Exercise Hints and Solutions for Chapter 20

Agent-based and Individual-Based Modeling: A Practical Introduction

Exercise 1

There is a region around survival-prob = 0.978 and scout-prob = 0.28 where all three criteria are met sometimes, but there may not be a combination that consistently reproduces all three.

Exercise 2

Students should produce a table very similar to Table 20.1:

Measure of model fit	Parameter set		
	1	2	3
Difference in mean	13*	49	41
Difference in standard deviation	58	3*	75
Maximum error	316	250*	404
Mean squared error	24400*	28800	30800
Number of years with results within 100 units of observed	6*	4	5

*Best value.

Parameter set 1 is best for three of the five measures, including mean squared error (probably the most meaningful measure); set 2 is best for two measures. Parameter set 1 is probably best.

Students should be aware that there are only three significant figures in the observed data (and few types of field data have even that much precision), so they should not compare model results at more than three significant figures of precision.

(The "Observed" resource abundance in Table 20.3 actually came from a simulation with r = 1.8, K = 3150, and $N_0 = 2500$; with +/-250 units of random noise. These "true" parameter values happen to also produce a mean square error of 24400.)

Exercise 3

Students can conduct a BehaviorSpace experiment like this:

The output variables are updated in this procedure:

```
to update-output
```

```
set nnd mean [min [distance myself] of other turtles] of turtles
```

```
set mnd max [max [distance myself]
of other turtles] of turtles
```

set mean-heading average-heading

```
set polarization mean [abs subtract-
headings heading mean-heading] of
turtles
```

end

We found all the calibration to be criteria to be met under most combinations of parameters in the ranges:

- vision: 7-10
- minimum-separation: 1.25-2.5
- max-align-turn: 10-20
- max-cohere-turn: 5-15
- max-separate-turn: 5-15.

▶ Experiment	×
Experiment name CalibrationExercise	
Vary variables as follows (note brackets and quotation mark	s):
["population" 8]	_
[VISION [I 5 IU]] ["minimum-separation" [0 1 25 5]]	
["max-align-turn" [0 5 201]	
["max-cohere-turn" [0 5 20]]	
["max-separate-turn" [0 5 20]]	T
Einher list values to use, for example: "my-silider" 12 7 8] or specify start, increment, and end, for example: "my-silider" [0 1 10] (note additional brackets) to go form 0.1 at a time, to 10. You may also vary max-pxcor, min-pxcor, max-pycor, min-pycor, rand	dom-seed.
Repetitions 1	
un each combination this many times	
Measure runs using these reporters:	
nnd	
polarization mpd	
	–
ne reporter per line; you may not split a reporter across multiple lines	
Measure runs at every step	
if unchecked, runs are measured only when they are over	
Setup commands: Go comma	nds:
setup no	
▼	~
► Stop condition: ► Final co	ommands:
the run stops if this reporter becomes true run at the er	nd of each run
Time limit 1000	
stop after this many steps (0 = no limit)	
OK Cancel	1

Exercise 4

This exercise implies that the model should be run with just one hunter (whereas the code in Chapter 2 includes two hunters); it is OK to use two, though the results will be different. Students need to create sliders for the three parameters and change the code to use the global variables from the sliders and to stop when 50 mushrooms have been found. Then a big BehaviorSpace experiment can report the number of ticks until the 50th mushroom is found. We ran an experiment covering wide ranges of all three parameters:

```
["switch-time" [0 5 100]]
["broad-search-angle" [1 10 181]]
["local-search-angle" [1 15 361]]
```

(We avoided search angles of zero because it's remotely possible for them to cause a hunter to search forever without finding another mushroom.)

The problem is interpreting the output systematically to find good parameter combinations. The model turns out to be highly stochastic, with random variation in results that is high compared to the effects of parameter values. One approach is to sort the results in descending order of hunting

time, and look for patterns in the parameter values that produced the shortest times. Just looking at the sorted results turns out not to be very useful, as each parameter has values over a wide range even within the model runs producing lowest hunting times:

switch-time	broad-search-angle	local-search-angle	ticks to find 50 mushrooms
40	21	136	320
20	41	106	382
80	91	106	391
45	91	121	425
65	41	121	426
90	21	346	445
45	71	166	450
15	1	91	470
15	71	151	471
45	131	106	473
45	41	181	482

However, it is useful look at the model runs producing the best hunting times statistically. First we can examine the range of model results by histogramming the hunting times over all model runs in the BehaviorSpace experiment. The histogram (below) shows that hunters sometimes took over 9000 ticks to find 50 mushrooms, but most often needed fewer than 1500 ticks.



We can define good model runs as those finishing in fewer than 1000 ticks, and look at the parameter values producing those runs. Again, histograms are useful. The following three histograms show the distribution of parameter values in the runs finishing in less than 1000 ticks.



These show that, in model runs producing best results, the broad search angle was most often less than about 80° (a rather wide range), the local search angle varied widely, and the time at which the hunter switches back to broad search also occurred over a broad range but values between 1 and 25 ticks were relatively more common. (Statistically minded students could also look for cross-correlations within the parameter combinations producing good results; we found high switching times to be associated with high broad-search angles and small local-search angles.)

A second approach is to look for statistical relations between parameter values and model results, over all model runs. Linear regression plots are a good place to start, although the above plots indicate that effects of the parameters are unlikely linear at low values. We produced the following regression relations, which indicate that the broad search angle is the only parameter with a strong effect. (Statistics such as P values may indicate that all parameters are "significant" but that is largely an artifact of using so many "observations"—model runs—in the analysis.)

In the following plots, the Y axis is the number of ticks at which the 50th mushroom was found, so good parameter values produce low Y values. They indicate that best results occur when the broad search angle is below about 60°; and the local search angle is intermediate (perhaps 90-180°) or high (near 360°) but its effect is small. It is impossible to draw any conclusion about switch time from its plot.



Simple Regression Plot

TICKS = 898.91 + 7.9119 * BROAD 95% conf and pred intervals





Simple Regression Plot

TICKS = 1558.3 + 1.2125 * SWITCH 95% conf and pred intervals