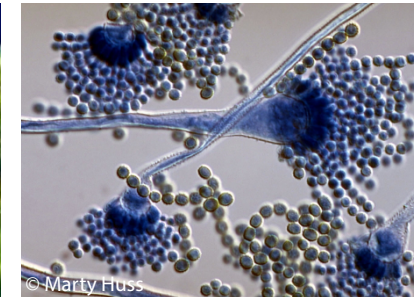
Some news on the population genetics of partial asexuals

PhD Thesis at INRA Rennes



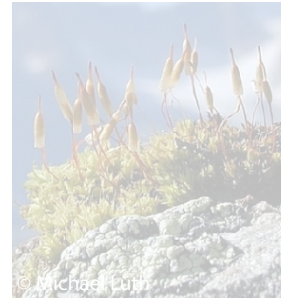
Organisms

examples for different rates of asexuality

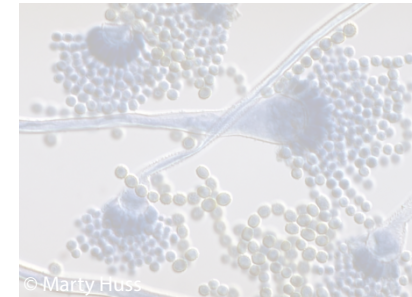
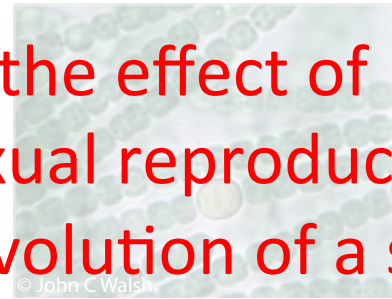
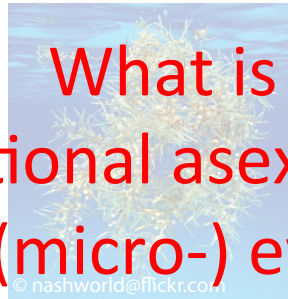


Organisms

examples for different rates of asexuality



What is the effect of optional asexual reproduction on the (micro-) evolution of a species?

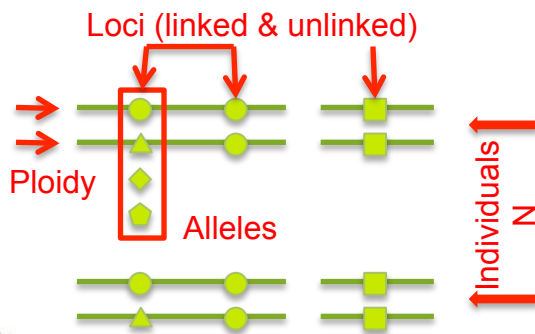
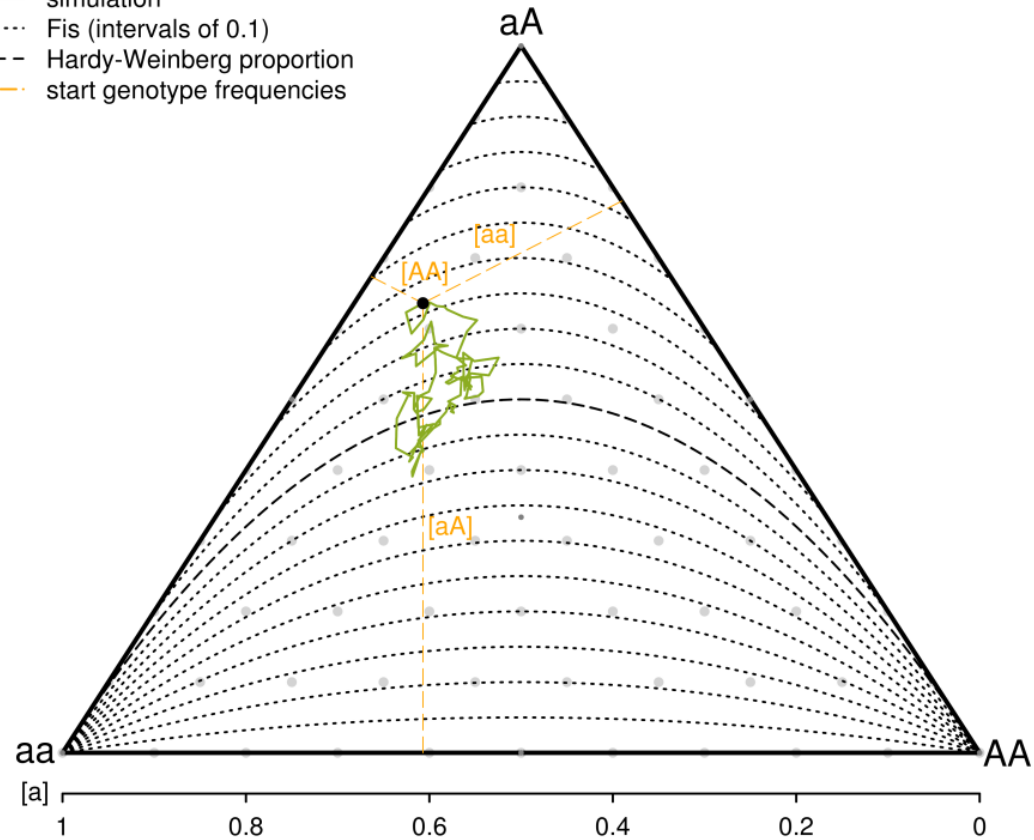


