## Mathematics 1180 MATHEMATICS FOR LIFE SCIENTISTS Computer Assignment III Due January 28, 2002

We will compare the solution of a one dimensional equation describing a disease,

$$\frac{di}{dt} = \alpha i (1-i) - \mu i, \tag{1}$$

with a system of differential equations describing the same and related processes. i is the fraction of sick people,  $\alpha$  the infection rate and  $\mu$  the removal rate. We can use the usual series of commands involving **dsolve** to find and plot the solution (remember to give your solution a name other than i).

To follow the dynamics of the actual numbers of infected and susceptible people, we can study a **system** of differential equations for i and s (the number of susceptible people)

$$\frac{di}{dt} = \alpha i s - \mu i$$

$$\frac{ds}{dt} = -\alpha i s + \mu i.$$
(2)

To plot the solution of this system, we will use a new command DEplot as follows:

The first line tells Maple to use the DEtools package. The next line defines the differential equation in the form demanded by DEplot. Be careful about the various types of brackets and parentheses. The DEplot command requires at least 4 arguments, the differential equation, a list of variables, the time range, and the initial conditions. Initial conditions are listed in the same order as the variables:  $[0, i_0, s_0]$  tells Maple to start from  $(t = 0, i = i_0, s = s_0)$  and must appear inside curly brackets (and must include an initial condition for t).

**DEplot** can take numerous optional arguments. Included here are limits for the variables i and s (both can run from 0 to 1), and scene, which tells Maple to put t on the x-axis and i on the y-axis. The step size option, stepsize=0.1, forces Maple to use a smaller step size in its solving routine (a fancy version of Euler's method). Without this command, Maple breaks the range into 6 pieces and gives a pretty pathetic graph; by setting the step size to 0.1, we break the region from t = 0 to t = 6 into 60 pieces and get a nice graph. The linecolor option produces a color that prints up better.

## PROBLEMS

- 1. Set  $\alpha = 8$ ,  $\mu = 6$  and  $i_0 = 0.01$  (the initial condition). Graph the solution of equation 1 for  $0 \le t \le 6$ . Describe in words what is happening. What is the fraction of infected people at t = 0, t = 1, t = 2, t = 3 and t = 5? Mark these points on your graph.
- 2. We now compare the solution of equation 1 with the two dimensional system equation 2 using the same parameter values.
  - **a.** Use DEplot to produce a graph of *i* as a function of time for equation 2. Use  $s(0) = s_0 = 0.99$  as your initial condition for *s* (can you see why?). How does the solution compare with 1?
  - **b.** Modify the **scene** command to produce a graph of *s* as a function of time. How is *s* related to *i*? Can you see why?
  - c. Modify the scene command to produce a graph of s as a function of i for equation 2 (a solution in the phase-plane). From the graphs of i and s as functions of t, mark where t = 0, t = 1, t = 2, t = 3 and t = 5 occur in the phase-plane.
- 3. Now consider the following modification of the equations.

$$\frac{di}{dt} = \alpha i s - \mu i 
\frac{ds}{dt} = -\alpha i s + \beta s,$$
(3)

The  $\alpha$  term is the same as above, but  $\mu$  now represents death due to the disease. The  $\beta$  term describes births of new susceptible offspring from susceptible parents.

- **a.** Convince yourself and other relevant people that equation 3 is identical to the predator-prey equations in the book. Which is the predator?
- **b.** Set  $\beta = 1$  and follow the steps in **2**. Can you explain why the dynamics are so completely different from **2**?
- **c.** Draw the nullclines on your figure (unless you are cleverer than us, you'll have to do this by hand).
- **d.** Try the same steps as in **b** with  $\beta = 0.1$ . Describe what happens. Does it remind you of anything? What would happen if  $\beta$  were made very small?