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> #OK to post homework
#Shreya Ghosh, 11-15-2021, Assignment 21
> read "/Users/shreyaghosh/Documents/DMB.txt"
First Written: Nov. 2021
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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 Zeilberger)

*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());

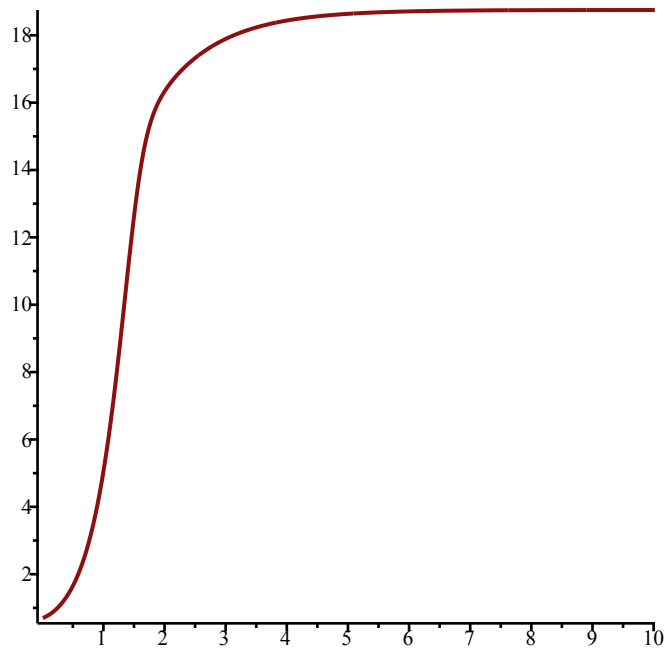
For help with any of them type: Help(ProcedureName);

(1)

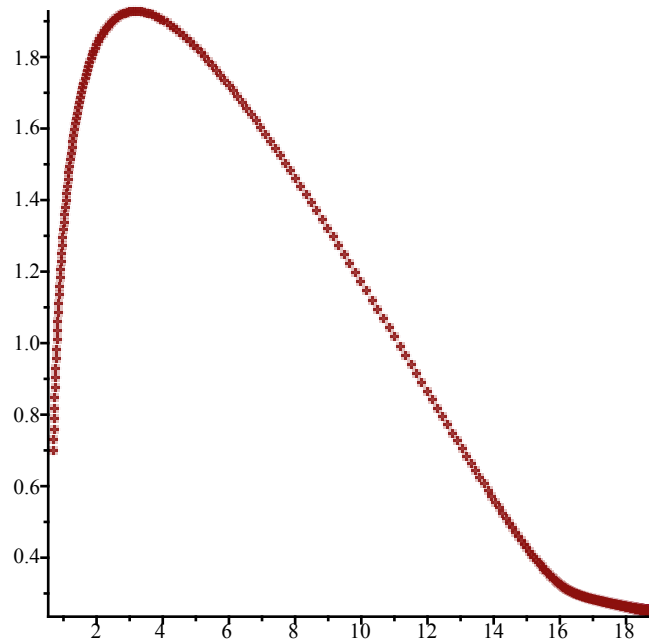
```
> #ChemoStat
> F := ChemoStat(N, C, 5, 4)
```

$$F := \left[\frac{5CN}{C+1} - N, -\frac{CN}{C+1} - C + 4 \right] \quad (2)$$

```
> TimeSeries(F, [N, C], [0.7, 0.7], 0.01, 10, 1)
```



> *PhaseDiag*(*F*, [*N*, *C*], [0.7, 0.7], 0.01, 10)



> *SEquP*(*F*, [*N*, *C*])

{[18.75000000, 0.2500000000]}

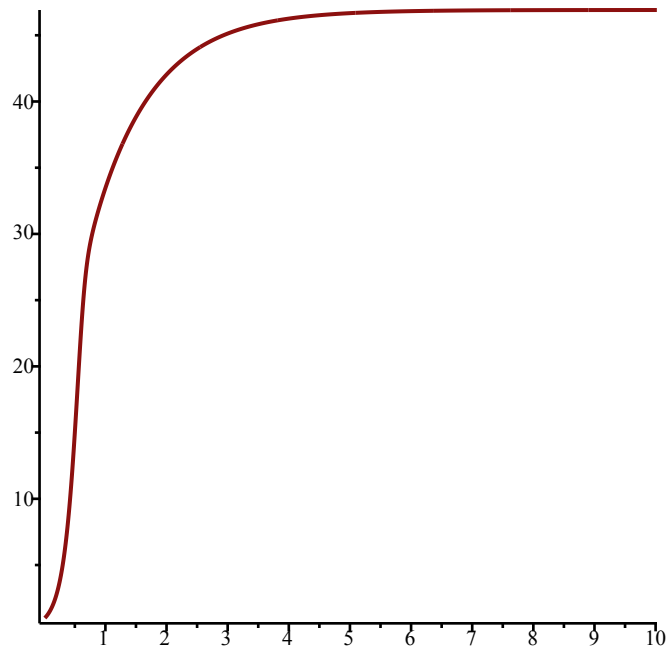
(3)

> *F* := *ChemoStat*(*N*, *C*, 12, 4)

