



On the Role of Stress-Induced Mutagenesis in Evolution

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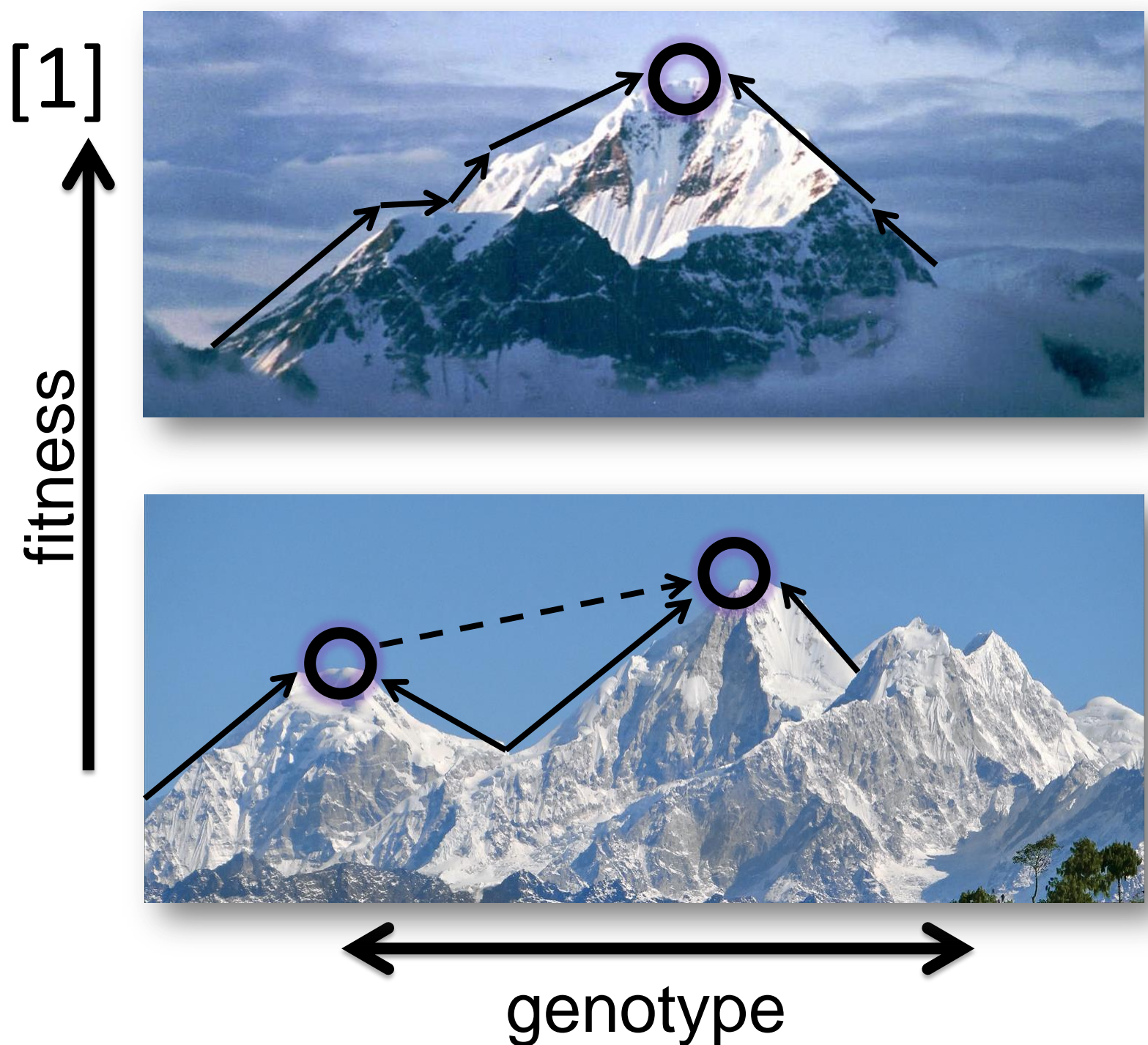
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Background

Mutagenesis is induced by stress responses in various species of bacteria and even in eukaryotes (Galhardo et al. 2007, Foster 2007).

