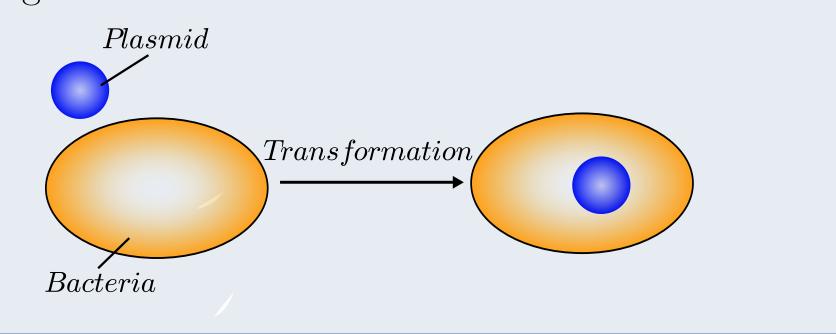
Simulation and Theory of Bacterial Transformation

JD Russo, Jiajia Dong

Department of Physics and Astronomy, Bucknell University

Introduction

- Ubiquitous threat of antibiotic resistant bacteria
- Investigate effect of different cellular transformation rates on antibiotic resistant bacterial population growth



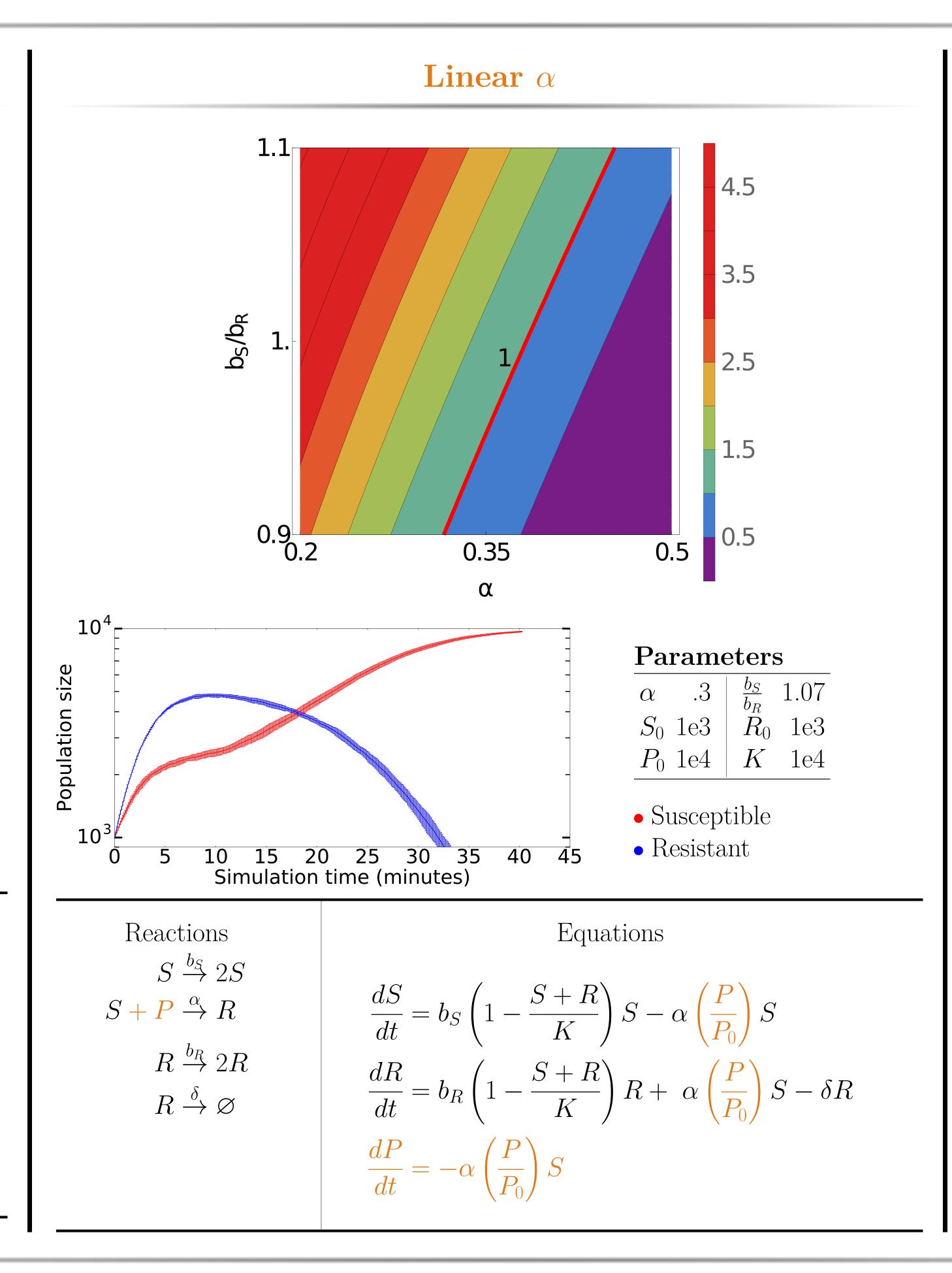
Biological Background

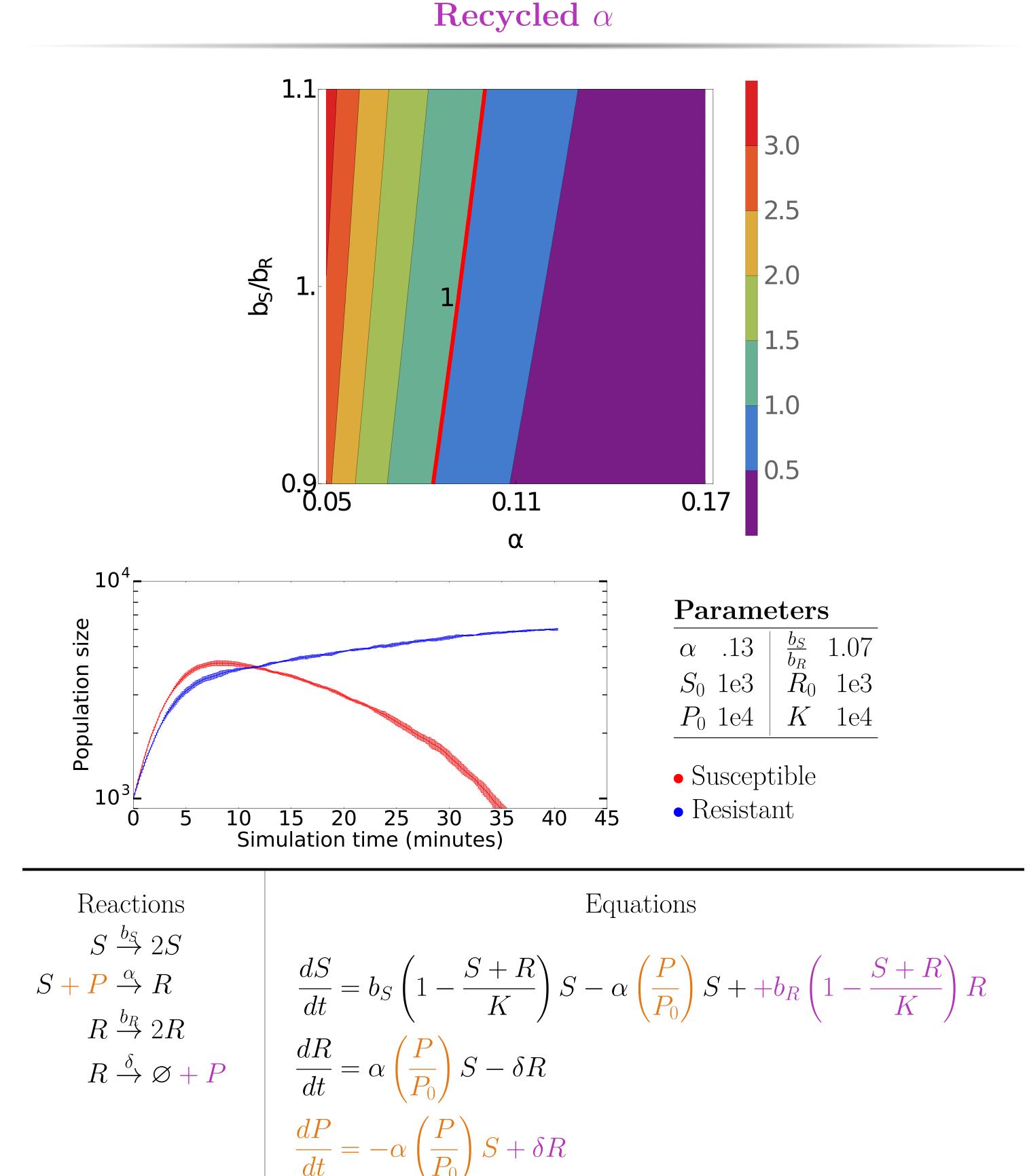
- Plasmids
- Fitness cost
- Transformation

Simulation Methods

- Combined approach of Kinetic Monte Carlo simulation and numerical modeling
- Gillespie algorithm
- Well-mixed population
- Carrying capacity
- Constant, Linear, Recycled α
- Symmetric division
- Realtime conversion

Constant α 0.17 0.11 Parameters S_0 1e3 $\stackrel{\circ}{R}_0$ 1e3 $P_0 \ 1e4 \ K \ 1e4$ Susceptible Resistant Simulation time (minutes) Reactions Equations $S \stackrel{b_S}{\rightarrow} 2S$ $\frac{dS}{dt} = b_S \left(1 - \frac{S+R}{K} \right) S - \alpha S$ $S \xrightarrow{\alpha} R$ $R \stackrel{b_R}{\rightarrow} 2R$ $\frac{dR}{dt} = b_R \left(1 - \frac{S+R}{K} \right) R + \alpha S - \delta R$ $R \xrightarrow{\delta} \varnothing$





Conclusions

- S/R transition point depends on both rate and mechanism
- Population extinction in linear case

Future Work

- Simulation on a lattice
- Adding antibiotics
- Asymmetric division

Acknowledgements/References

Thank you etc etc

- Source 1
- Source 2