

```
> #OK to post
>
> #Anne Somalwar, hw21, 11.15.2021
>
> read "C:/Users/aks238/OneDrive - Rutgers University/Documents/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);*

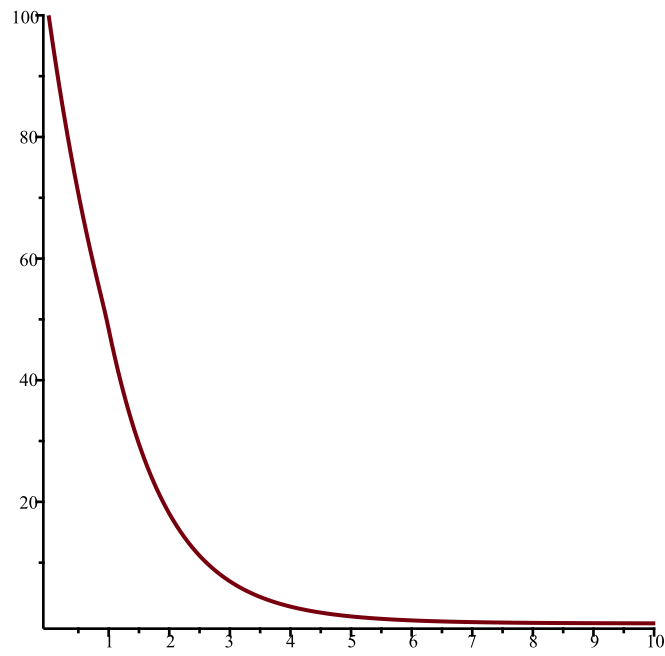
*For a list of the functions that give examples of Discrete-time dynamical systems (some famous),
type: HelpDDM());*

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

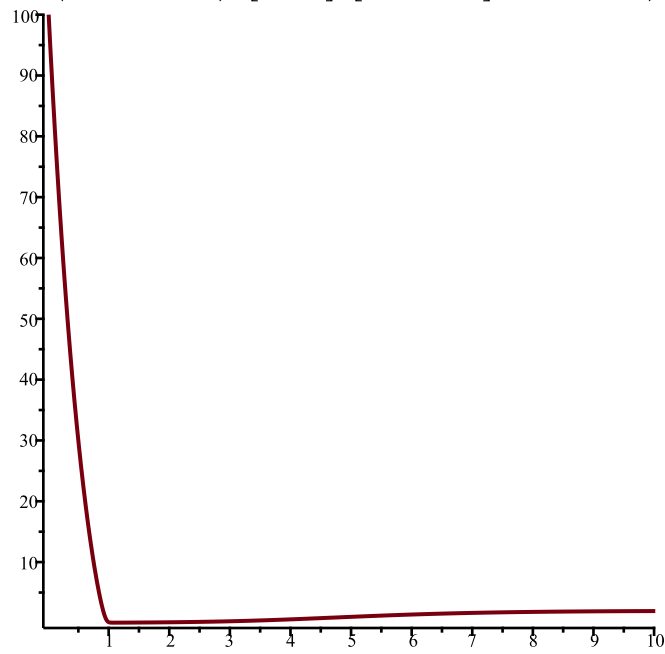
*For a list of the functions continuous-time dynamical systems (some famous) type: HelpCDM());
