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Simulating Spread of Antimicrobial Resistant Bacteria in the Pig Pen - try our online tool.

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Background

Antimicrobial resistance is an increasing challenge. For example, 40% of *Escherichia coli* in Danish pigs are resistant to tetracycline.

The purpose of the MiniResist project is to find treatment protocols that **reduce antimicrobial resistance development**. The project includes a field trial with +2,000 pigs. This poster presents an app that can be used for exploring the effects of different treatment strategies.

Model

The experiments inform a mathematical model that incorporates multiple pigs and multiple bacterial strains interacting in a pen.

Antimicrobial (AM) dependent growth of individual bacterial strains in individual pigs is simulated, including dynamic effects of the excretion to and uptake of bacteria from the common pen environment. The model app includes AM treatments where length, dose, and time can be **specified by the user**.

MiniResist ← → C 🔒 https://kagr.shinyapps.io/MiniResist/ MiniResist Overall Runs Sample Pigs About Welcome to the MiniResist WebApp. This app simulates how AntiMicrobial Resistant Bacteria (AMRB) spread between pigs in a pig Fraction of Anti Microbial Resistant Bacteria (AMRB) in the pen pen following antimicrobial treatments. For an introduction and user guide press the 'About' pane to the right. Calculate 0.8 Select number of treatments: 2 0.6 AMRB

Growth rates of more than 50 *E. coli* strains, collected from Danish farms, were estimated for different AM concentrations to inform the model.

The mathematical model captures the temporal effects of the field studies. The model was then integrated into an free online WebApp using shiny and shinyapps in R.

Results

The figure presents how a shorter duration of AM use may reduce the peak fraction of resistant bacteria (compare black to grey line).

Several further results are discussed in our article [1]. I.e. how diarrhea, excretion rates, and transmission rates affect the level of AM resistant bacteria.

Star	day of treatment 1: [day 1-135]
5	
Star	day of treatment 2:
35	

Length of each treatment [1-14 days]:



Treatment dose per day:

• High (200%)

Normal (100%)

Low (50%)

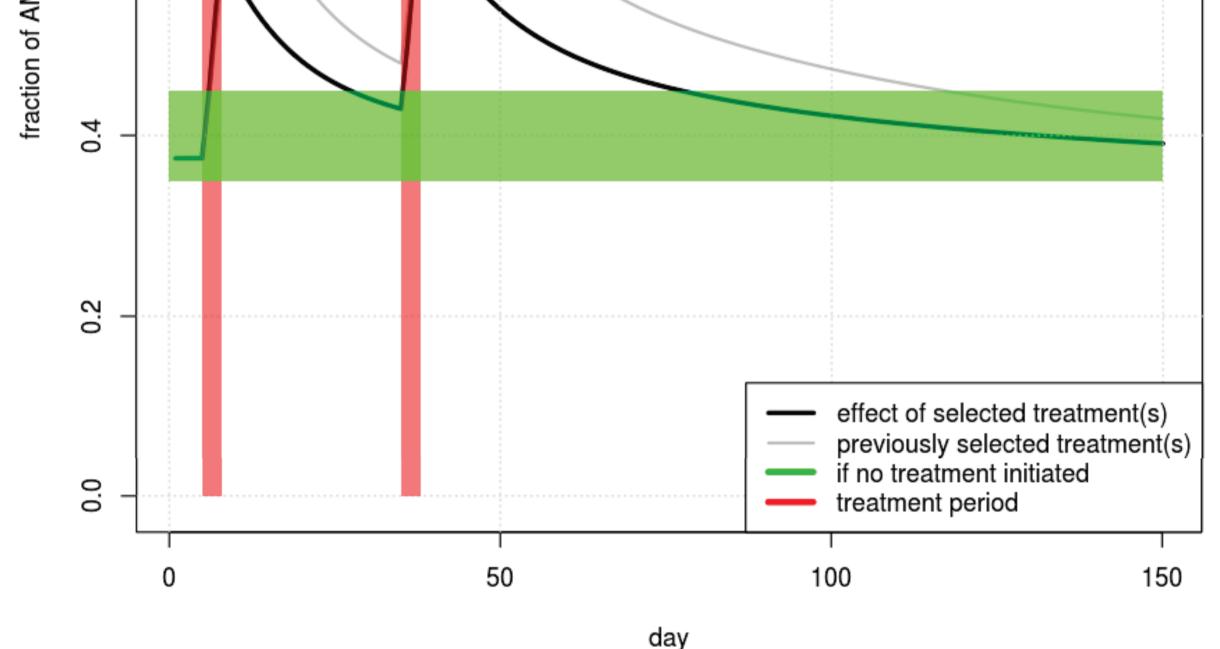
Initial fraction of resistant bacterial strains:



Do bacteria cause diarrhoea?:

One of the susceptible strains causes diarrhoea

One of the resistant strains causes diarrhoea



Conclusion

AM resistance development based on different treatments strategies can be studied using our app. The tool presented is freely available online at kagr.shinyapps.io/MiniResist.

The model captures all information regarding presence of AM bacteria obtained by our field study.

This model can be used to devise treatment strategies to minimize the level of AM resistant bacteria. However, any proposed strategy should be validated by field experiments.



Try our online WebApp!

It predicts the level of antimicrobial resistant bacteria based on treatments that you define.

https://kagr.shinyapps.io/MiniResist/



Read the article! The research presented in this poster is published in: [1]

"How Fitness Reduced Antimicrobial Resistant Bacteria Survive and Spread: A Multiple Pig - Multiple Bacterial Strain Model."

PLoS ONE 9(7): e100458. doi:10.1371/journal.pone.0100458



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