

> read `Users/Deven/Desktop/Fall 2021/Dynamic Models of Biology/DMB.txt`
First Written: Nov. 2021

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)

> # Deven Singh
Assignment 21
#OK TO POST

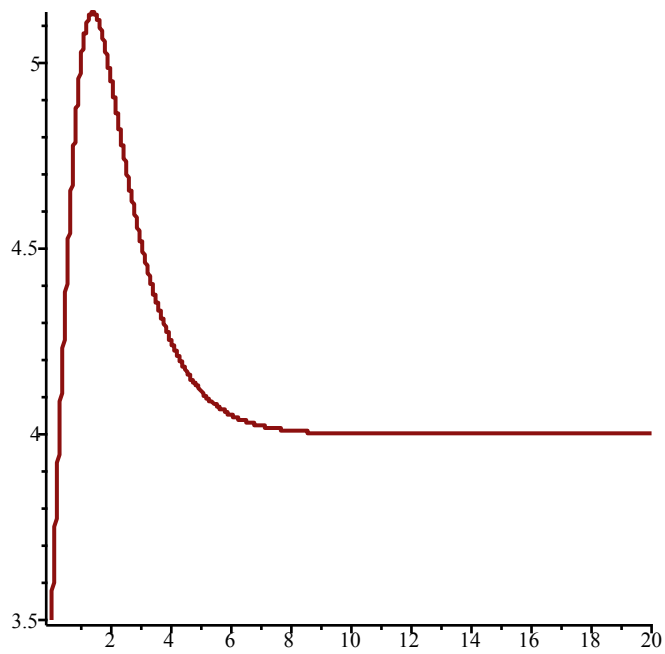
> FI := ChemoStat(N, C, 2, 3);

$$FI := \left[\frac{2CN}{C+1} - N, -\frac{CN}{C+1} - C + 3 \right] \quad (2)$$

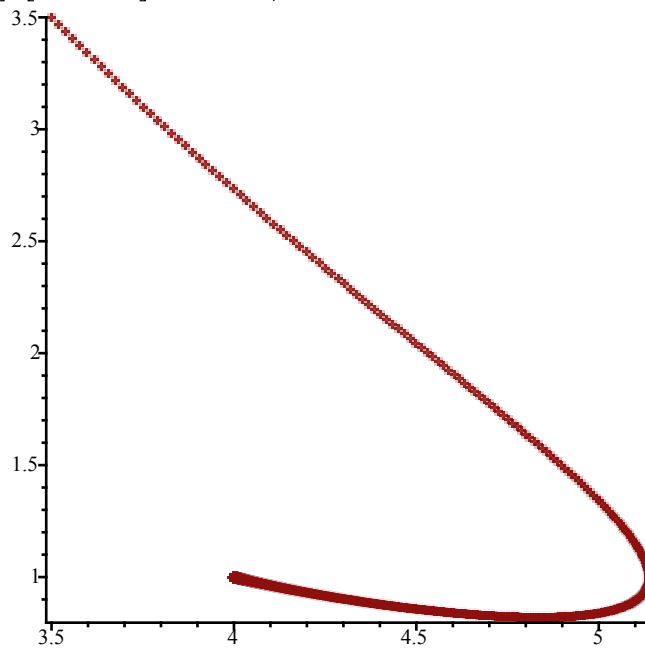
> SEquP(FI, [N, C]);

{[4., 1.]} (3)

> TimeSeries(FI, [N, C], [3.5, 3.5], .01, 20, 1);



> *PhaseDiag*(*F1*, [*N*, *C*], [3.5, 3.5], 0.01, 20);



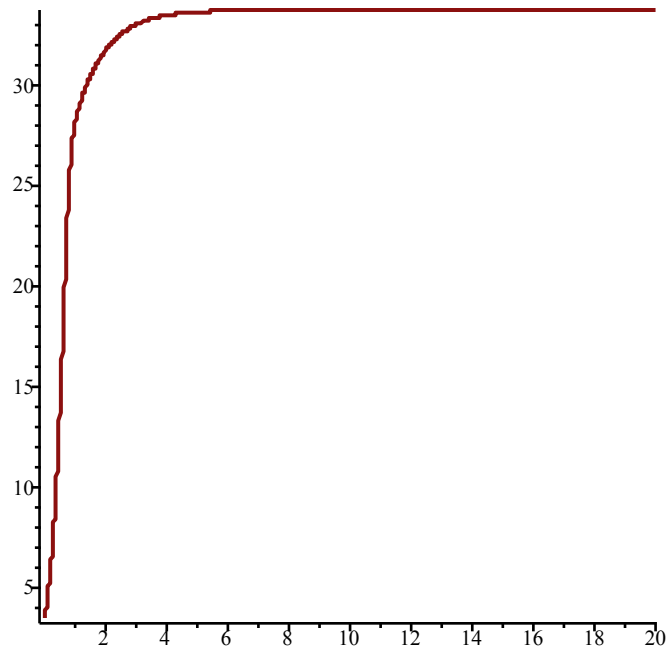
> *F2* := *ChemoStat*(*N*, *C*, 5, 7);

