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> read("C:/Users/mudas/Documents/DMB.txt")
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```

This is DMB.txt, A Maple package to explore Dynamical models in Biology (both discrete and continuous) accompanying the class Dynamical Models in Biology, Rutgers University. Taught by Dr. Z. (Doron Zeilbeger)

*The most current version is available on WWW at:
<http://sites.math.rutgers.edu/~zeilberg/tokhniot/DMB.txt> .
Please report all bugs to: DoronZeil at gmail dot com .*

*For general help, and a list of the MAIN functions,
type "Help():". For specific help type "Help(procedure_name);"*

*For a list of the supporting functions type: Help1();
For help with any of them type: Help(ProcedureName);*

*For a list of the functions that give examples of Discrete-time dynamical systems (some famous),
type: HelpDDM());
For help with any of them type: Help(ProcedureName);*

*For a list of the functions continuous-time dynamical systems (some famous) type: HelpCDM());
For help with any of them type: Help(ProcedureName);*

```
> #1
> g := Orbk(3, z, 2 z[1] - z[3], [1, 1, 2], 1000, 1000)
```

```
g :=
[
43466557686937456435688527675040625802564660517371780402481729089536555417949\
05189040387984007925516929592259308032263477520968962323987332247116164299644\
0906533187938298969649928516003704476137795166849228875 ]
```

(1)

(2)

> $h := \text{Orbk}(3, z, 2z[1] - z[3], [1, 1, 2], 999, 999)$
 $h :=$ (3)

[
 26863810024485359386146727202142923967616609318986952340123175997617981700247\
 88168933836965448335656419182785616144335631297667364221035032463485041037768\
 0367334151172899169723197082763985615764450078474174626]

> # Compute g divided by h to get 1.61803398875

>
 #2

> $F := \left[\frac{5}{2}x \cdot (1-x) \cdot \left(1 - \frac{1}{2}x\right) \right]$

$$F := \left[\frac{5x(1-x)\left(1 - \frac{x}{2}\right)}{2} \right]$$
 (4)

> #a)

> $\text{EquP}(F, [x])$
 $\{[0], [1], [2]\}$ (5)

> #b)

> $\text{SEquP}(F, [x])$
 $\{[1.]\}$ (6)

> #c)

> # Since 1 is the stable equilibrium point, it is expected that $t = 100$ would be 1.
 # Check:

> $x1 := \text{dsolve}\left(\left\{\text{diff}(x(t), t) = \frac{5}{2}x(t) \cdot (1-x(t)) \cdot \left(1 - \frac{1}{2}x(t)\right), x(0) = 0.1\right\}, x(t)\right)$

$$x1 := x(t) = \frac{19 e^{\frac{5t}{2}}}{81 \left(-\frac{19 e^{\frac{5t}{2}}}{81} - 1 \right) \left(-\frac{1}{\sqrt{1 + \frac{19 e^{\frac{5t}{2}}}{81}}} - 1 \right)}$$
 (7)

> $\text{eval}(x1, t=100)$

$$x(100) = \frac{19 e^{250}}{81 \left(-\frac{19 e^{250}}{81} - 1 \right) \left(-\frac{1}{\sqrt{1 + \frac{19 e^{250}}{81}}} - 1 \right)}$$
 (8)

> # This evaluates to 1.

>
 #3)

$$\begin{aligned}
 > F1 := \left[\frac{5}{2}x \cdot (1-x) \cdot \left(1 - \frac{1}{2}x\right) \right] \\
 & \qquad \qquad \qquad F1 := \left[\frac{5x(1-x)\left(1 - \frac{x}{2}\right)}{2} \right]
 \end{aligned} \tag{9}$$

$$\begin{aligned}
 > \#a) \\
 > FP(F1, [x]) \\
 & \qquad \qquad \qquad \left\{ [0], \left[\frac{3}{2} - \frac{\sqrt{105}}{10} \right], \left[\frac{3}{2} + \frac{\sqrt{105}}{10} \right] \right\}
 \end{aligned} \tag{10}$$

$$\begin{aligned}
 > \#b) \\
 > SFP(F1, [x]) \\
 & \qquad \qquad \qquad \{ [0.475304923] \}
 \end{aligned} \tag{11}$$

$$\begin{aligned}
 > \#c) \\
 > Orb(F1, [x], [0.1], 1000, 1000) \\
 & \qquad \qquad \qquad [[0.4753049232]]
 \end{aligned} \tag{12}$$