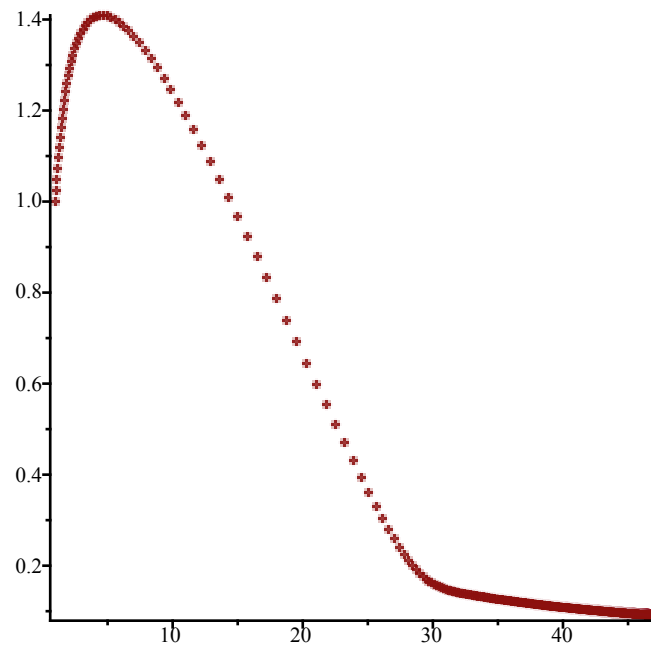
$$F := \left[\frac{12CN}{C+1} - N, -\frac{CN}{C+1} - C + 4 \right]$$

(4)

> *TimeSeries*(*F*, [*N*, *C*], [1, 1], 0.01, 10, 1)



> PhaseDiag(F, [N, C], [1, 1], 0.01, 10)



> SEquP(F, [N, C])

{[46.90909091, 0.090909091]}

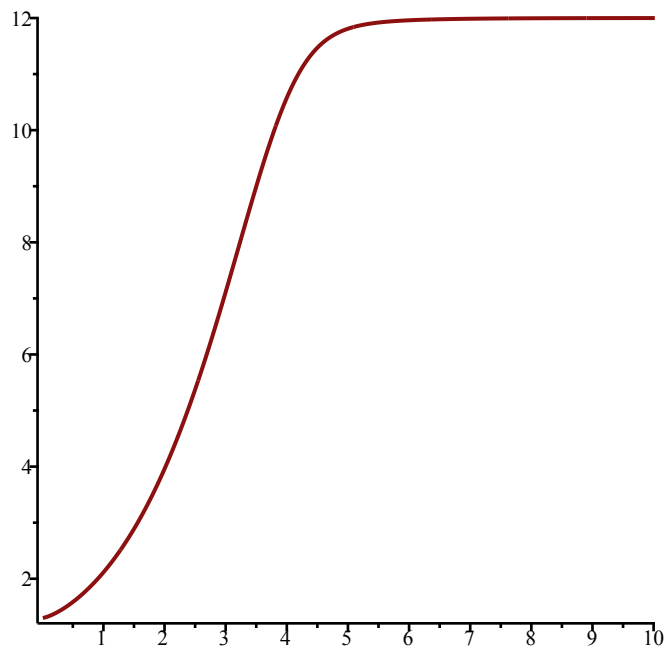
(5)

> F := ChemoStat(N, C, 2, 7)

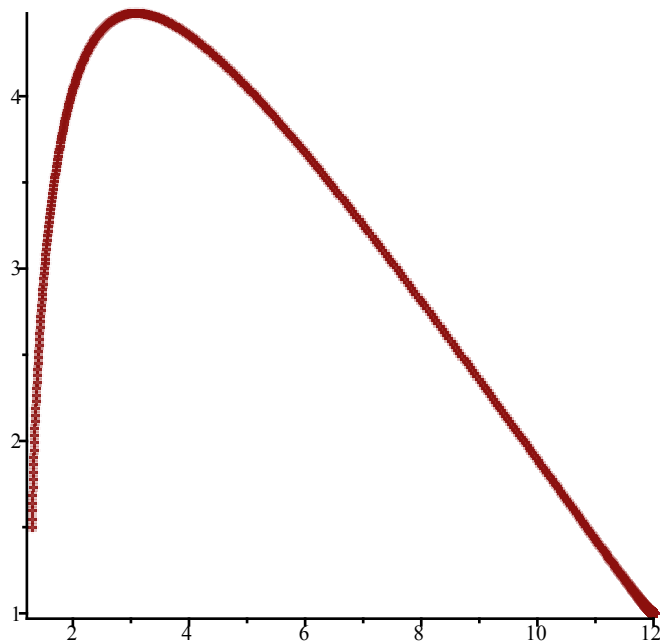
$$F := \left[\frac{2CN}{C+1} - N, -\frac{CN}{C+1} - C + 7 \right]$$

(6)

> TimeSeries(F, [N, C], [1.3, 1.5], 0.01, 10, 1)



> PhaseDiag(F, [N, C], [1.3, 1.5], 0.01, 10)



> SEquP(F, [N, C])

{[12., 1.]}

(7)

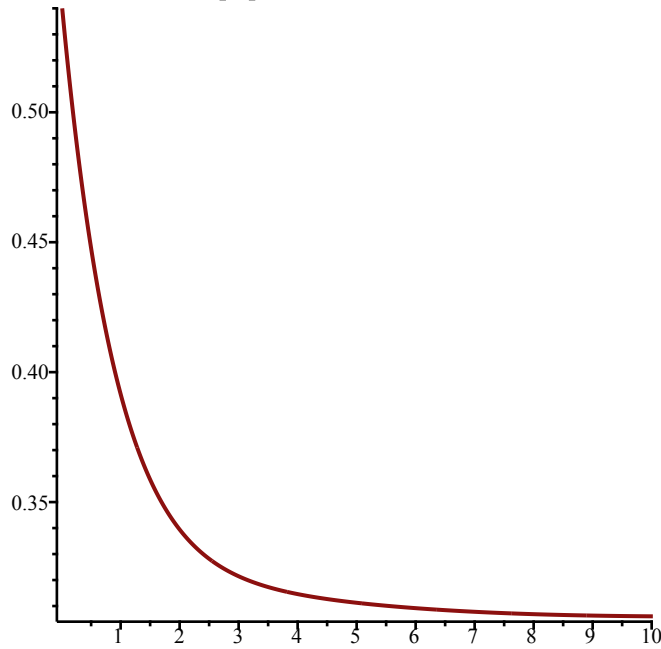
> #The stable equilibrium points for ChemoStat coincide with the horizontal asymptotes found in the time series