$$F2 := \left[\frac{5CN}{C+1} - N, -\frac{CN}{C+1} - C + 7 \right] \quad (4)$$

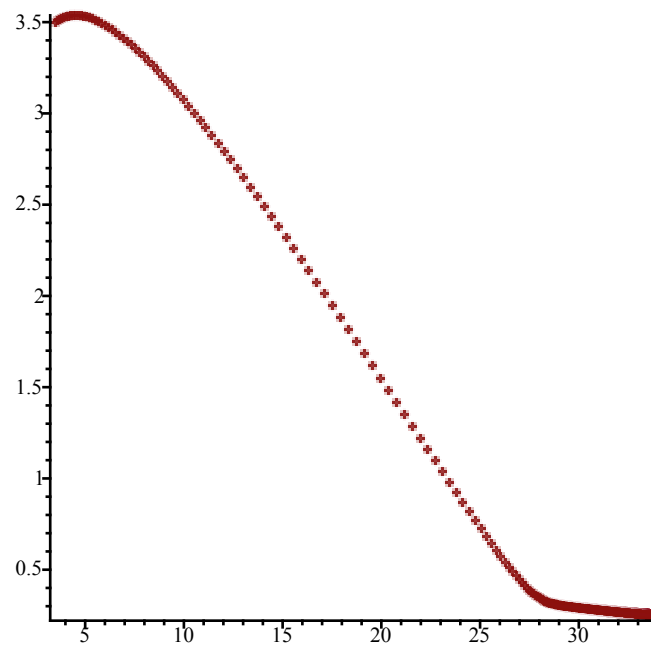
> *SEquP*(*F2*, [*N*, *C*]);

$$\{ [33.75000000, 0.2500000000] \} \quad (5)$$

> *TimeSeries*(*F2*, [*N*, *C*], [3.5, 3.5], .01, 20, 1);



> *PhaseDiag*(*F2*, [*N*, *C*], [3.5, 3.5], 0.01, 20);



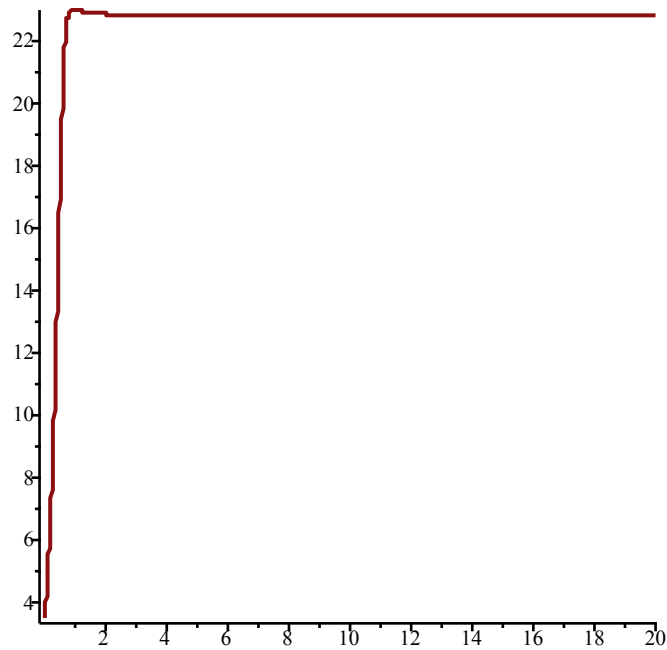
> *F3* := *ChemoStat*(*N*, *C*, 6, 4);