For help with any of them type: Help(ProcedureName);*

(1)

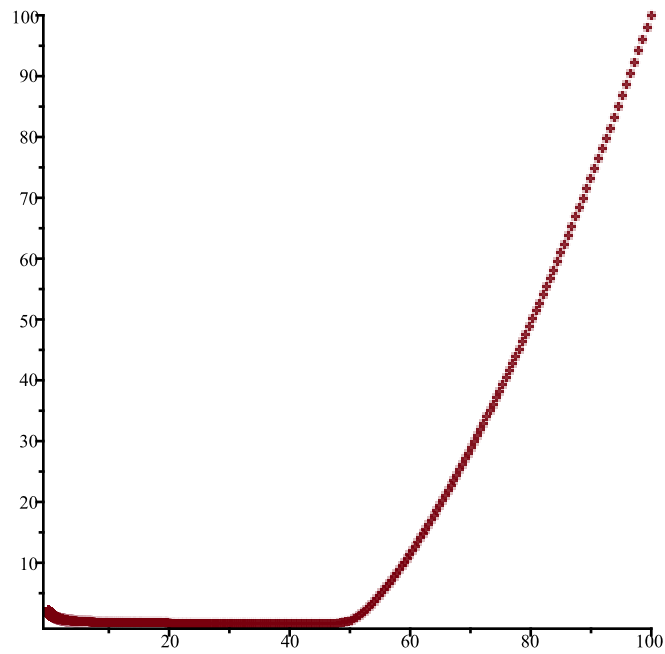
```
>
> #1
>
>
>
> TimeSeries(ChemoStat(N, C, 0.3, 2), [N, C], [100, 100], 0.01, 10, 1)
```



> *TimeSeries*(ChemoStat($N, C, 0.3, 2$), [N, C], [100, 100], 0.01, 10, 2)



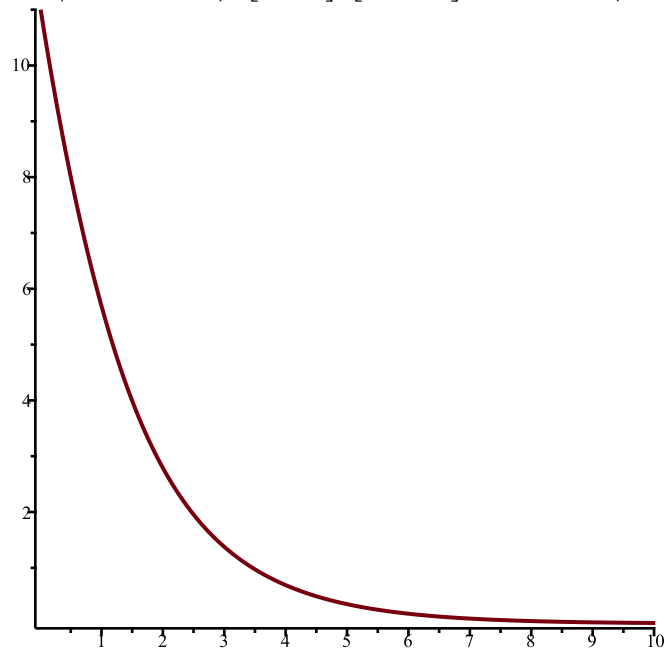
>
> *PhaseDiag*(ChemoStat($N, C, 0.3, 2$), [N, C], [100, 100], 0.01, 10)



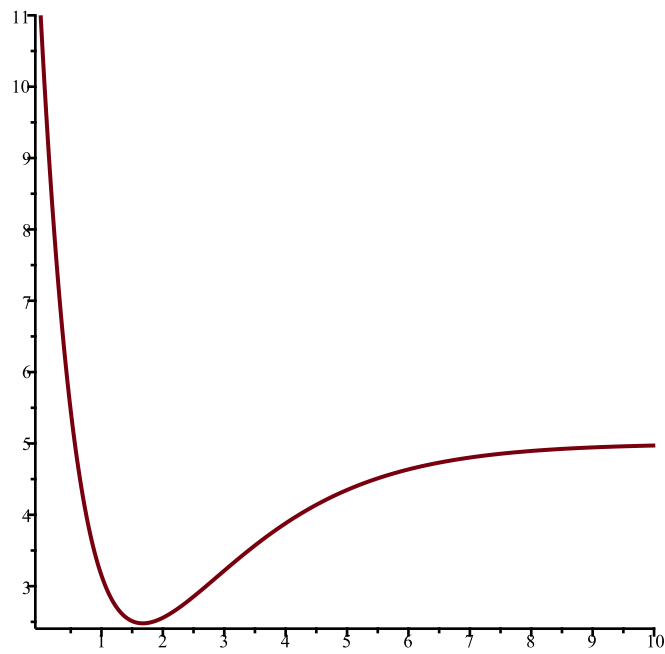
> *SEquP*(ChemoStat(*N*, *C*, 0.3, 2), [*N*, *C*])
 { [0., 2.], [1.028571429, -1.428571429] }

(2)