> #GeneNet

> F := GeneNet(0, 0.4, 0.6, 1, m1, m2, m3, p1, p2, p3)

$$F := \begin{bmatrix} -m1 + \frac{0.4}{1 + p3}, -m2 + \frac{0.4}{1 + p1}, -m3 + \frac{0.4}{1 + p2}, -0.6 p1 + 0.6 m1, -0.6 p2 + 0.6 m2, \\ -0.6 p3 + 0.6 m3 \end{bmatrix} \quad (8)$$

> TimeSeries(F, [m1, m2, m3, p1, p2, p3], [0.54, 0.37, 0.28, 0.99, 0.64, 0.33], 0.01, 10, 1)

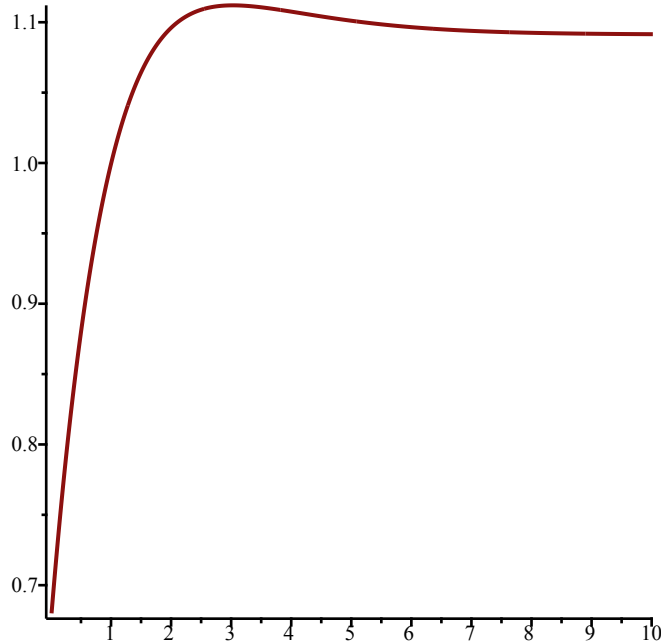


> SEquP(F, [m1, m2, m3, p1, p2, p3])
 {[0.3062257748, 0.3062257748, 0.3062257748, 0.3062257748, 0.3062257748, 0.3062257748]} (9)

> F := GeneNet(1, 0.2, 0.74, 2, m1, m2, m3, p1, p2, p3)

$$F := \left[-m1 + \frac{0.2}{p3^2 + 1} + 1, -m2 + \frac{0.2}{p1^2 + 1} + 1, -m3 + \frac{0.2}{p2^2 + 1} + 1, -0.74 p1 \right. \\ \left. + 0.74 m1, -0.74 p2 + 0.74 m2, -0.74 p3 + 0.74 m3 \right] \quad (10)$$

> TimeSeries(F, [m1, m2, m3, p1, p2, p3], [0.68, 0.44, 0.22, 0.36, 0.19, 0.14], 0.01, 10, 1)

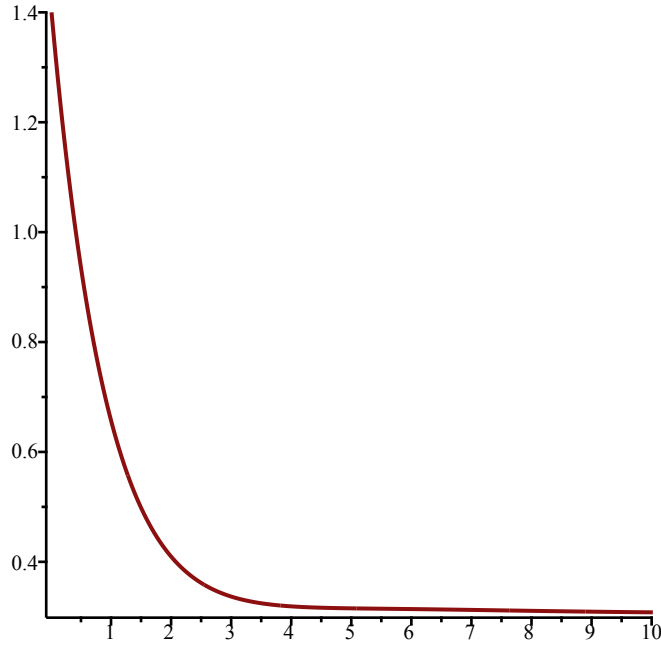


> SEquP(F, [m1, m2, m3, p1, p2, p3])
 {[1.091286433, 1.091286433, 1.091286433, 1.091286433, 1.091286433, 1.091286433]} (11)

> $F := \text{GeneNet}(0, 0.4, 0.66, 1, m1, m2, m3, p1, p2, p3)$

$$F := \left[-m1 + \frac{0.4}{1 + p3}, -m2 + \frac{0.4}{1 + p1}, -m3 + \frac{0.4}{1 + p2}, -0.66 p1 + 0.66 m1, -0.66 p2 + 0.66 m2, -0.66 p3 + 0.66 m3 \right] \quad (12)$$