$$F3 := \left[\frac{6CN}{C+1} - N, -\frac{CN}{C+1} - C + 4 \right] \quad (6)$$

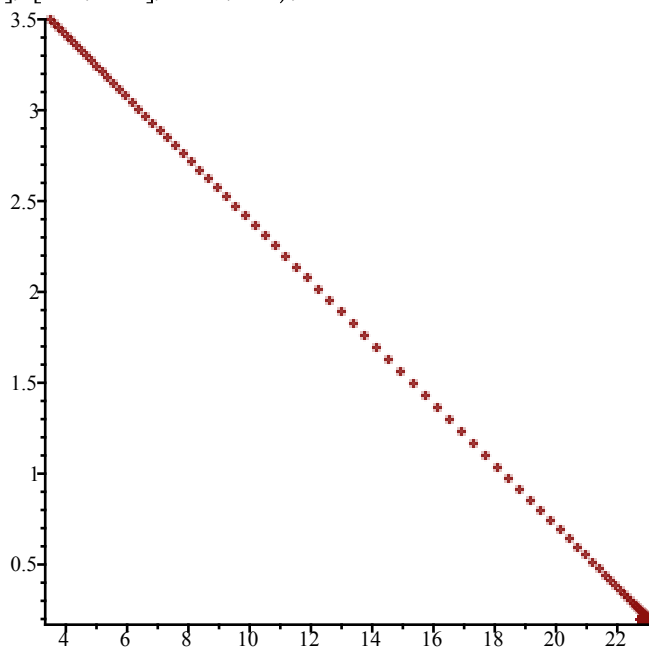
> *SEquP*(*F3*, [*N*, *C*]);

$$\{ [22.80000000, 0.2000000000] \} \quad (7)$$

> *TimeSeries*(*F3*, [*N*, *C*], [3.5, 3.5], .01, 20, 1);



> `PhaseDiag(F3, [N, C], [3.5, 3.5], 0.01, 20);`



> `Help(GeneNet);`

GeneNet(a0,a,b,n,m1,m2,m3,p1,p2,p3): The continuous-time dynamical system, with quantities $m1, m2, m3, p1, p2, p3$, due to M. Elowitz and S. Leibler

described in the Ellner-Guckenheimer book, Eq. (4.1) (chapter 4, p. 112)

and parameters a_0 (called alpha_0 there), a (called alpha there), b (called beta there) and n . Try:

*`GeneNet(0,0.5,0.2,2,m1,m2,m3,p1,p2,p3);` **(8)***

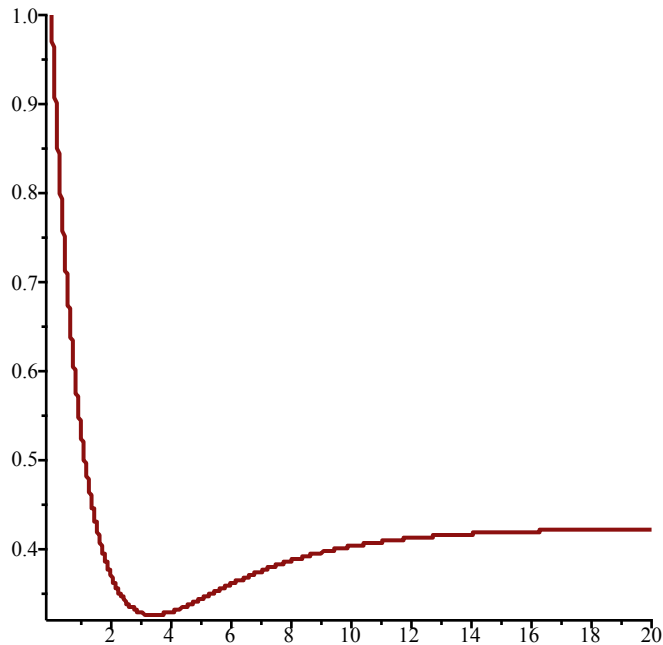
> `G1 := GeneNet(0, 0.5, 0.2, 2, m1, m2, m3, p1, p2, p3);`

$$G1 := \left[-m1 + \frac{0.5}{p3^2 + 1}, -m2 + \frac{0.5}{p1^2 + 1}, -m3 + \frac{0.5}{p2^2 + 1}, -0.2 p1 + 0.2 m1, -0.2 p2 \right] \quad \mathbf{(9)}$$

$+ 0.2 m_2, -0.2 p_3 + 0.2 m_3]$

> *SEquP*(*G1*, [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*]);
 {[0.4238537991, 0.4238537991, 0.4238537991, 0.4238537991, 0.4238537991, 0.4238537991]} (10)