Bruno de Finetti

and his diagrams



- start point
- simulation
- ⋯ Fis (intervals of 0.1)
- - Hardy-Weinberg proportion
- - start genotype frequencies



Markov Chains

the principal idea



- mathematically: a (recursive) sequence
 - an example:
 - $a_1 = 1$ ⇒ generation 1 (first observation)
 - $a_{n+1} = a_n + 3$ ⇒ reproduction “rule”
 - gives {1; 4; 7; 10; 13; 16; 19; 22; 25; 28; 31; 34; 37; ...}
 - this series converges to ∞ with a constant positive “speed”
 - none of its members is divisible by 3, but every second is divisible by 2, and (with a skip) every fourth by 4, every fifth by 5, every seventh by 7 ...
- to incorporate partial asexuality, we need to change the rules for reproduction!

Some more detail

Model equations




μ ... mutation rate
 c ... asexuality rate

	do not mutate	mutate	
<i>asexual</i>	$p^c(aa) = (1-\mu)^2 * p(aa)$ $p^c(AA) = (1-\mu)^2 * p(AA)$ $p^c(aA) = ((1-\mu)^2 + \mu^2) * p(aA)$	$+ \mu (1-\mu) * p(aA) \quad + \mu^2 * p(AA)$ $+ \mu (1-\mu) * p(aA) \quad + \mu^2 * p(aa)$ $+ 2\mu (1-\mu) * p(aa) \quad + 2\mu (1-\mu) * p(AA)$	
<i>sexual</i>	$p^s(aa) = ((1-\mu) * p(aa) + \mu * p(AA) + 0.5 * p(aA))^2$ $p^s(AA) = ((1-\mu) * p(AA) + \mu * p(aa) + 0.5 * p(aA))^2$ $p^s(aA) = 2 * (((1-\mu) * p(aa) + \mu * p(AA) + 0.5 * p(aA)) + ((1-\mu) * p(AA) + \mu * p(aa) + 0.5 * p(aA)))$	<p style="text-align: right;">a A a A</p>	which gametes are produced?
<i>total</i>	$p'(aa) = c * p^c(aa) + (1-c) * (p^s(aa) + 0.5 * p^s(aA))^2$ $p'(AA) = c * p^c(AA) + (1-c) * (p^s(AA) + 0.5 * p^s(aA))^2$ $p'(aA) = c * p^c(aA) + 2 * (1-c) * (p^s(aa) + 0.5 * p^s(aA)) * (p^s(AA) + 0.5 * p^s(aA))$	<p style="text-align: right;"><i>asexual</i> <i>sexual</i></p>	which gametes combine?

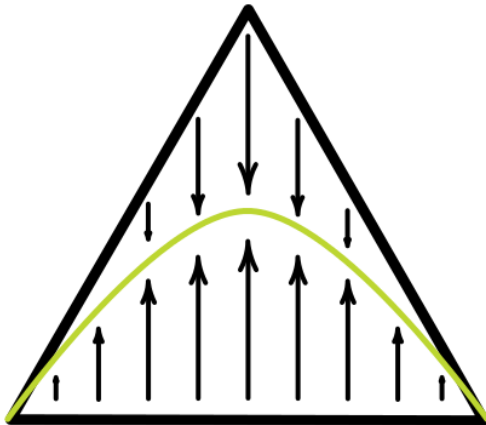
Parameter effects

Explaining what we see

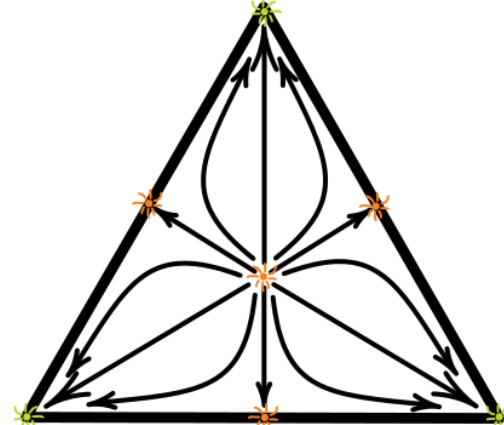
Equilibria:

-  attractive + stable
-  attractive + instable
-  not attractive + instable

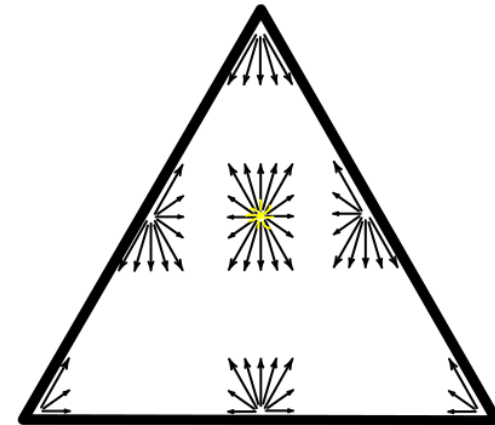
Recombination



Drift

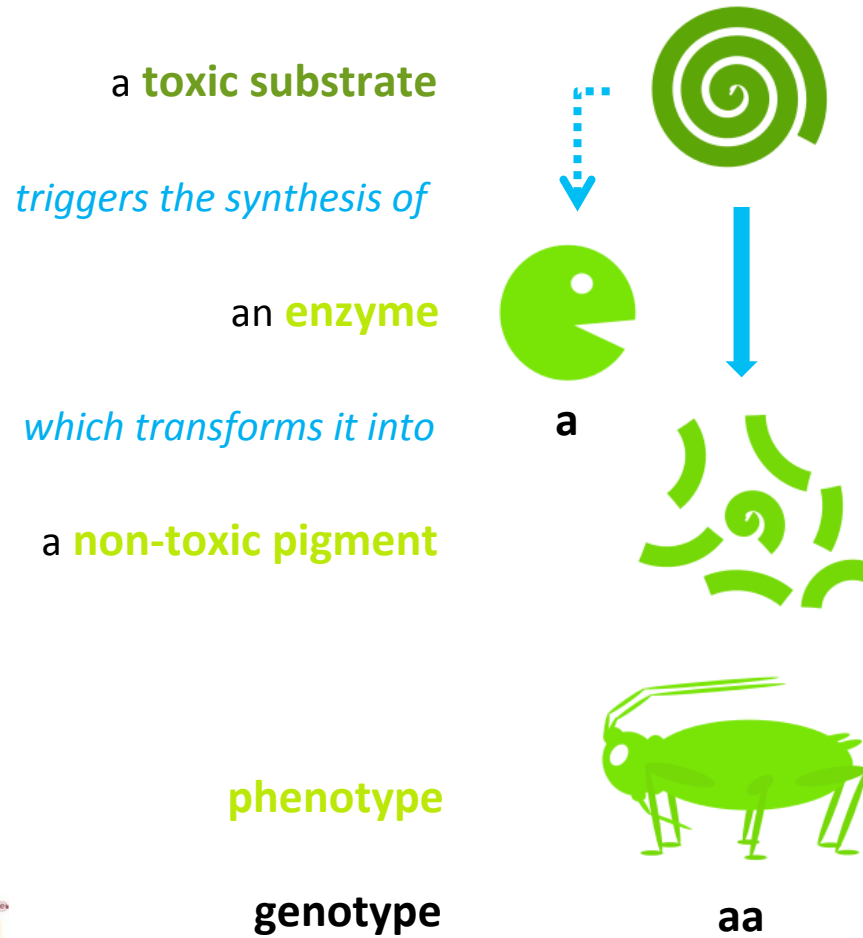