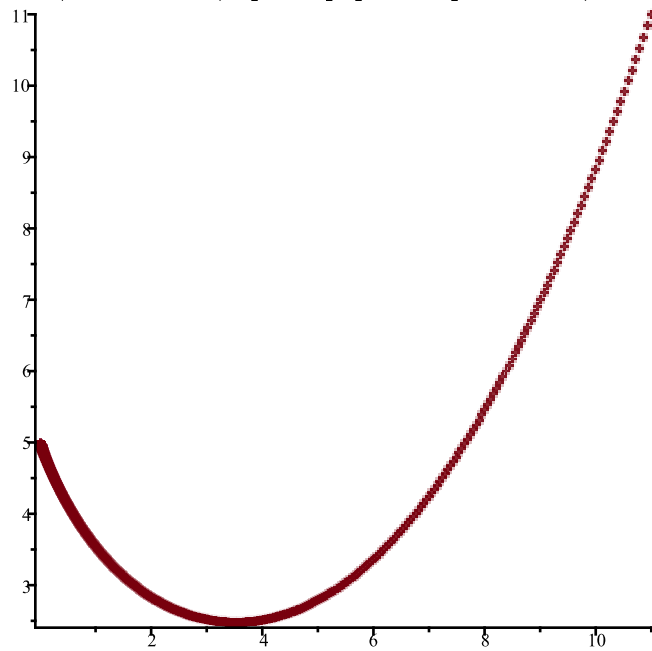
> *TimeSeries*(ChemoStat(*N*, *C*, 0.4, 5), [*N*, *C*], [11, 11], 0.01, 10, 1)



> *TimeSeries*(ChemoStat(*N*, *C*, 0.4, 5), [*N*, *C*], [11, 11], 0.01, 10, 2)



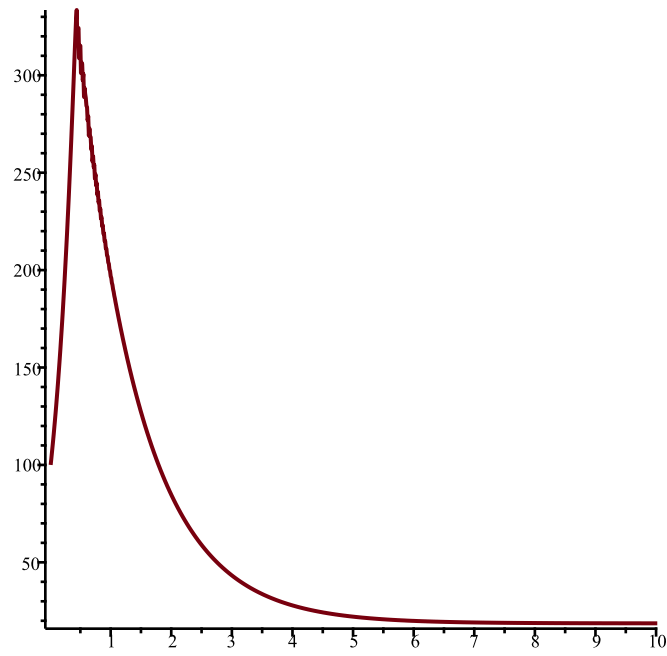
```
> PhaseDiag(ChemoStat(N, C, 0.4, 5), [N, C], [11, 11], 0.01, 10)
```



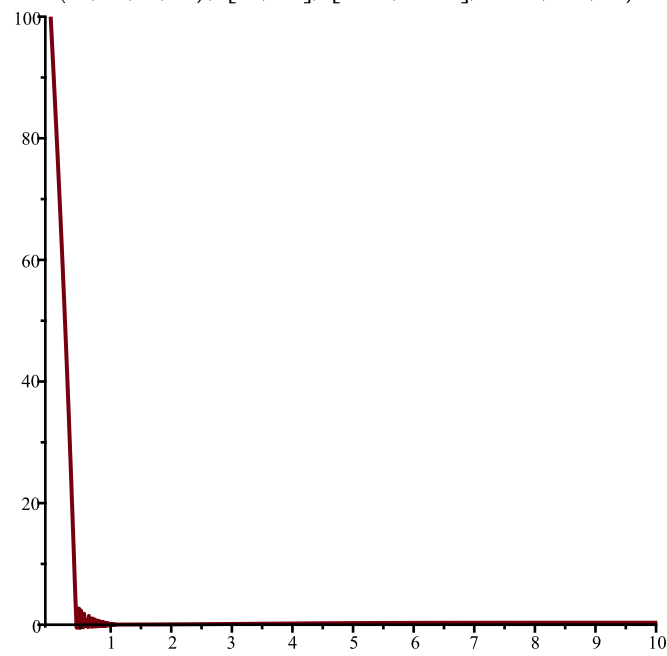
```
> SEquP(ChemoStat(N, C, 0.4, 5), [N, C])
      { [0., 5.], [2.666666667, -1.666666667] }
```

```
> TimeSeries(ChemoStat(N, C, 4, 5), [N, C], [100, 100], 0.01, 10, 1)