In a previous work (Ram & Hadany 2012) we studied the evolution of stress-induced mutagenesis in constant and changing environments. We showed that **stress-induced mutagenesis (SIM)** is favored by selection over constant rate mutagenesis because it generates beneficial mutations when they are most needed.

Complex adaptations require **two or more mutations that are jointly advantageous but separately deleterious**, and therefore presents an open evolutionary question, first described by Sewall Wright (1931) and popularized using the **fitness landscape** metaphor:



Sign	Name	Estimate	Citation
s	Selection coefficient	0.001-0.01	5,6
H	Double mutant advantage	1-10	7
U	Genomic mutation rate	0.003-0.0004	7,8
μ	Beneficial site mutation rate	$U/5000$	6
τ	Mutation rate increase	1-100	9
N	Population size	10^5 - 10^{10}	10

Table 1. Model parameters and estimated values for bacteria.

Adaptation rate results

The adaptation rate v as a function of the mutation rate fold-increase τ is approximated with normal mutagenesis (NM), constitutive mutagenesis (CM) and stress-induced mutagenesis (SIM) by:

$$v_{NM} \approx 4NH\mu^2 \left(1 - \frac{U}{s}\right)$$

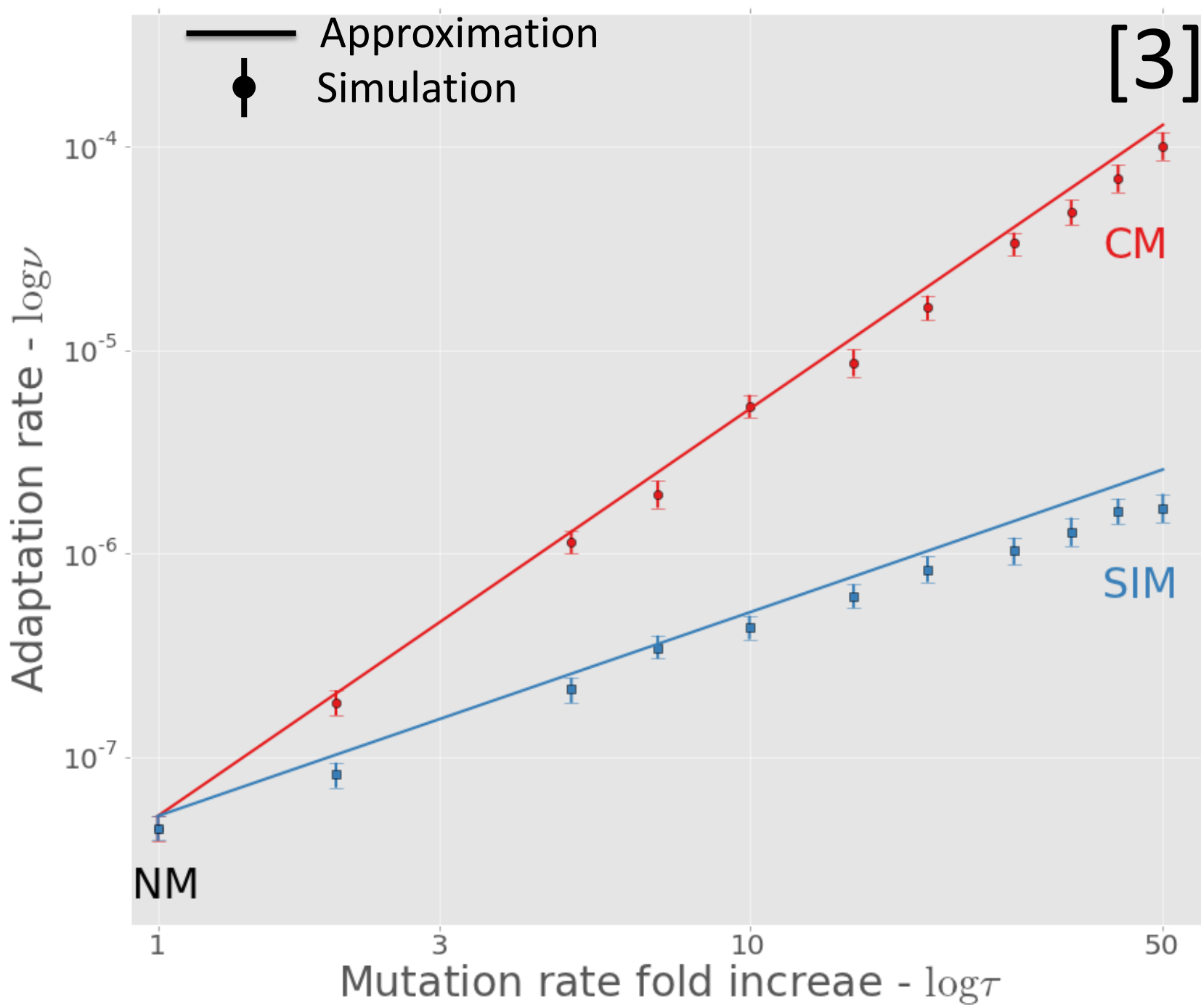
$$v_{CM} \approx v_{NM} \cdot \tau^2 \left(1 - \frac{\tau U}{s}\right)$$