Mutation



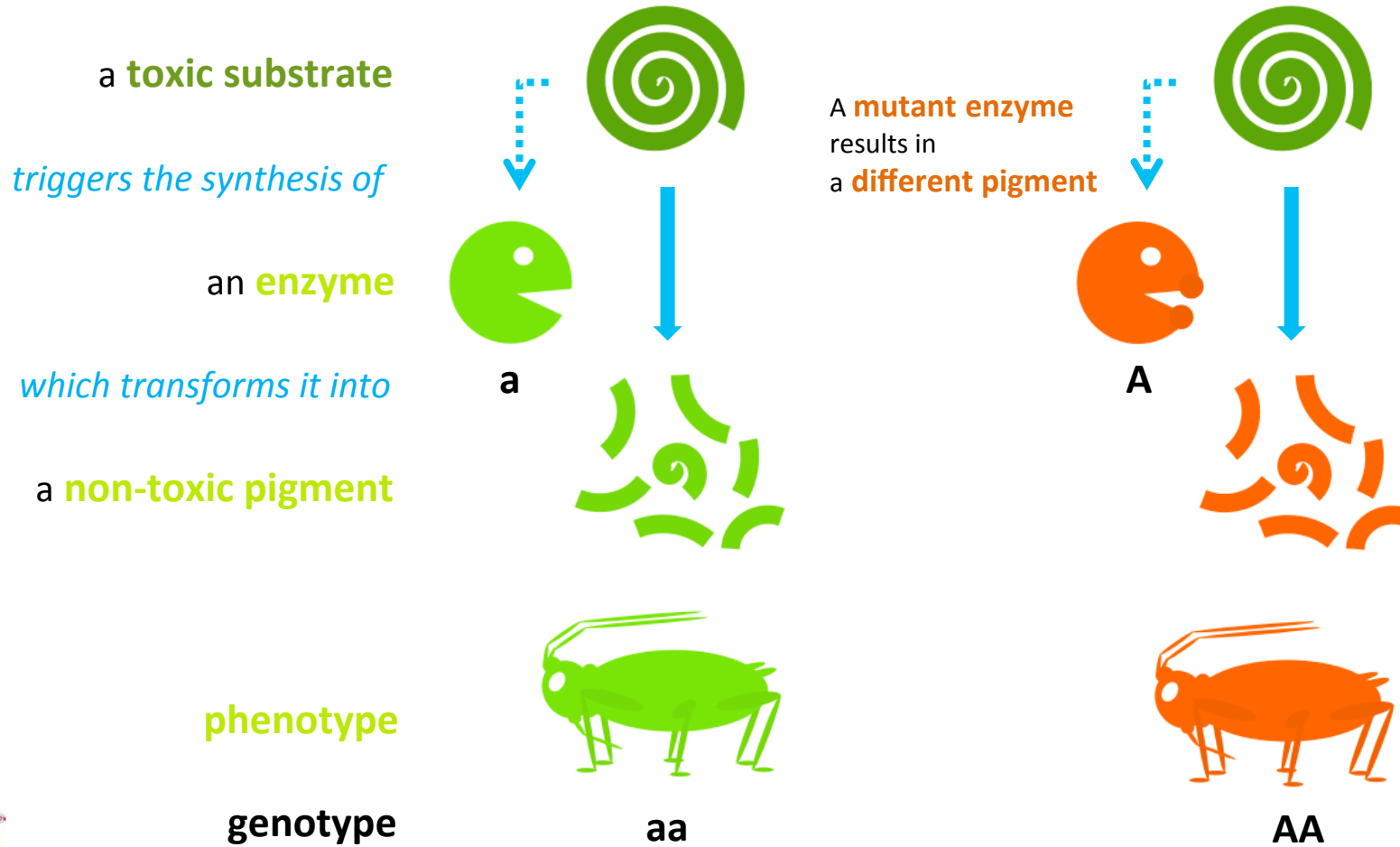
A hypothetical scenario

color synthesis in aphids



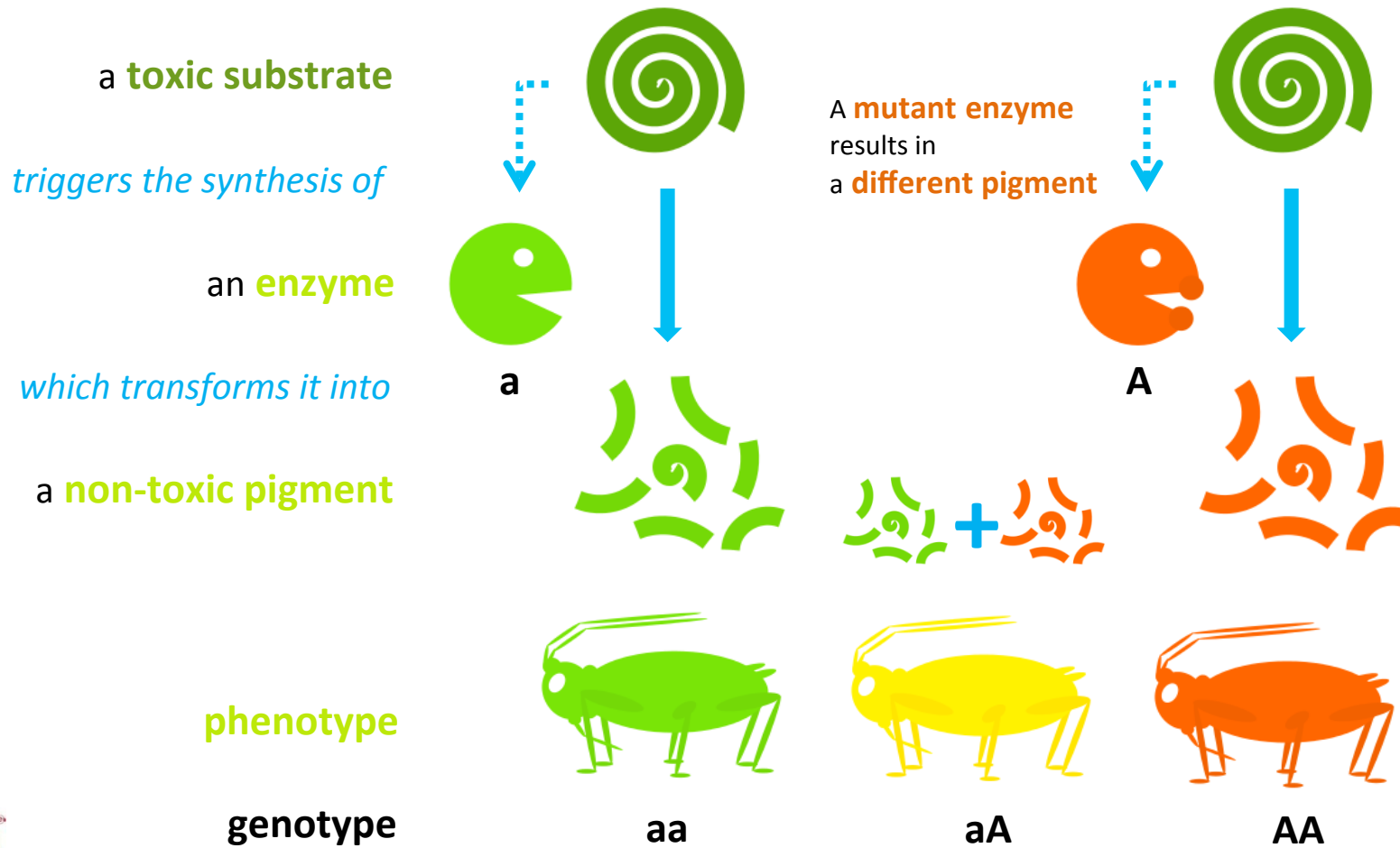