```

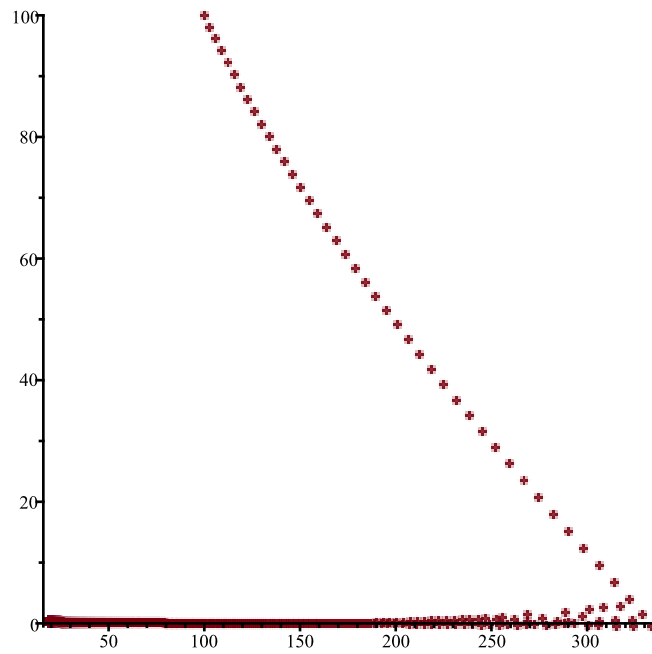
(3)



> *TimeSeries*(ChemoStat($N, C, 4, 5$), [N, C], [100, 100], 0.01, 10, 2)



> *PhaseDiag*(ChemoStat($N, C, 4, 5$), [N, C], [100, 100], 0.01, 10)

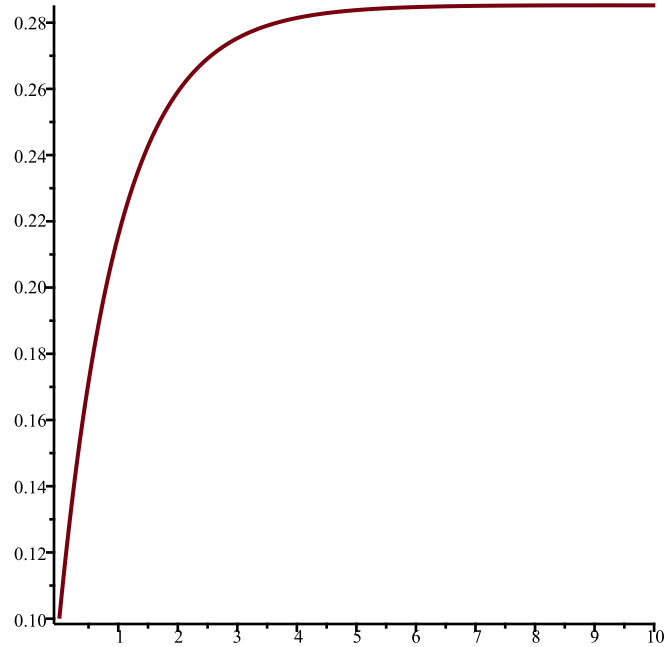


> *SEquP(ChemoStat(N, C, 4, 5), [N, C])*
{ [18.66666667, 0.3333333333] }

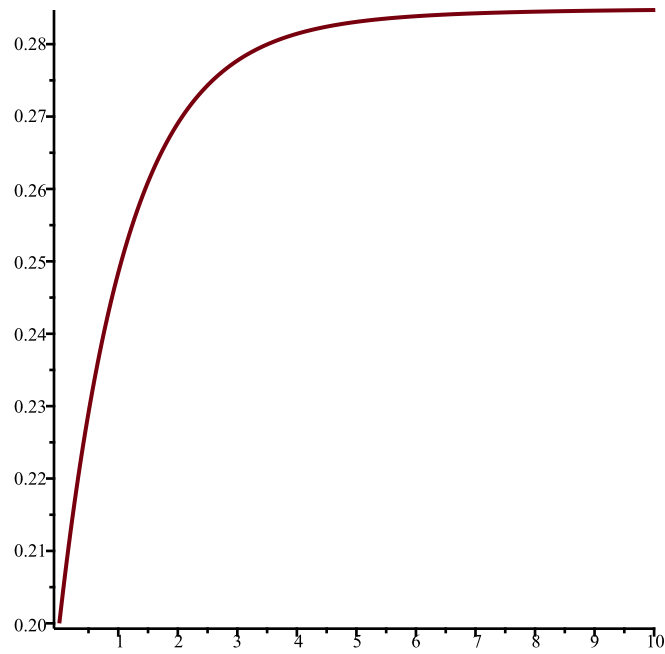
(4)

>
>
>
>

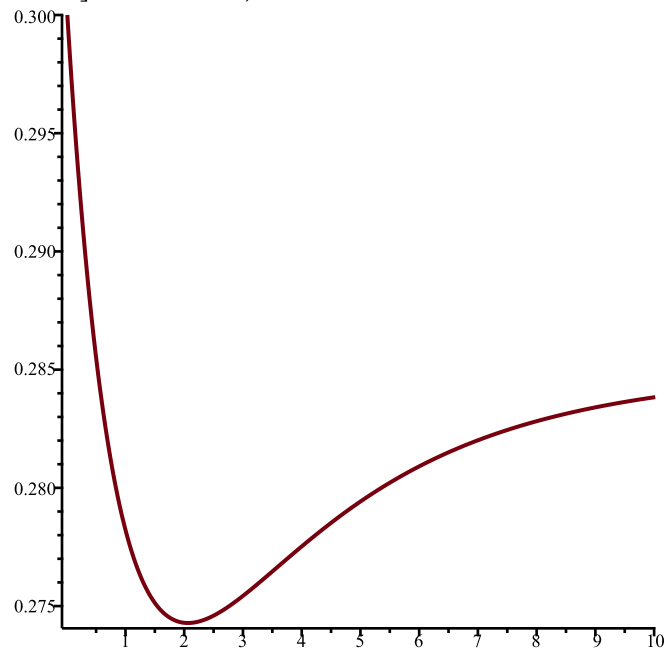
> *TimeSeries(GeneNet(0.1, 0.2, 0.3, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.1, 0.2, 0.3, 0.4, 0.5, 0.3], 0.01, 10, 1)*