> $\text{TimeSeries}(F, [m1, m2, m3, p1, p2, p3], [1.4, 1.66, 0.12, 0.72, 0.88, 1.1], 0.01, 10, 1)$



> $\text{SEquP}(F, [m1, m2, m3, p1, p2, p3])$
 $\{ [0.3062257748, 0.3062257748, 0.3062257748, 0.3062257748, 0.3062257748, 0.3062257748] \} \quad (13)$

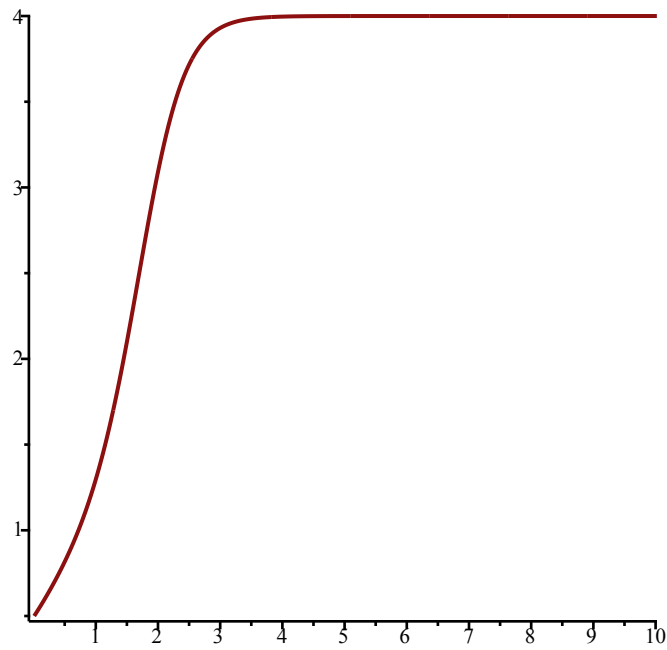
> *#The stable equilibrium points for GeneNet coincide with the horizontal asymptotes found in the time series. The phase diagrams cannot be drawn because the function is not in R2*

> *#Lotka*

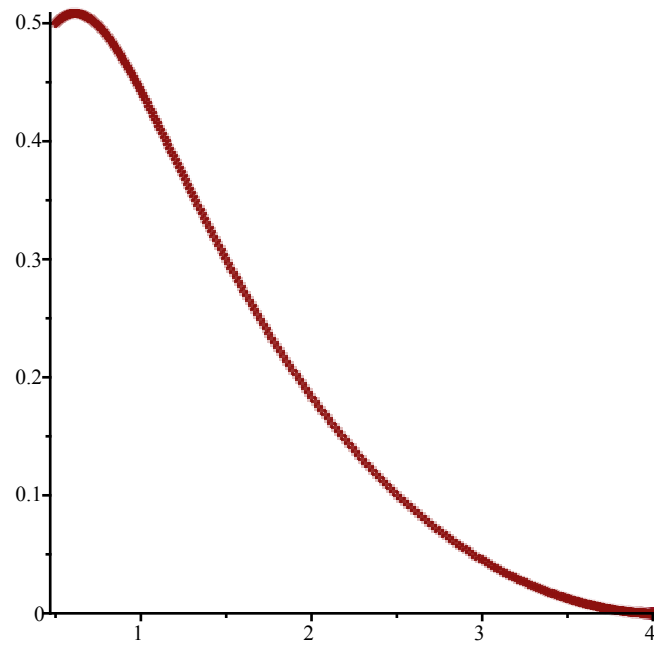
> $F := \text{Lotka}(3, 4, 1, 6, 4, 9, N1, N2)$

$$F := \left[\frac{3 N1 (4 - N1 - 4 N2)}{4}, \frac{N2 (6 - N2 - 9 N1)}{6} \right] \quad (14)$$

> $\text{TimeSeries}(F, [N1, N2], [0.5, 0.5], 0.01, 10, 1)$



> PhaseDiag(F, [N1, N2], [0.5, 0.5], 0.01, 10)



> SEquP(F, [N1, N2])

$$\{[0., 6.], [4., 0.]\}$$

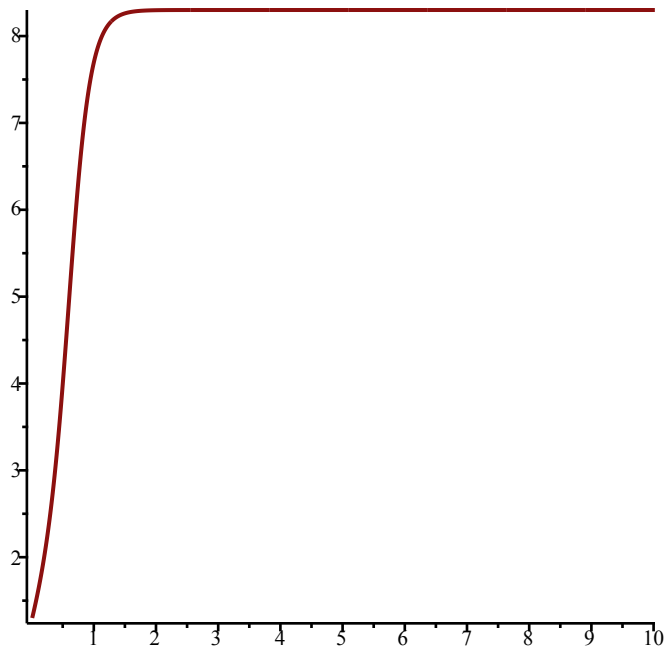
(15)