A hypothetical scenario

color synthesis in aphids



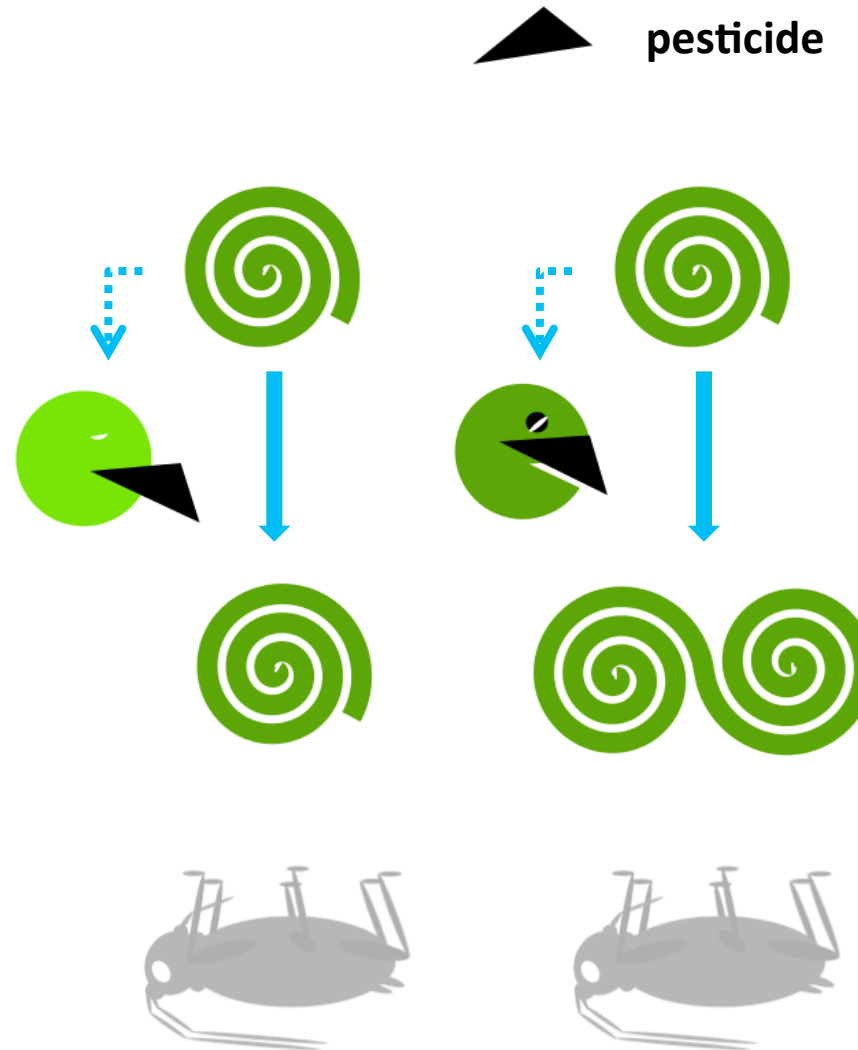
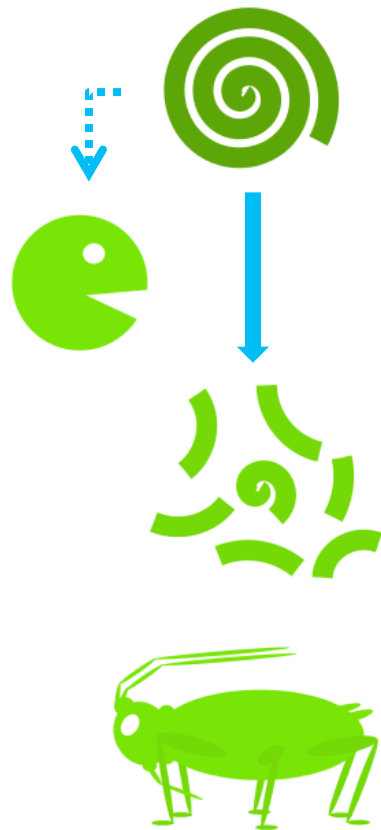
A hypothetical scenario