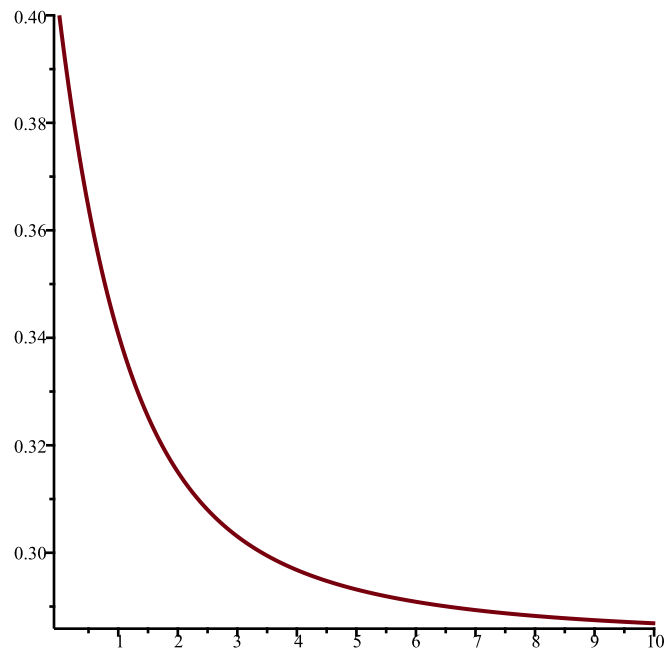
> *TimeSeries(GeneNet(0.1, 0.2, 0.3, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.1, 0.2, 0.3, 0.4, 0.5, 0.3], 0.01, 10, 2)*



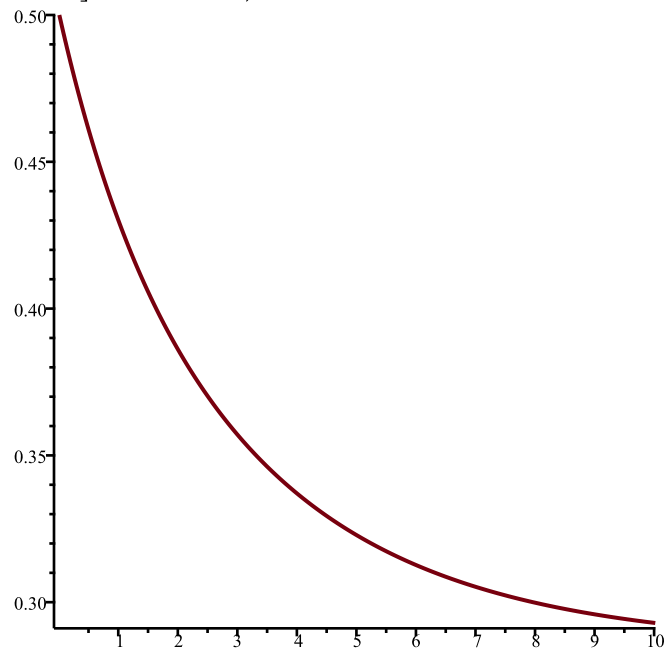
> *TimeSeries*(GeneNet(0.1, 0.2, 0.3, 2, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [0.1, 0.2, 0.3, 0.4, 0.5, 0.3], 0.01, 10, 3)



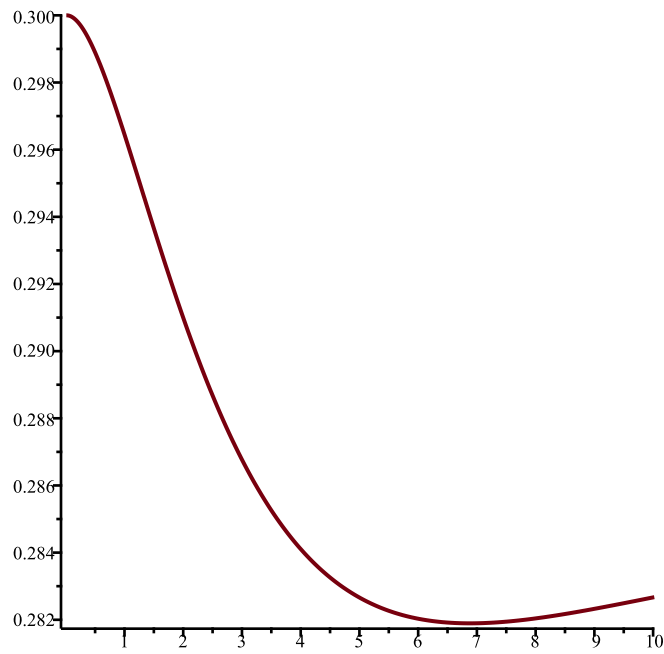
> *TimeSeries*(GeneNet(0.1, 0.2, 0.3, 2, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [0.1, 0.2, 0.3, 0.4, 0.5, 0.3], 0.01, 10, 4)