> F := Lotka(5.5, 8.3, 4, 9, 2.3, 6, N1, N2)

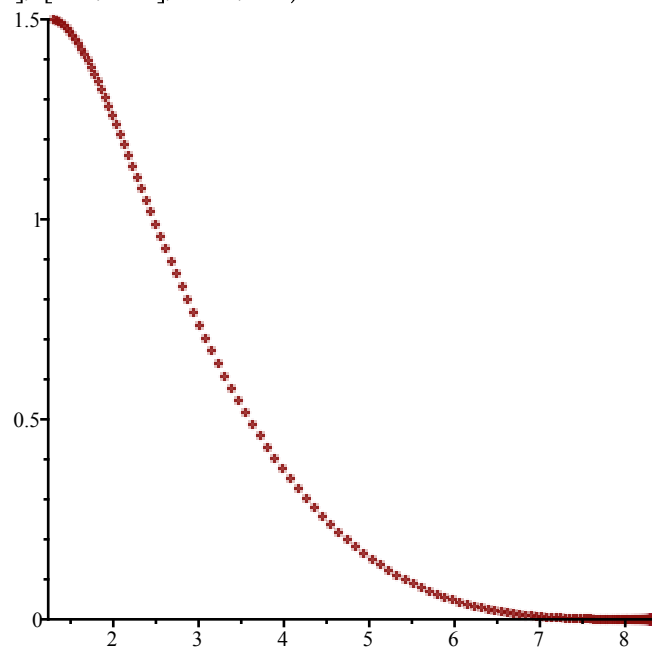
$$F := \left[0.6626506024 N1 (8.3 - N1 - 2.3 N2), \frac{4 N2 (9 - N2 - 6 N1)}{9} \right]$$

(16)

> TimeSeries(F, [N1, N2], [1.3, 1.5], 0.01, 10, 1)



> PhaseDiag(F, [N1, N2], [1.3, 1.5], 0.01, 10)



> SEquP(F, [N1, N2])

{[0., 9.], [8.300000000, 0.]}

(17)

> #The stable equilibrium points for Lotka coincide with the horizontal asymptotes found in the time series

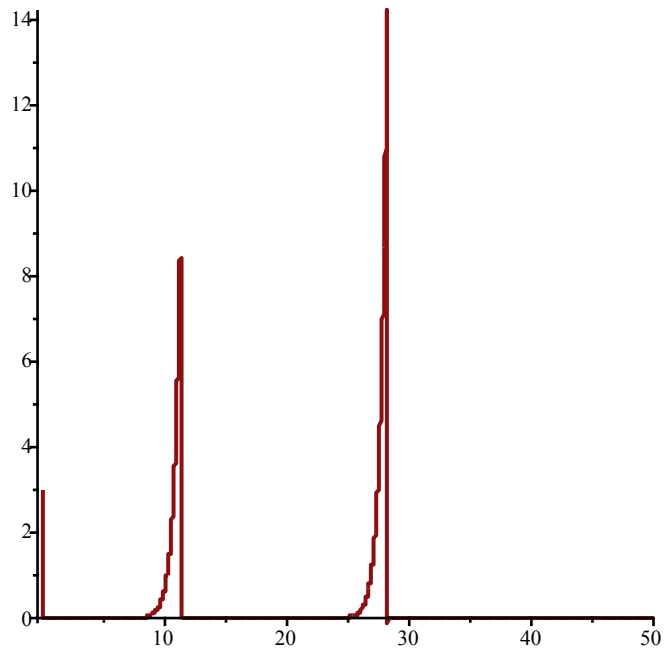
> #Volterra

> F := Volterra(2, 6, 3, 8, x, y)

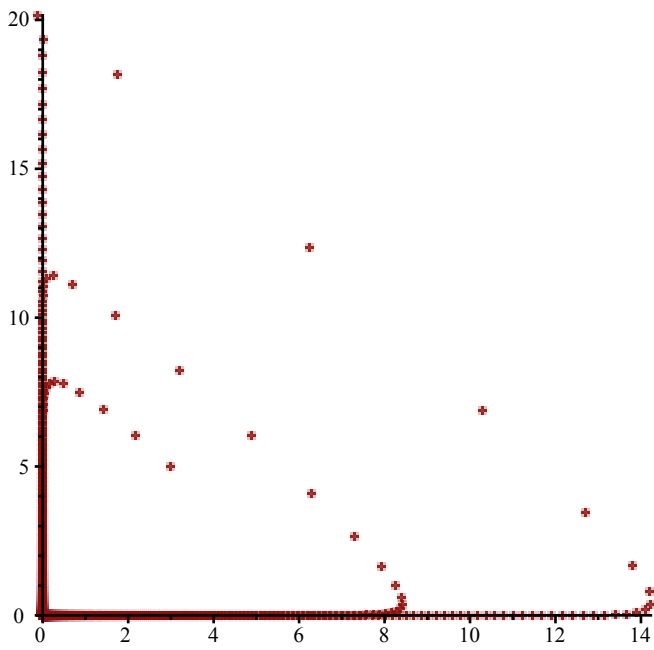
F := [-6xy + 2x, 8xy - 3y]

(18)

> TimeSeries(F, [x, y], [3, 5], 0.01, 50, 1)



> *PhaseDiag*(*F*, [*x*, *y*], [3, 5], 0.01, 50)



> *SEquP*(*F*, [*x*, *y*])

