Current Status
Third year of a five-year project

Funding Sources and/or Partners
State Wildlife Grants, Purdue University, Nongame Fund

Project Personnel
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Background and Objectives
The main goal of this project is to determine whether interstate highways in Indiana are restricting the movements of wild animal species of various sizes, which may result in declines of these species due to losses in genetic diversity. We will investigate the ability of six mammal species to cross the interstate by analyzing genetic information and movement patterns of wild populations.

Methods
We have been trapping six common species of mammals, including raccoons, opossums, gray squirrels, fox squirrels, chipmunks and white-footed mice, from forests on opposite sides of interstate highways in Indiana. Even though these species are common, we hope the results obtained will be transferable to less-common species, threatened or endangered wildlife. For sampling, we have...
Animals that are related share a lot of the same DNA, and the more related they are, the more DNA they have in common. Therefore, a population of animals is likely to share DNA at a relatively high rate because there will be a lot of relatives living near each other.

Populations on either side of an interstate can be compared for evidence of gene sharing and relatedness. Using this information, we can determine if the interstate is separating two populations or if the animals on either side are actually one population. If the animals on the two sides of the interstate are two populations, then the interstate may be creating a barrier to animal movements. This effect may be more evident in species that are more reluctant to cross over roads or to use culverts to cross under roads (such as chipmunks and squirrels) than species that are not as reluctant (such as raccoons and opossums). However, if we determine that the animals from either side of the interstate form one population, then the animals are successfully crossing the interstate and gene flow is occurring.

Detailed data on the movement patterns of animals help us understand the day-to-day response of these animals to highways. That understanding of those daily movement responses, in combination with the genetic data, are critical to building computer models that we will use to predict the consequences of interstate highways in Indiana on the long-term persistence of viable populations of selected forests that are separated by the interstate, but that otherwise have good habitat for these species. These are so-called “paired” sites, because each site actually has two parts—one on either side of the interstate.

We set trapping grids in these habitats to catch the mammals that inhabit the forest close to the interstate. We use box traps, which are like wire cages, to trap the animals. The animal is drawn into the trap by bait in the back of the trap, and it steps on a treadle that shuts the door.

Each animal is handled carefully to reduce the stress of its experience as much as possible. We marked the animals with numbered ear tags, and we took a small genetic sample from each animal, usually a tiny piece of the tip of the ear. Collars containing radio telemetry transmitters were attached to 10 adult chipmunks, gray squirrels, fox squirrels and raccoons within each trapping grid.

All animals are released at the trap location, which we record to help us determine the spatial arrangement of the animals in the forest.

We take the tissue samples back to the wildlife genetics research lab at Purdue University to extract DNA and determine the genetic characteristics of each individual. Combining the genetic information with location data will help us determine the effect the interstate has on the local populations by allowing us to look for evidence of genetic exchange between populations on either side of the road.
wildlife. The radio telemetry collars provide us detailed data on the movement patterns of the 40 animals we are tracking at each grid. Animal locations are estimated using four automated telemetry receivers and towers that are mounted on trailers. This system allows us to estimate the location of each collared animal within a grid once every couple of minutes.

Every two or three weeks we move the four trailers between grids so we collect movement data at all of our study sites. In the lab we analyze the recorded movements of the animals to determine if there are consistent patterns in how frequently these animals turn or move slowly or quickly. We compare the movement patterns of these animals when they are close to highways to times when they are not. We then use those movement patterns to create virtual animals inside the computer.

By simulating the movements of thousands of virtual animals we estimate the strength of the barrier effects of different highways for different species. Then we compare the predictions of the computer models with the genetic data to investigate the details of how highways restrict the movement of different wild animal species.

Progress

This past year, we trapped the second set of three paired sites in central Indiana, which included study sites along I-65 near Columbus and Seymour as well as a site along I-70 near Greencastle. These trapping sites from the second year of field work occurred on private lands where we worked with permission from the landowners. We have now trapped all of the paired sites and gathered nearly all of our needed tissue samples from wild mammals. We will return to collect additional samples from gray and fox squirrels at a few sites in early 2011 to bolster our sample sizes at those locations.

While live-trapping for tissue sample collection, we placed radio-telemetry collars on the target wildlife species at two of the three paired sites in central Indiana, and the trailers with the towers and the automated receiving units have been moved between those study sites to collect data on animal locations.

During this past year we finished writing the computer program we will use to simulate animal movements and assess the strength of the barrier effects of the highway upon the wildlife species we are studying. The lab analysis of the collected DNA samples is in progress, and the genetic work will be completed during spring 2011.

The future of this project includes the collection of additional movement data using the automated telemetry system. This will take place throughout 2011. During 2011 we will also begin to construct virtual versions of our wildlife species in the computer software and run simulation models. We expect to have useful information to share with our funding partners by 2012.

Cost: $1,042,067 for the complete five-year study