

Researchers assess use of drug-susceptible parasites to fight drug resistance

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Andrew Park, an associate professor at the University of Georgia, led the research team that developed a model for evaluating a potential new strategy in the fight against drug-resistant diseases. Credit: Dot Paul/University of Georgia

Researchers at the University of Georgia have developed a model for



evaluating a potential new strategy in the fight against drug-resistant diseases.

The strategy would take advantage of parasite refugia—host populations that have not been treated with drugs, thereby serving as "safe zones" where parasites don't develop drug resistance. When parasites from refugia mix with their drug-resistant counterparts in the general population, they could reduce the incidence of drug-resistance overall, which may help prolong a drug's effectiveness.

The research, just published in the Royal Society journal *Biology Letters*, offers a way to assess when such an approach is likely to work—information that could help in the increasingly urgent hunt for alternatives to the current suite of parasite-fighting drugs.

"Once resistance emerges, you might squeeze a little bit of life out of a drug by tweaking it, but often, very quickly, that entire class of drugs will become useless," said Andrew Park, an associate professor in the UGA Odum School of Ecology and College of Veterinary Medicine, who led the research.

"Right now, we're really struggling to manage diseases like MRSA and extensively drug-resistant TB," he said. "We're at a point where new classes of drugs don't grow on trees anymore; it can take 15 to 20 years to develop them. It's a challenge just to keep pace. Refugia represent a management strategy that's being considered as a way to slow down the development of drug resistance, particularly in animal health."

Park and his colleagues created a model to predict the effects of refugia on two important outcomes: the overall prevalence of infection within a population and the proportion of those infections that are drug-resistant.

The model incorporates two main variables. One is the level of drug



<u>coverage</u>—the proportion of the overall population that has been treated against parasites. The other is the degree of mixing between the treated and untreated groups.

In the case of a disease like heartworm, which is transmitted by mosquitoes and infects both wild animal populations and companion animals, there is some natural variation in both drug coverage and mixing. Both variables can also be controlled, to an extent.

"There might be variation in terms of how compliant animal owners are in giving heartworm preventive and whether dogs are kept indoors a lot versus just allowed to run around outside," Park said.

In this case, the mixing is unintentional rather than planned. "Given that it's happening, we should at least understand what the consequences might be for drug resistance," he said.

The model yielded predictions about how changing the levels of drug coverage and mixing would affect prevalence and resistance, and also shed light on the evolutionary processes at work.

"The dogma is that increasing contact between treated and untreated groups would be predicted to simply increase prevalence and decrease frequency of resistance, but we found that the relationships were much more complex than that," Park said. "And we gained a lot of insight into the different roles of selection—how different strains are favored because of the host environment—and gene flow, the movement of drug resistant and drug susceptible parasites between host groups."

Park said that his team's model serves as a general outline for considering the use of refugia as a management strategy, providing a blueprint for future models to predict outcomes in specific host-parasite systems.



"With refugia, the idea is to dilute overall <u>drug resistance</u>," Park said. "We can say, yes, we're putting parasites into the population, but they're the right kind of parasites, and in some cases that may be better than having it just flooded with the wrong kind of parasites that we can never treat. But we need this kind of model to understand all these interactions in order to know when that's a good idea."

More information: Andrew W. Park et al. Refugia and the evolutionary epidemiology of drug resistance, *Biology Letters* (2015). DOI: 10.1098/rsbl.2015.0783

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