\emptyset

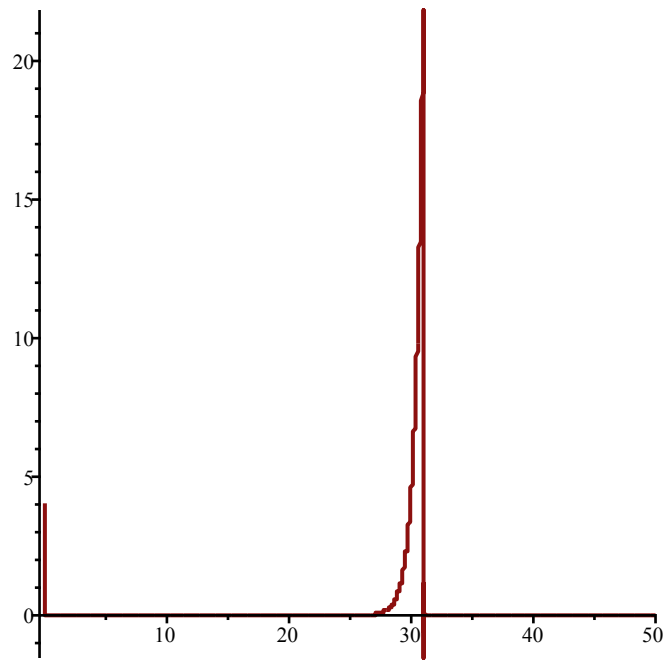
(19)

> *F* := *Volterra*(1.6, 3.5, 2.6, 7.3, *x*, *y*)

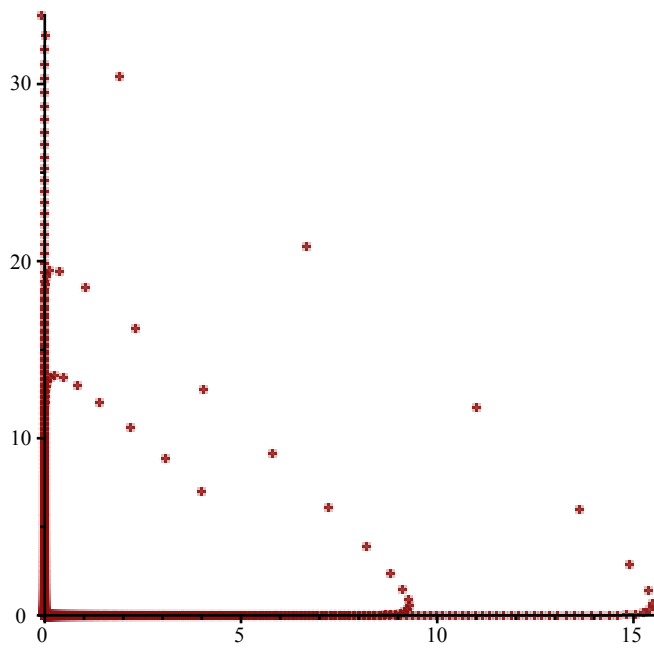
F := [1.6 *x* - 3.5 *x* *y*, -2.6 *y* + 7.3 *x* *y*]

(20)

> *TimeSeries*(*F*, [*x*, *y*], [4, 17], 0.01, 50, 1)



> *PhaseDiag*(*F*, [*x*, *y*], [4, 7], 0.01, 50, 1)



> *SEquP*(*F*, [*x*, *y*])

\emptyset

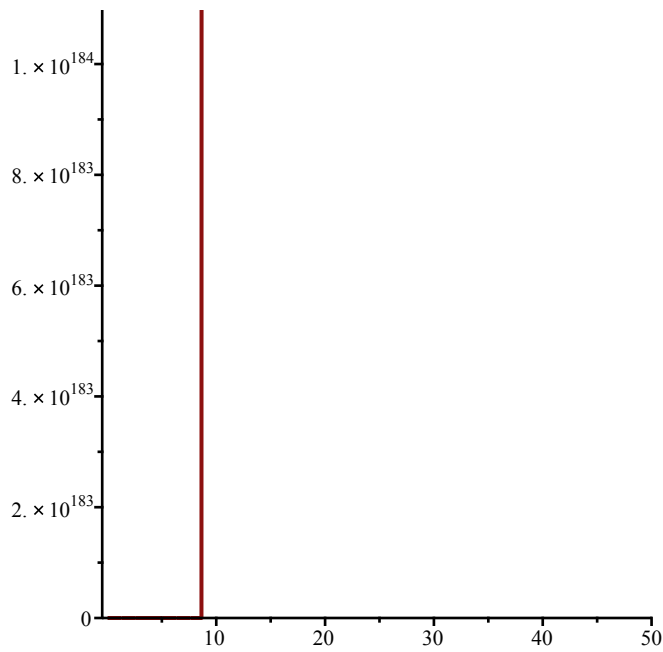
(21)

> *F* := *Volterra*(7.4, 3.2, 8.2, 4.4, *x*, *y*)

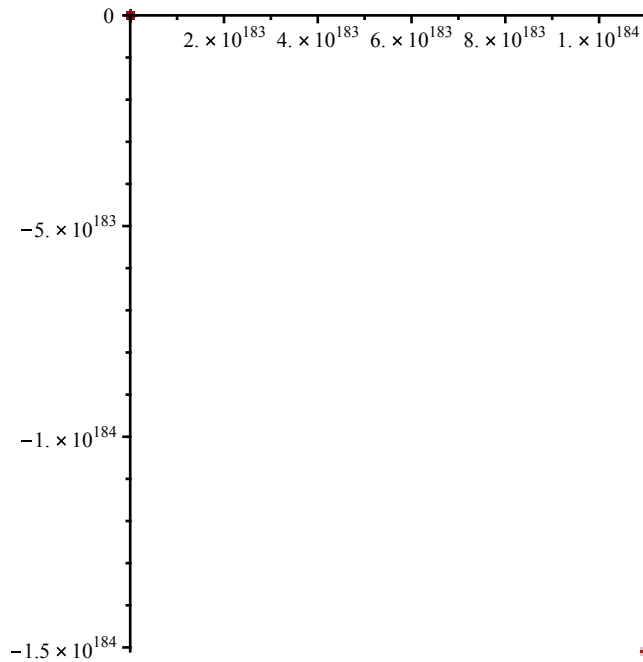
F := [7.4 *x* - 3.2 *x y*, -8.2 *y* + 4.4 *x y*]