$$\begin{aligned}
 > \#4 \\
 > \#a) \\
 > Orb\left(\left(HW3(u, v, w), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 1, 1\right)\right) \\
 & \qquad \qquad \qquad \left[\left[\frac{1}{4}, \frac{1}{2}, \frac{1}{4}\right]\right]
 \end{aligned} \tag{13}$$

$$\begin{aligned}
 > \# v \text{ represents } Aa. \text{ The proportion of the second generation that would have } Aa \text{ is } 0.5. \\
 > \#b) \\
 > Orb\left(\left(HW3(u, v, w), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 999, 999\right)\right) \\
 & \qquad \qquad \qquad \left[\left[\frac{1}{4}, \frac{1}{2}, \frac{1}{4}\right]\right]
 \end{aligned} \tag{14}$$

$$\begin{aligned}
 > \# v \text{ represents } Aa. \text{ The proportion of the second generation that would have } Aa \text{ is } 0.5. \\
 > \#5 \\
 > A := Matrix([\ [.1, .2, .1], \ [.1, .1, .1], \ [.1, .1, .1]]) \\
 & \qquad \qquad \qquad A := \begin{bmatrix} 0.1 & 0.2 & 0.1 \\ 0.1 & 0.1 & 0.1 \\ 0.1 & 0.1 & 0.1 \end{bmatrix}
 \end{aligned} \tag{15}$$

$$\begin{aligned}
 > \# Change the matrix position M[1][2] to 0.2 to represent AA females twice as likely to mate with Aa male \\
 > \#a) \\
 > Orb\left(HW3g(u, v, w, A), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 1, 1\right) \\
 & \qquad \qquad \qquad [[0.2750000001, 0.5000000001, 0.2250000000]]
 \end{aligned} \tag{16}$$

$$\begin{aligned} &> F := SIRS(s, i, 0.099999, 0.5, 100, 1000) \\ & \quad F := [-0.099999 s i + 500.0 - 0.5 s - 0.5 i, 0.099999 s i - 100 i] \end{aligned} \quad (26)$$

$$\begin{aligned} &> SEquP(F, [s, i]) \\ & \quad \{[1000., 0.]\} \end{aligned} \quad (27)$$

$$\begin{aligned} &> F := SIRS(s, i, 0.100001, 0.5, 100, 1000) \\ & \quad F := [-0.100001 s i + 500.0 - 0.5 s - 0.5 i, 0.100001 s i - 100 i] \end{aligned} \quad (28)$$

$$\begin{aligned} &> SEquP(F, [s, i]) \\ & \quad \{[999.9900001, 0.00004975074627]\} \end{aligned} \quad (29)$$

> # The number of infected individuals is still 0 close at 0.1 (shown by $\beta = 0.099999$), and changes to non-zero once past 0.1 (shown by $\beta = 0.100001$). However, there are no stable fixed points at 0.1 itself. Therefore, $\beta = 0.1$ is the cutoff for non-zero number of infected people.

> #8

> #a)

$$\begin{aligned} &> F := GeneNet(0, 1, 0.2, 2, m1, m2, m3, p1, p2, p3) \\ F & := \left[-m1 + \frac{1}{p3^2 + 1}, -m2 + \frac{1}{p1^2 + 1}, -m3 + \frac{1}{p2^2 + 1}, -0.2 p1 + 0.2 m1, -0.2 p2 \right. \\ & \quad \left. + 0.2 m2, -0.2 p3 + 0.2 m3 \right] \end{aligned} \quad (30)$$

$$\begin{aligned} &> SEquP(F, [m1, m2, m3, p1, p2, p3]) \\ & \quad [0.6823278038, 0.6823278038, 0.6823278038, 0.6823278038, 0.6823278038, 0.6823278038] \end{aligned} \quad (31)$$

> # The horizontal asymptote is at 0.6823278038. By definition, the height of the asymptote is the value of the stable equilibrium point.

> #b)

$$\begin{aligned} &> F := GeneNet(0, 3, 0.2, 2, m1, m2, m3, p1, p2, p3) \\ F & := \left[-m1 + \frac{3}{p3^2 + 1}, -m2 + \frac{3}{p1^2 + 1}, -m3 + \frac{3}{p2^2 + 1}, -0.2 p1 + 0.2 m1, -0.2 p2 \right. \\ & \quad \left. + 0.2 m2, -0.2 p3 + 0.2 m3 \right] \end{aligned} \quad (32)$$

$$\begin{aligned} &> SEquP(F, [m1, m2, m3, p1, p2, p3]) \\ & \quad \{[1.213411663, 1.213411663, 1.213411663, 1.213411663, 1.213411663, 1.213411663]\} \end{aligned} \quad (33)$$

> # The horizontal asymptote is at 1.213411663. By definition, the height of the asymptote is the value of the stable equilibrium point.

