

# **Background and Motivation**

- Pseudomonas aeruginosa is an opportunistically pathogenic bacterium that causes 51,000 infections per year in the United States.
- *P. aeruginosa* is quickly becoming antibiotic-resistant, rendering current treatments impractical.
- Bacteriophage (viruses that selectively infect and kill bacteria) could be an alternative.
- **Research Question:** What assumptions are necessary to accurately capture the interaction between phages and bacterial cells in a controlled environment?

# **Experimental Setup**

- We considered optical density (OD) measurements of a strain of P. aeruginosa over 18 hours.
- Optical density is a quantitative, unitless measure that indicates the degree of light scattering through a sample.
- Six experiments were conducted, in each of which varying amounts of a novel P. aeruginosa phage were added, while the initial amounts of bacterial cells remained constant between tests.
- **Research Question:** How significantly does bacterial debris contribute to optical density readings?

# Model Development



#### The SIMPL Model:

$$\frac{\mathrm{d}S}{\mathrm{d}t} = \underbrace{(1-\mu)\rho S\left(1-\frac{S+I}{K_1}\right)}_{\text{reproduction without mutation}} - \underbrace{\alpha SP}_{\text{infection}}, \\ \frac{\mathrm{d}I}{\mathrm{d}t} = \underbrace{\alpha SP}_{\text{infection}} - \underbrace{\eta I}_{\text{burst}}, \\ \frac{\mathrm{d}M}{\mathrm{d}t} = \underbrace{\rho_M M\left(1-\frac{S+I}{K_1}-\frac{M}{K_2}\right)}_{\text{reproduction}} + \underbrace{\mu\rho S\left(1-\frac{S+I}{K_1}\right)}_{\text{reproduction with mutation}}, \\ \frac{\mathrm{d}P}{\mathrm{d}t} = \underbrace{\beta\eta I}_{\text{burst}} - \underbrace{\sigma SP - \sigma IP}_{\text{attachment}}, \\ \frac{\mathrm{d}L}{\mathrm{d}t} = \underbrace{\eta I}_{\text{burst}} - \underbrace{\gamma L}_{\text{decay}}, \end{aligned}$$

# **A SIMPL Model of Phage-Bacteria Interactions Accounting for Mutation and Competition**

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# **Qualitative Analysis** 3.5 2.5 2 -2 -0.5 Equilibrium 3.5 e 2.5 1.5 0.4 0.2

- The system has three equilibria: (1) bacteria-free, (2) phage-free, (3) coexistence.
- In the SIP subsystem, two transcritical bifurcations occur at ab = 1 and ab = 1 + ac, where  $ab = \frac{\alpha\beta}{2}$  and  $ac = \frac{(1-\mu)\rho}{2}$ .
- No steady-state is globally stable; depending on the initial amount of bacteriophage, there are scenarios where phage win out despite a parameter combination where bacteria can survive.
- By a theorem of Thieme, the long-term behavior of the SIMPL system is asymptotic to the SIP subsystem.

# Parameter Estimation and Optimization Problem

 Take the total bacterial contribution to optical  $\frac{dB}{dt} = \left(\rho S + \rho_M M\right) \left(1 - \frac{S+I}{K_1}\right) - \rho$ 

where  $0 \leq \epsilon < 1$ .

Optimization problem:

$$\min_{\boldsymbol{\theta}\in T_{ad}} J(B, \boldsymbol{\theta}) := \frac{\omega}{2} \sum_{i=1}^{6} \sum_{j=1}^{N_t} (B_i(t^j) - B_i^j)^2 + \frac{\nu}{2} \|\boldsymbol{\theta}\|_2^2, \text{ subject to (1)}$$

- We numerically solved the optimization problem in MATLAB using **fmincon**, MultiStart, ode15s.
- We assessed the accuracy of our parameter estimates using the relative  $l^2$  error percentage:

$$EP = \sqrt{\frac{\sum_{i=1}^{6} \sum_{j=1}^{N_t} (B_i(t^j) - B_i^j)^2}{\sum_{i=1}^{6} \sum_{j=1}^{N_t} (B_i^j)^2}} \times 100$$

(1)



I density, 
$$B = S + I + M + \epsilon L$$
,  
 $o_M\left(\frac{M}{K_2}\right) - (1 - \epsilon)\eta I - \epsilon \gamma L$ , (2)

#### Results



- as live cells.
- infection, (3) lysis, (4) mutation.

# Conclusions

- *Pseudomonas aeruginosa* is a deadly bacterium that is mutating to become largely resistant to antibiotic treatments.
- Though bacteriophage therapy is a promising alternative treatment method, we need a solid understanding of the dynamics between bacterial and phage populations to successfully implement this therapy.
- Mathematical models like the one developed in this project provide important insight into this interaction and can be made simple while keeping results accurate. • We concluded that bacterial debris contributes roughly 31% as much as live bacterial cells to optical density measurements.
- A cocktail of multiple strains of bacteriophage will likely be needed for effective clinical use.

# **Future Work**

- Extend our model to an in vivo setting by incorporating the removal of bacterial cells due to a patient's immune response.
- Consider a combination treatment of bacteriophage and antibiotics to develop an optimized treatment strategy.
- Explore new methods to convert optical density to bacterial counts.

# **References & Acknowledgements**

#### References:

- 1. C Peterson, et al. A SIMPL Model of Phage-Bacteria Interactions Accounting for Mutation and Competition. Submitted Aug 2024, Revised Jan 2025.
- 2. J Serralta. Isolation and Characterization of a Novel Bacteriophage for Pseudomonas aeruginosa. MS thesis. University of North Texas Health Science Center at Fort Worth, 2023.

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• Our model fit the observed optical density data with an error of 8.5%. • Lysed bacterial cells contribute to optical density measurements about 31% as much

• Observed four distinct phases in phage-bacteria interaction: (1) susceptible, (2)

#### Investigate the effect of bacterial debris on bacterial carrying capacity.

Acknowledgements: This work was funded by the NSF Math for Human Health RTG (DMS-

