How to Run the Rhino EMT's Political-Ecological Simulation Model

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Needed software and files

It is necessary to download two free software packages: the Java SE Development Kit 8, and gnuplot. See

www.oracle.com/technetwork/java/javase/downloads/jdk8-download-2133151.html, and

https://sourceforge.net/projects/gnuplot/. In addition, a text editor such as Word-pad will be needed.

You will also need to download the jini v2.1 starter kit from https://mvnrepository. com/artifact/com.sun.jini/jini-starterkit or similar. Verify that this package installs into the folder c:/jini2_1.

The id system consists of a set of Java class files. These may be found on the web at www4.uwm.edu/people/haas/downloads. The needed .zip files are idsrce.7z, and batfiles.7z.

To run the article's analyses, one needs to download wrkshpinputs.7z, wrkshppars.7z, wrkshppars.7z, wrkshpplsts.7z, wrkshplsts.7z, and emat.dfn.

If you need a PostScript to PDF translator, a free one, ps2pdf is part of the Ghostscript package available at

www.ghostscript.com/download/gsdnld.html. After installing, the path environment variable needs to be edited to include the string:

C:\Program Files\gs\gs9.19\lib;C:\Program Files\gs\gs9.19\bin. Then, at a command prompt, an Encapsulated PostScript file may be converted to a PDF file with the command

ps2pdf yourfile.eps yourfile.pdf where "yourfile" is the name of the file being converted.

General setup of an IBM

- 1. An IBM operates as follows. First, individuals are initialized along with their environment. Then, time is incremented by some amount and each individual's status is *updated* based on the model's equations describing how the individual is affected by this passage of time (which may include the individual's activities such as eating). This is incremented as each individual is updated again. These two activities are repeated until the simulation's stop time is reached.
- 2. The key idea of an IBM is that aggregate behavior patterns of a group of animals will *emerge* through the collective behavior of individuals. This is how nature works and hence an IBM is, in this way, a more faithful representation of an ecological process than an aggregate model such as a predator-prey differential equation system is.

Integrating a poaching operation model with an IBM

One way to do this is to build an *agent-based* submodel of wildlife trafficking and have it interact with a submodel composed of a population dynamics IBM of the species being traded. We call the result an *economic-ecological* model.

Overview of the rhino IBM

Spatial effects are coarsely modeled by defining a patch partition of the region and keeping track of each animal's patch membership.

Step-by-step details of the rhino IBM





Example: An economic-ecological model of South African rhinos

Input files

- id command files: krhino.id, trdrs.id, middle.id, poachers.id, anti-poach.id, and rhinoeco.id
- 2. Parameter files: trdrs-hyp.par, middle-hyp.par, poachers-hyp.par, anti-poach-shoot.par, and rhinoeco-hyp.par
- 3. Data files: knprhinos.dat, allknpveg.dat, patches.dat, and knpremoved.dat

Commands to run the example

- 1. Open a command prompt window by clicking All Programs > Accessories > Command Prompt. Change folder with cd ../../workshop (or similar).
- Set TNMNDS to 40, and NMNETS to 5. Delete all *.class files with the command del *.class, and then recompile the entire id package with compile Run_id.java.
- 3. Run the example with idalone krhino.id.
- 4. To see a list of files sorted by most recently created, enter the command dir /o-d.

Output files

- prices.plt
- 2. krhinoest.html
- 3. finalpop.dat
- average abundance: sq.plt, all abundance realizations: sqabund.plt (names are set in trdrs.id)

Plotting support

Use **gnuplot** to create complex graphics of the IBM output. In this example, running the batch file **abunds.gp** creates a **gnuplot** plot of rhino abundance over time across all management strategies.

This is accomplished in two steps. First, enter the command gnuplot at the command prompt. Second, at the gnuplot command prompt, enter the command load "abunds.gp".