color synthesis in aphids



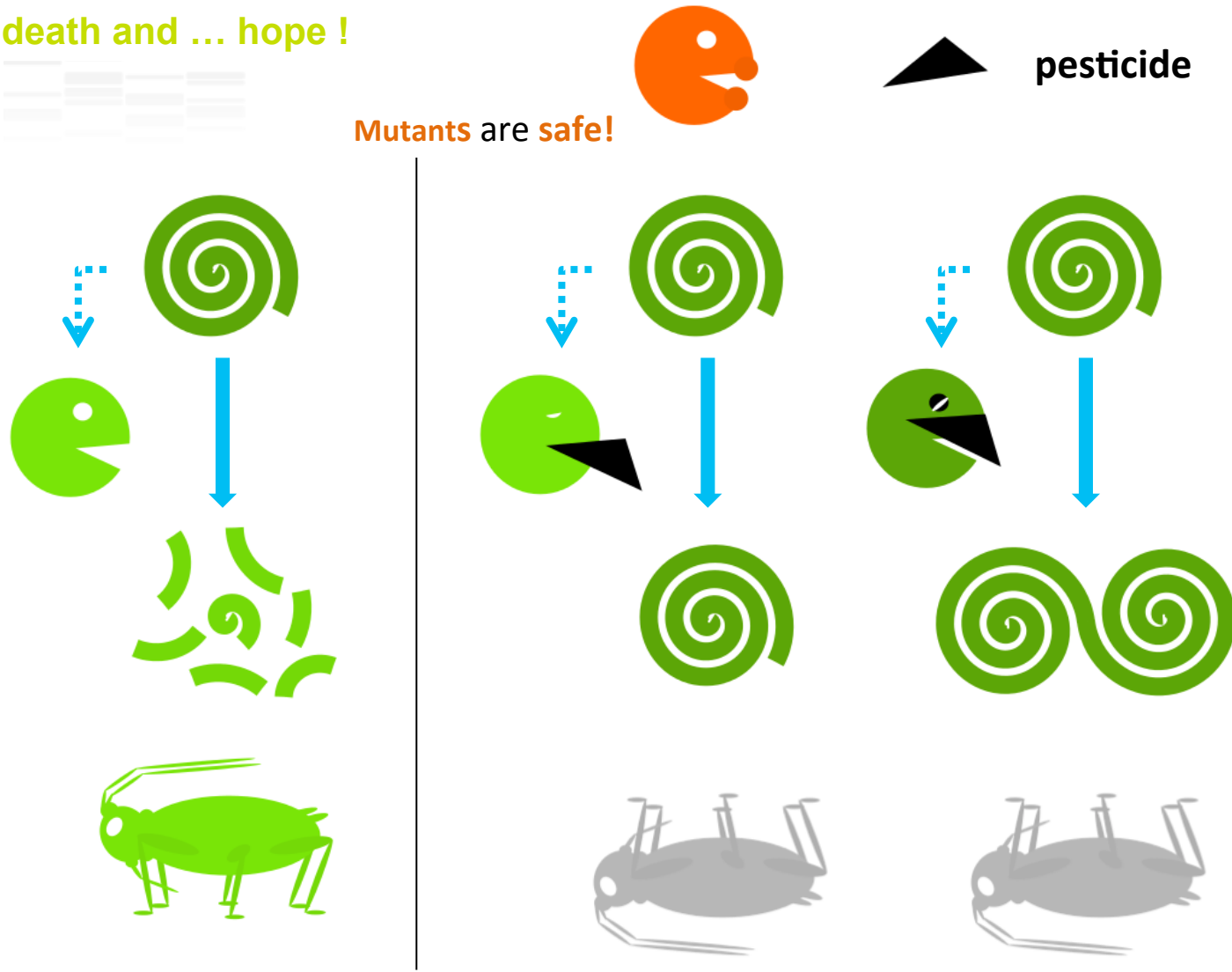
Pesticide

death and ... hope !



