The mutation rate itself may change the order of adaptive mutations Alexandre de Aquino Soares<sup>1</sup>, Lucas Wardil<sup>1</sup>, Ronald Dickman<sup>1</sup> and Louis Bernard Klaczko<sup>2</sup> <sup>1</sup>Universidade Federal de Minas Gerais, <sup>2</sup>Universidade Estadual de Campinas

### Introduction

Yampolsky and Stoltzfus [1] presented evidence that a mutation bias, when opposite to a fitness bias, changes the probabilities of fixation of the alternative alleles depending on the overall mutability *m*. Our objective is to understand this effect, and from it to understand how different paths to a common final genotype may be biased with *m*, if so.

# Single step

Model



Dynamics



#### Conclusions

- Mutability may radically change the odds of fixation
- The turning point is sensitive to the organism;
- It may be not too sensitive to the selective pressure;
- At very low *m*:

• Clonal interference is very rare;

• High mutation rate is favored;

• Who "tries" to fix more often, wins.

• At very high *m*:

• Clonal interference is the rule;

• High fitness is favored;

• Who competes better, wins.

• The most fit allele fixes faster;

• Dynamics is the most variable for intermediate *m*.

## Multiple steps

#### Model



#### Final state



#### Effect





#### Predictability

### Summary



• Mutability again changes the odds of fixation

• Paths with decreasing rates are favored for low *m*;

#### Reference

1. Yampolsky, L.; Stoltzfus, A. EVOLUTION & DEVELOPMENT 2001, 3, 73 – 83.