(22)

> *TimeSeries*(*F*, [*x*, *y*], [14, 17], 0.01, 50, 1)



> PhaseDiag(F, [x, y], [14, 17], 0.01, 50)



> SEquP(F, [x, y])

\emptyset

(23)

> #The Volterra system does not have stable equilibria

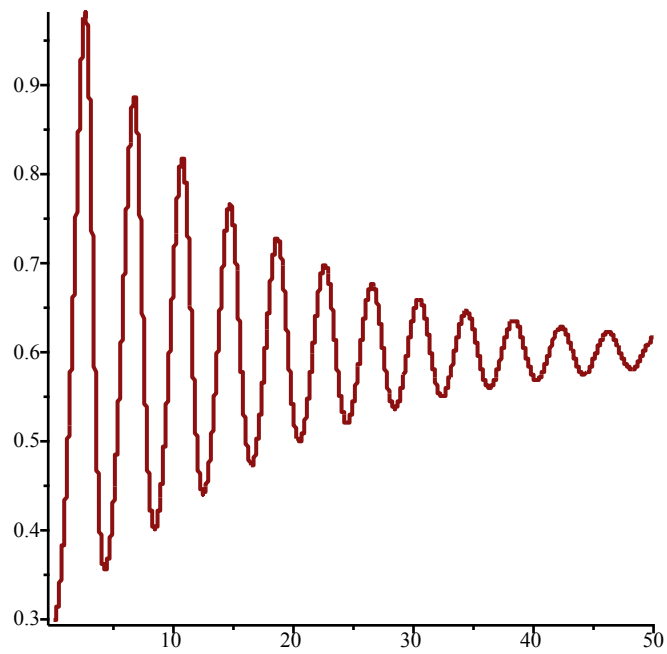
> #VolterraM

> F := VolterraM(1, 2, 3, 4, 5, x, y)

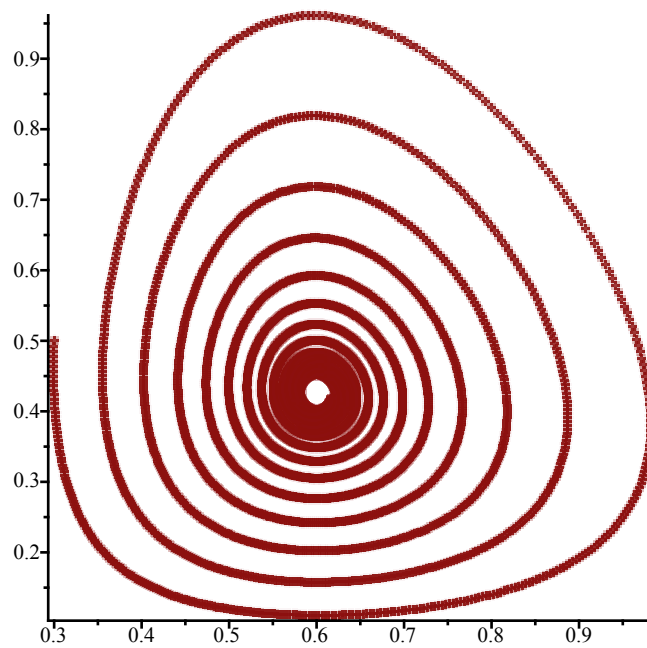
$$F := \left[x \left(1 - \frac{x}{4} \right) - 2xy, 5xy - 3y \right]$$

(24)

> TimeSeries(F, [x, y], [.3, .5], 0.01, 50, 1)



> *PhaseDiag*(*F*, [*x*, *y*], [.3, .5], 0.01, 50)



> *SEquP*(*F*, [*x*, *y*])

{[0.6000000000, 0.4250000000]}

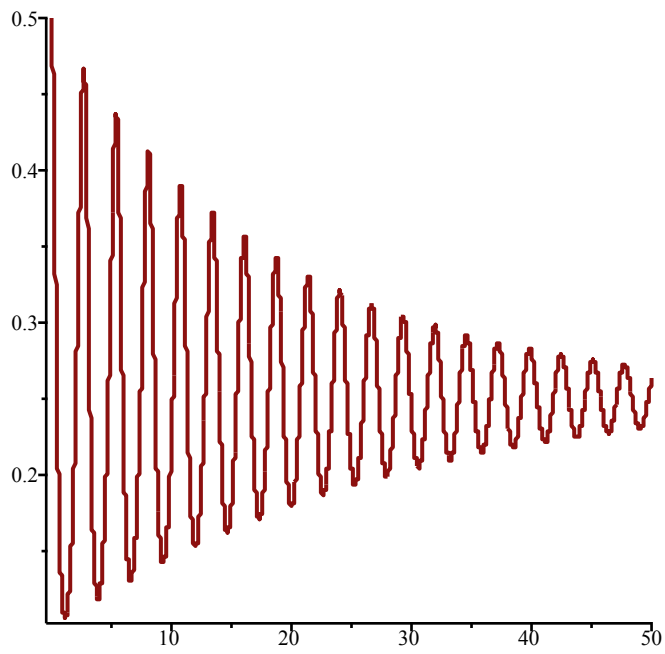
(25)