$$v_{SIM} \approx v_{NM} \cdot \tau(1 - \tau U)$$

SIM increases the adaptation rate of complex traits:

solid lines are analytic approximations, markers are results of simulations (see below), error bars are 95% CI. Both axes are in log scale – the slope of the CM line is twice as steep as the slope of the SIM line.

The difference between the approximations and the simulations is explained by appearances of AB on deleterious backgrounds in the simulations (“rise of the living dead”).



SIM is more efficient than CM

Adaptability: adapting to new conditions

- Constitutive mutagenesis (CM) increases the rate of complex adaptation τ^2 -fold (τ : mutation rate fold-increase).
- Stress-induced mutagenesis (SIM) increases the rate of complex adaptation τ -fold.

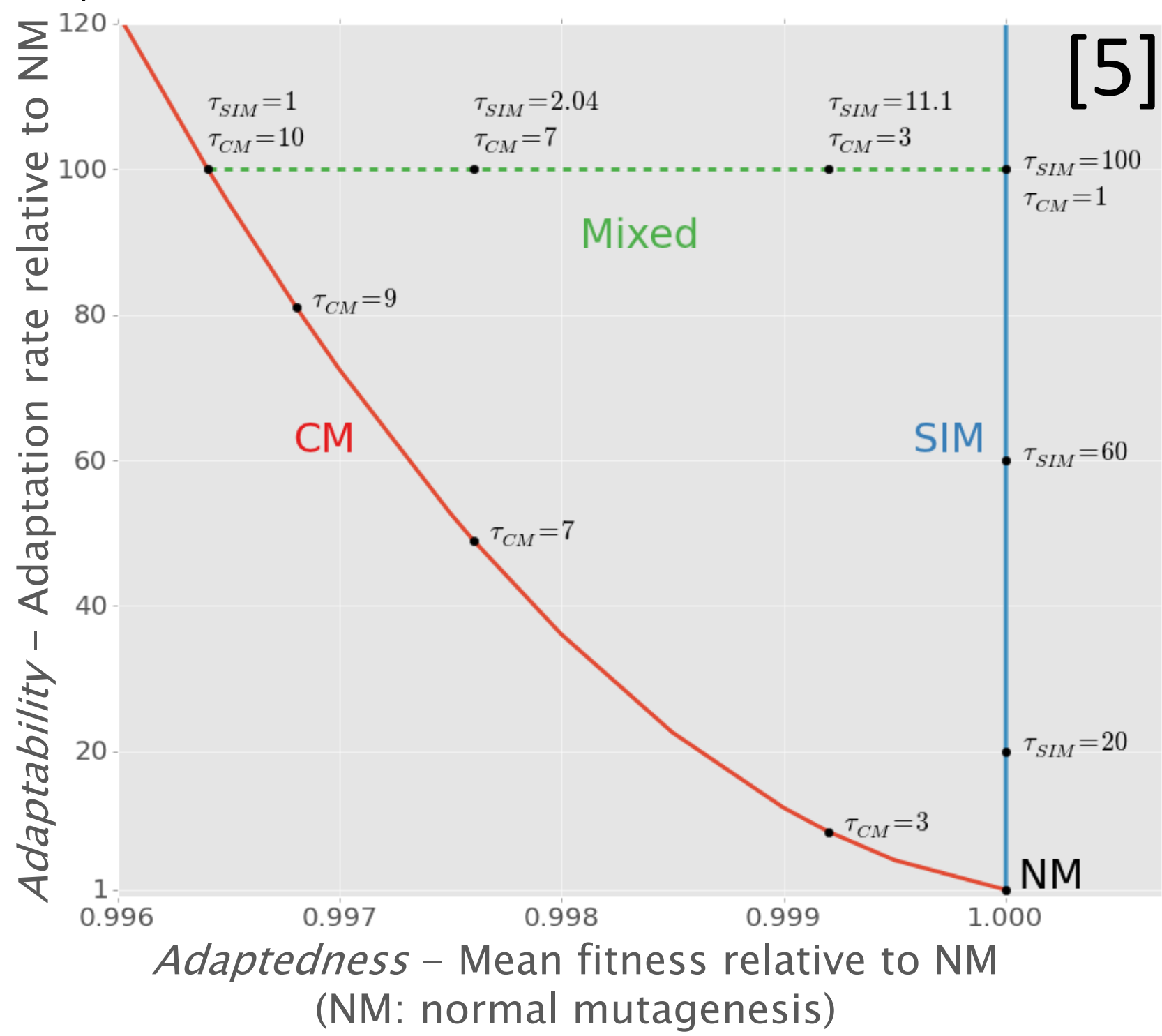
Adaptedness: staying adapted to existing conditions

- CM decreases the population mean fitness in stable environments $\approx(1-U\tau)$ -fold due to the accumulation of deleterious mutations (Kimura & Maruyama 1966).
- SIM slightly increases the population mean fitness (Ram & Hadany 2012).

Breaking the adaptability-adaptedness trade-off

Both CM and SIM increase the **adaptability** of populations, represented by their adaptation rate. SIM, in contrast to CM, doesn't reduce the **adaptedness** of populations, represented by the mean fitness in stable environments.

In **mixed strategies** all individuals increase their mutation rate τ_{CM} -fold and stressed individuals increase their mutation rate τ_{SIM} -fold. These strategies represent the space between the CM and SIM lines.