> *TimeSeries*(*G1*, [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [1, 1, 1, 1, 1, 1], .01, 20, 1);



> *Help*(*Lotka*);

Lotka(*r1*,*k1*,*r2*,*k2*,*b12*,*b21*,*N1*,*N2*): *The Lotka-Volterra continuous-time dynamical system, Eqs.*

(9a),(9b) (p. 224, section 6.3) of *Edelstein-Keshet*

with populations *N1*, *N2*, and parameters *r1*,*r2*,*k1*,*k2*, *b12*, *b21* (called there *beta_12* and *beta_21*)

Try:

Lotka(*r1*,*k1*,*r2*,*k2*,*b12*,*b21*,*N1*,*N2*);

Lotka(1,2,2,3,1,2,*N1*,*N2*);

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> *H1* := *Lotka*(1, 2, 2, 3, 1, 2, *N1*, *N2*);

$$H1 := \left[\frac{N1 (2 - N1 - N2)}{2}, \frac{2 N2 (3 - N2 - 2 N1)}{3} \right]$$

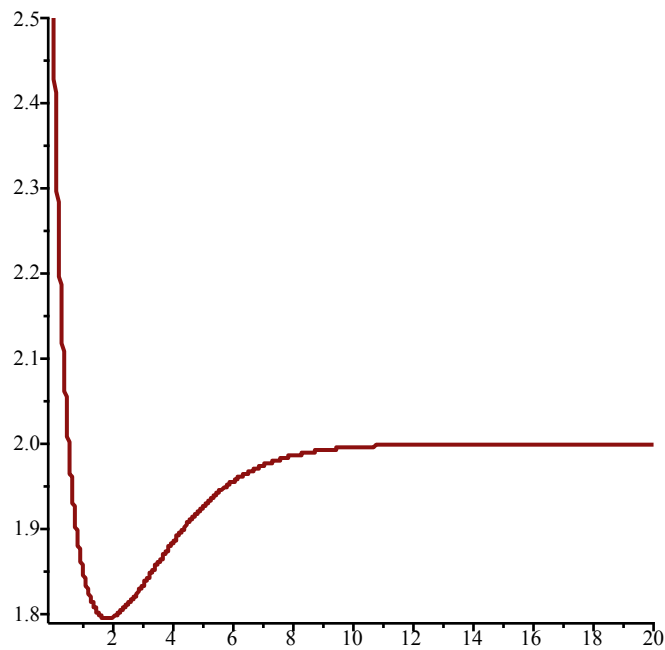
(12)

> *SEquP*(*H1*, [*N1*, *N2*]);

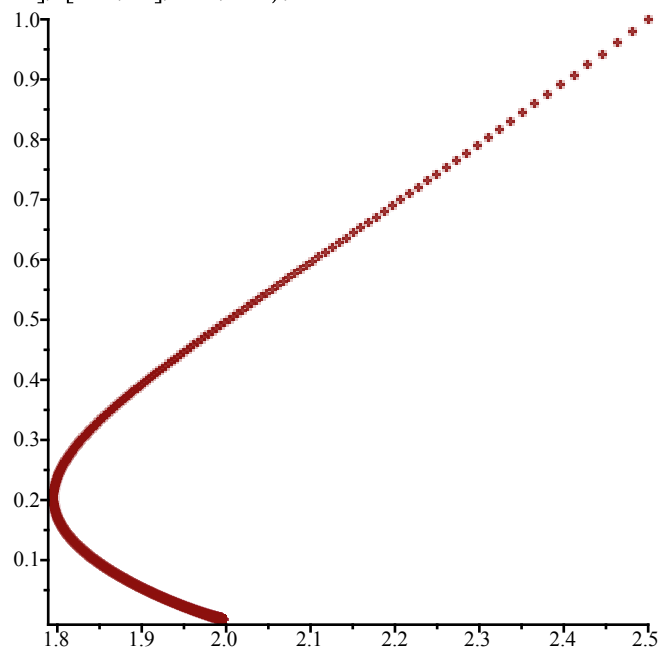
{[0., 3.], [2., 0.]}

(13)

> *TimeSeries*(*H1*, [*N1*, *N2*], [2.5, 1], .01, 20, 1);



> `PhaseDiag(HI, [N1, N2], [2.5, 1], .01, 10);`



> `Help(Volterra);`

Volterra(a,b,c,d,x,y): The (simple, original) Volterra predator-prey continuous-time dynamical system with parameters a,b,c,d