> #c)

$$\begin{aligned} &> F := GeneNet(0, 7.393, 0.2, 2, m1, m2, m3, p1, p2, p3) \\ F & := \left[-m1 + \frac{7.393}{p3^2 + 1}, -m2 + \frac{7.393}{p1^2 + 1}, -m3 + \frac{7.393}{p2^2 + 1}, -0.2 p1 + 0.2 m1, -0.2 p2 \right. \\ & \quad \left. + 0.2 m2, -0.2 p3 + 0.2 m3 \right] \end{aligned} \quad (34)$$

$$\begin{aligned} &> SEquP(F, [m1, m2, m3, p1, p2, p3]) \\ & \quad \{[7.393, 7.393, 7.393, 7.393, 7.393, 7.393]\} \end{aligned} \quad (35)$$

$$\{[1.777450148, 1.777450148, 1.777450148, 1.777450148, 1.777450148, 1.777450148]\} \quad (35)$$

> # There are no stable equilibrium points at $a = 7.394$. However, at $a = 7.393$, there is stable equilibrium. Therefore, there is no longer a stable equilibrium starting $a = 7.394$.

> # Check $a = 7.393$:

> $F := \text{GeneNet}(0, 7.393, 0.2, 2, m1, m2, m3, p1, p2, p3)$

$$F := \left[-m1 + \frac{7.393}{p3^2 + 1}, -m2 + \frac{7.393}{p1^2 + 1}, -m3 + \frac{7.393}{p2^2 + 1}, -0.2 p1 + 0.2 m1, -0.2 p2 + 0.2 m2, -0.2 p3 + 0.2 m3 \right] \quad (36)$$

> $\text{SEquP}(F, [m1, m2, m3, p1, p2, p3])$

$$\{[1.777450148, 1.777450148, 1.777450148, 1.777450148, 1.777450148, 1.777450148]\} \quad (37)$$

> # This shows that there is stable equilibrium at $a = 7.393$, but not at $a = 7.394$.

>

> #9

> $F := \text{ChemoStat}(N, C, 2.5, 2.7)$

$$F := \left[\frac{2.5 CN}{C + 1} - N, -\frac{CN}{C + 1} - C + 2.7 \right] \quad (38)$$

> $\text{SEquP}(F, [N, C])$

$$\{[5.083333333, 0.666666667]\} \quad (39)$$

> #a

> # The value of bacterial population density after a long time will be 5.08333333, since it is a stable fixed point.

> #b

> # The value of nutrient concentration after a long time will be 0.66666667, since it is a stable fixed point.

>

> #10

> $A := \text{Matrix}\left(\left[[0.2, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1], [0.1, 0.2, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1], \right.$

$$\left. \begin{aligned} & [0.1, 0.1, 0.2, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1], \left[\frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, 0.4, \frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, \right. \\ & \left. \frac{0.6}{8}, \frac{0.6}{8} \right], \left[\frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, 0.4, \frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8} \right], \left[\frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8}, \right. \\ & \left. \frac{0.6}{8}, 0.4, \frac{0.6}{8}, \frac{0.6}{8}, \frac{0.6}{8} \right], \left[\frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \right. \\ & \left. \frac{0.4}{8}, 0.6, \frac{0.4}{8}, \frac{0.4}{8} \right], \left[\frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \frac{0.4}{8}, \right. \\ & \left. \frac{0.4}{8}, 0.6 \right] \end{aligned} \right)$$

$A := [[0.2, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1],$

$[0.1, 0.2, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1],$

$[0.1, 0.1, 0.2, 0.1, 0.1, 0.1, 0.1, 0.1, 0.1],$

(40)


```
0.102564102564107, 0.102564102564107, 0.153846153846160, 0.153846153846160,  
0.153846153846160 ]]
```

```
> #a
```

```
> # The probability of a surfer being on web-page 1 is 0.0769230769230801, or approximately  
7.69 %
```

```
> #b
```

```
> # The probability of a surfer being on web-page 9 is 0.153846153846160, or approximately  
15.385 %
```

```
>
```