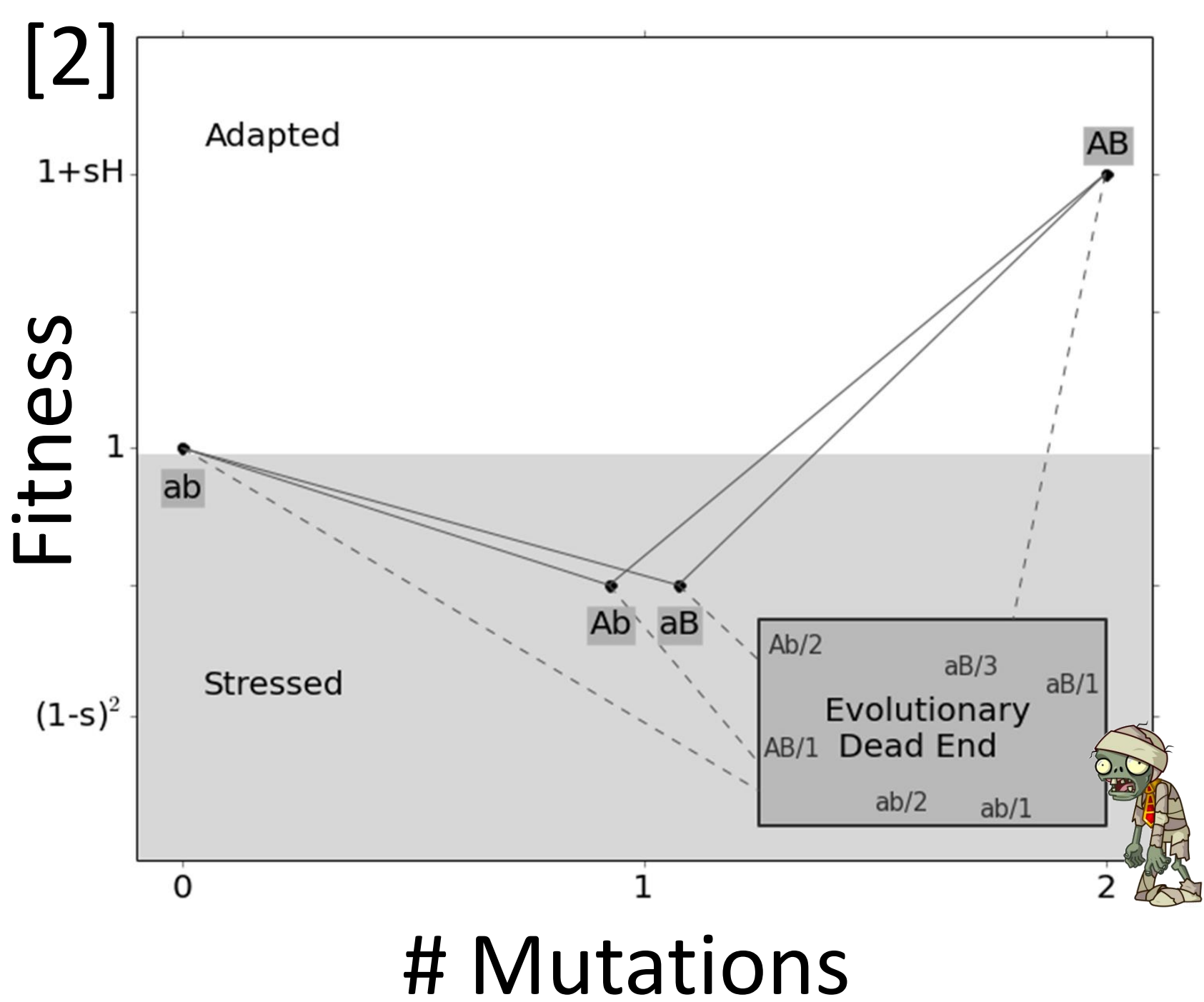


Model

Figure 2 describes a two-locus (A/a and B/b) model of **complex adaptation**.

Each node represents a genotype. Genotype **ab** is the **wildtype** local adaptive peak, **AB** is the **global adaptive peak** with the highest fitness, and the single mutants **Ab** and **aB** are **adaptive valleys** with fitness lower than the wildtype. “**Evolutionary Dead End**” represents genotypes with deleterious mutations that will not contribute to adaptation.

Lines define mutations and denote the relevant mutation rate: **U** for background deleterious mutations (dashed lines) and **μ** for mutations in the A/a and B/b loci (solid lines).