Pesticide

death and ... hope !



Selection

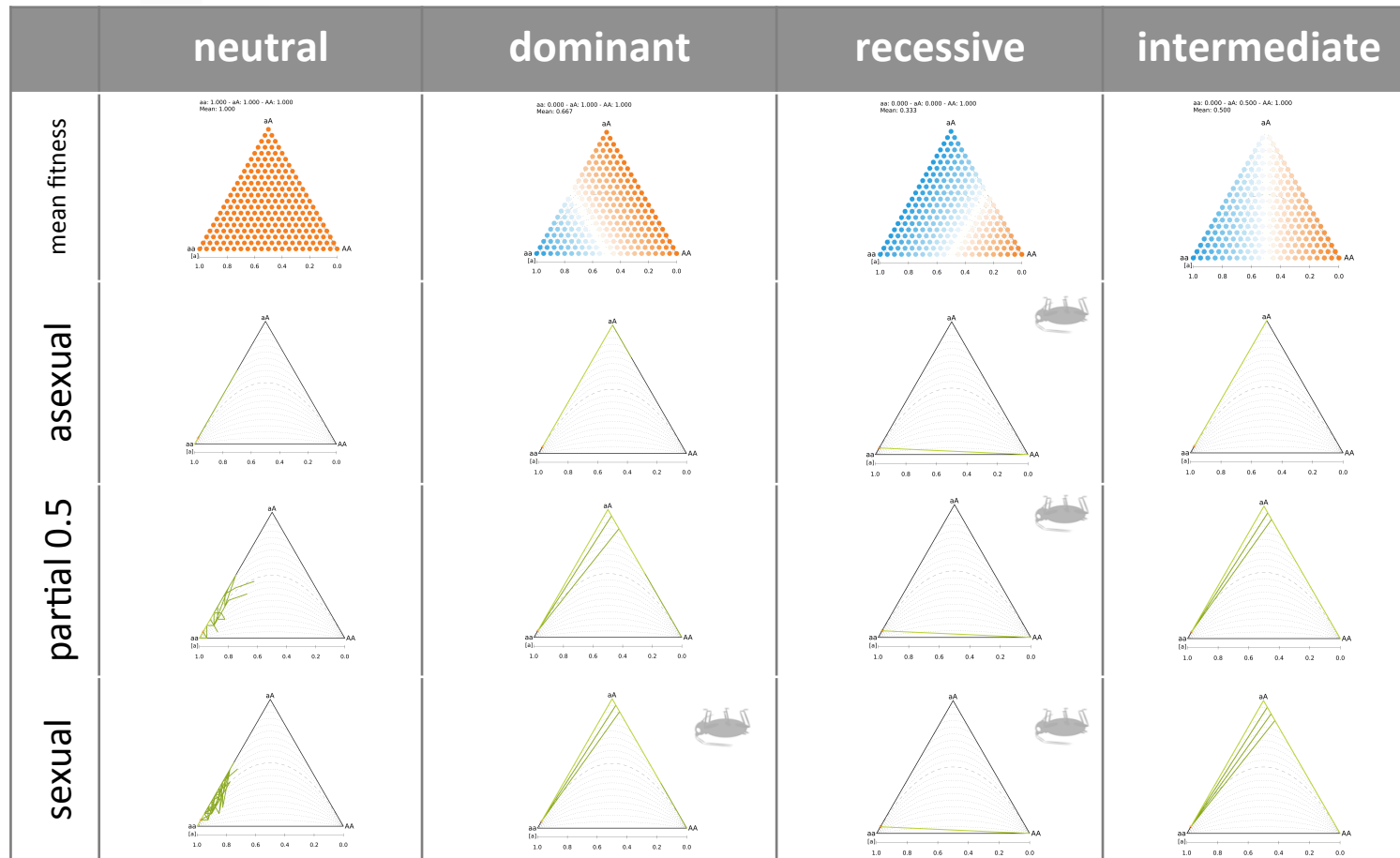
what does it look like

	wildtype aa natural	wildtype aa	heterozygous mutant aA	homozygous mutant AA	symbol
neutral					
dominant					
recessive					
intermediate					