> *F* := *VolterraM*(3, 6, 2, 5, 8, *x*, *y*)

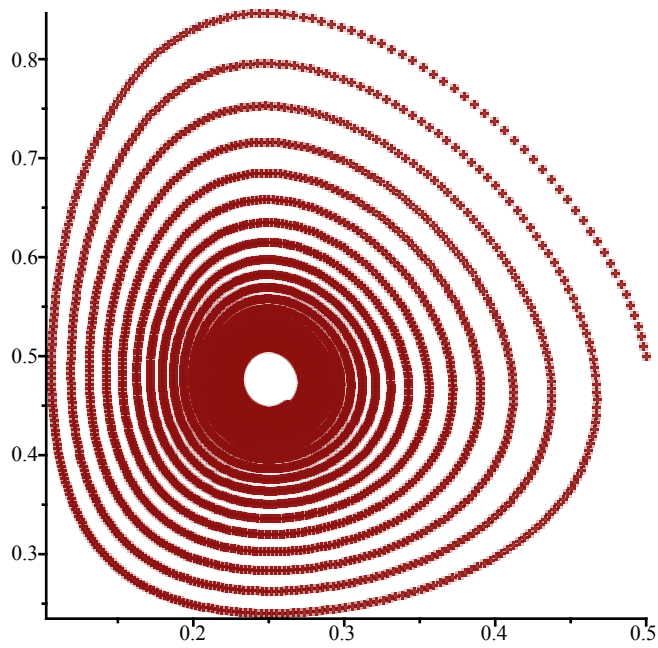
$$F := \left[3x \left(1 - \frac{x}{5} \right) - 6xy, 8xy - 2y \right]$$

(26)

> *TimeSeries*(*F*, [*x*, *y*], [0.5, 0.5], 0.01, 50, 1)



> *PhaseDiag*(*F*, [*x*, *y*], [0.5, 0.5], 0.01, 50)



> *SEquP*(*F*, [*x*, *y*])

{[0.2500000000, 0.4750000000]}

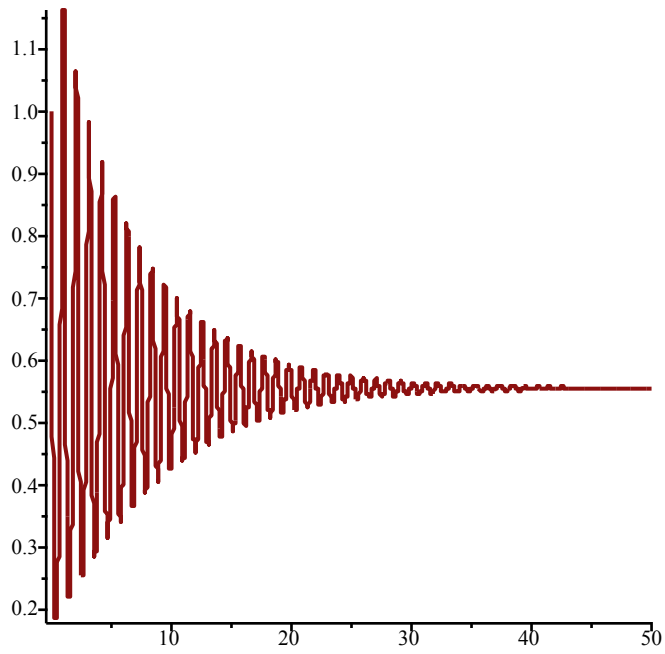
(27)

> *F* := *VolterraM*(8, 3, 5, 7, 9, *x*, *y*)

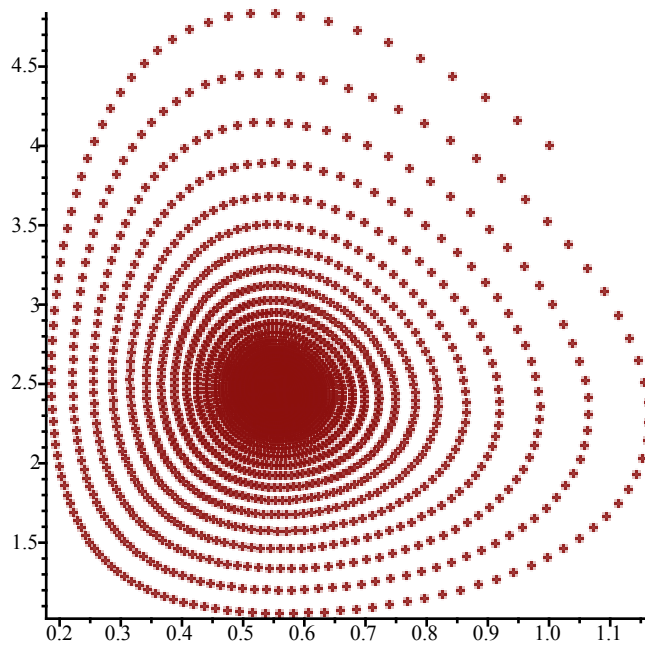
$$F := \left[8x \left(1 - \frac{x}{7} \right) - 3xy, 9xy - 5y \right]$$

(28)

> *TimeSeries*(*F*, [*x*, *y*], [1, 4], 0.01, 50, 1)



> *PhaseDiag*(*F*, [*x*, *y*], [1, 4], 0.01, 50, 1)



> *SEquP*(*F*, [*x*, *y*])

{[0.5555555556, 2.455026455]}

(29)

> #The modified Volterra system does have stable equilibria that corresponds to the horizontal asymptote found in the time series