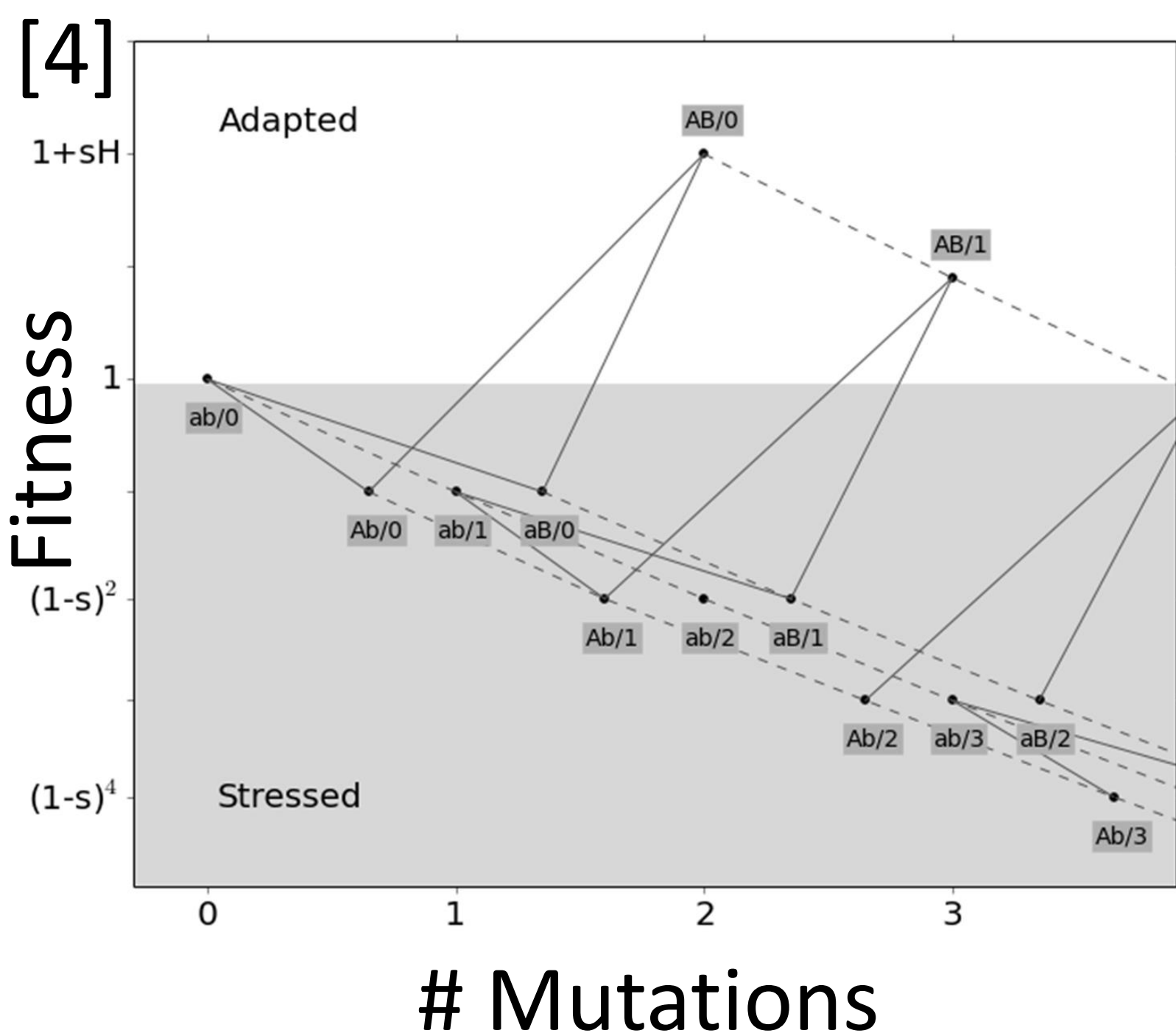


Simulations

Figure 4 describes our **multi-locus Wright-Fisher simulations**, which also account for **genotypes with deleterious mutations** – denoted by the number after the slash ($ab/2$ is ab with two deleterious mutations) – so there are no “**Evolutionary Dead Ends**”.

The figure shows up to three mutations for simplicity, the simulations have up to 25.

The simulations do not make assumptions on the distribution of deleterious alleles at the **mutation-selection balance**, but rather allow this balance to evolve before the adaptation process starts.



Summary

We compared the **adaptability** (ability to adapt) and **adaptedness** (ability to stay adapted) of populations with and without stress-induced mutagenesis.

We showed that **stress-induced mutagenesis increases the adaptability** of a population and that in contrast to constitutive mutagenesis, **it does not jeopardize its adaptedness**.

Literature cited

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