Simulation I

all non-resistant individuals die

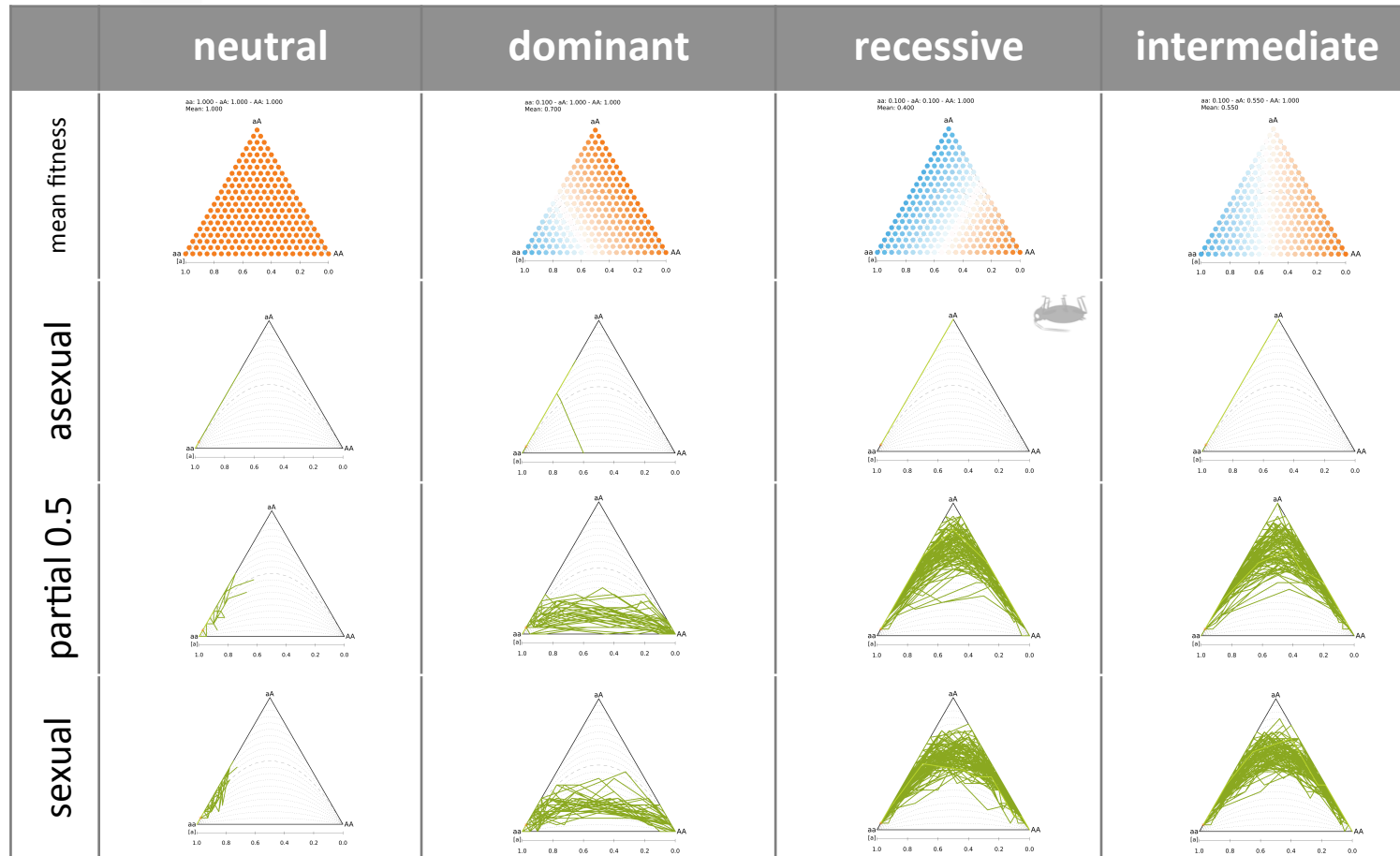
$\mu \dots 10^{-6}$
 $N \dots 20$



Simulation II

10% of the non-resistant individuals survive

$\mu \dots 10^{-6}$
 $N \dots 20$



Conclusions

pesticides and evolution

Pesticides

- responses to pesticide treatment depend on the mode of action of the pesticide and the reproductive mode of the pest species
- one copy of a “resistance allele” may be enough, in particular if the treatment is not even 100% effective against the non-resistant genotypes – pesticides are not a long-term option
- if a resistant genotype exists ... it will spread!

Evolution

- adaptive evolution depends on the underlying “fitness landscape” and the reproductive mode of the adapting species
- one adapted individual may be sufficient to very quickly invade a new environment, especially if there are also some non-adapted mating partners around
- “killing” a species takes quite some force

Outlook

still to come

extending the model

- more alleles – multidimensional fitness landscape
- more loci – crossing over & linkage disequilibrium
- modeling genome evolution

extending the analysis

- times to fixation
- redefining population genetic parameters
- Can we infer the rate of asexual reproduction from genetic data?

== __ ==

Thank you

everyone

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... in particular:

- Solenn Stoeckel and Jean-Pierre Masson
- Denis Roze (CNRS Roscoff) for some first ideas
- my “aphidophile” colleagues for the inspiration
- INRA and the Région Bretagne for funding my work