Given by Eqs. (7a) (7b) in Edelstein-Keshet p. 219 (section 6.2).

a,b,c,d may be symbolic or numeric

Try:

Volterra(a,b,c,d,x,y);

Volterra(1,2,3,4,x,y);

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> `A1 := Volterra(1, 2, 3, 4, x, y);`

(15)

$$AI := [-2xy + x, 4xy - 3y] \quad (15)$$

> SEquP(AI, [x, y]);

$$\emptyset \quad (16)$$

> Help(VolterraM);

VolterraM(a,b,c,d,x,K,y): The MODIFIED Volterra predator-prey continuous-time dynamical system with parameters a,b,c,d,K

Given by Eqs. (8a) (8b) in Edelstein-Keshet p. 220 (section 6.2).

a,b,c,d,K may be symbolic or numeric

Try:

VolterraM(a,b,c,d,K,x,y);

VolterraM(1,2,3,4,3,x,y);

(17)

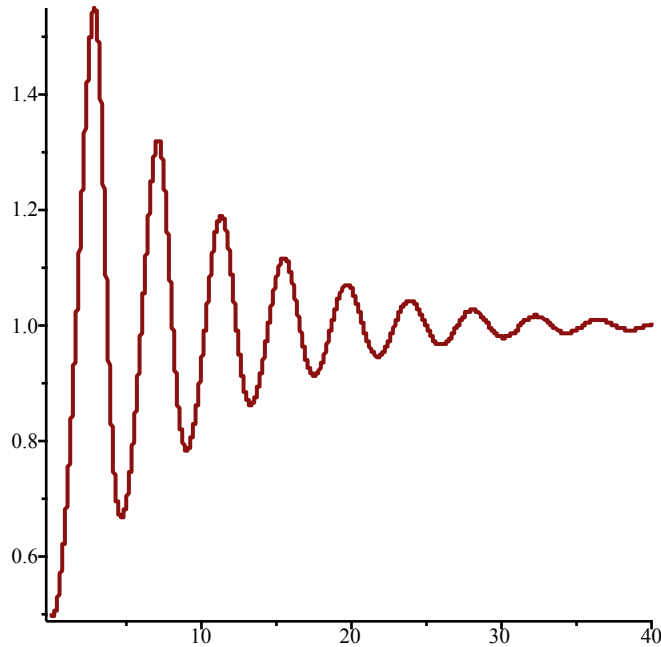
> BI := VolterraM(1, 2, 3, 4, 3, x, y);

$$BI := \left[x \left(1 - \frac{x}{4} \right) - 2xy, 3xy - 3y \right] \quad (18)$$

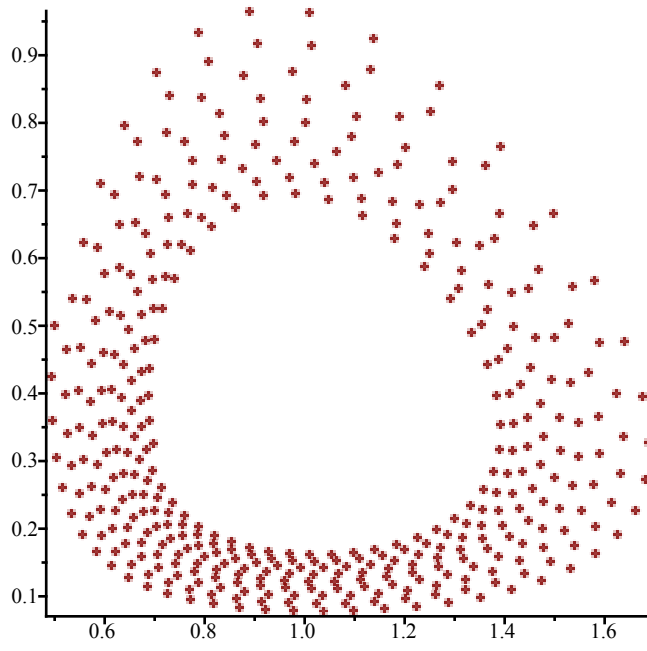
> SEquP(BI, [x, y]);

$$\{[1., 0.3750000000]\} \quad (19)$$

> TimeSeries(BI, [x, y], [0.5, 0.5], .01, 40, 1);



> PhaseDiag(BI, [x, y], [0.5, 0.5], 0.1, 40);



>