> *TimeSeries*(GeneNet(0.1, 0.2, 0.3, 2, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [0.1, 0.2, 0.3, 0.4, 0.5, 0.3], 0.01, 10, 5)



> *TimeSeries*(GeneNet(0.1, 0.2, 0.3, 2, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [0.1, 0.2, 0.3, 0.4, 0.5, 0.3], 0.01, 10, 6)

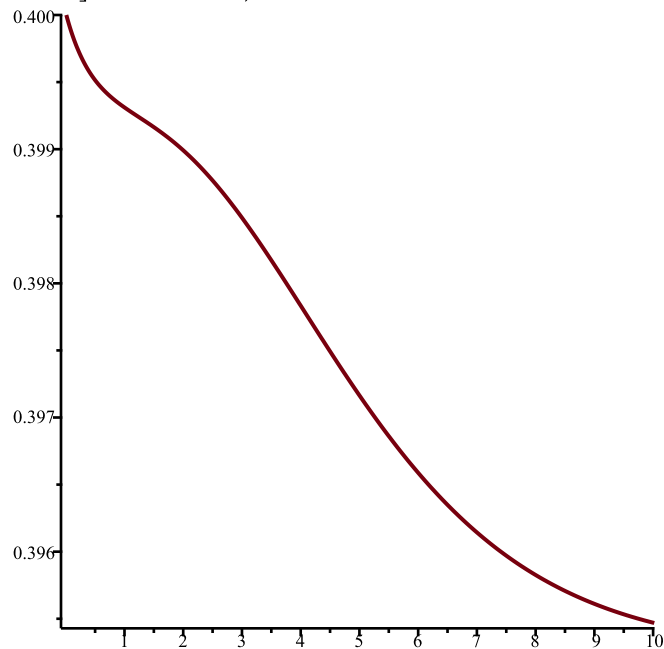


```
> SEquP(GeneNet(0.1, 0.2, 0.3, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3])
{[0.2849775630, 0.2849775630, 0.2849775630, 0.2849775630, 0.2849775630,
0.2849775630]}
```

(5)

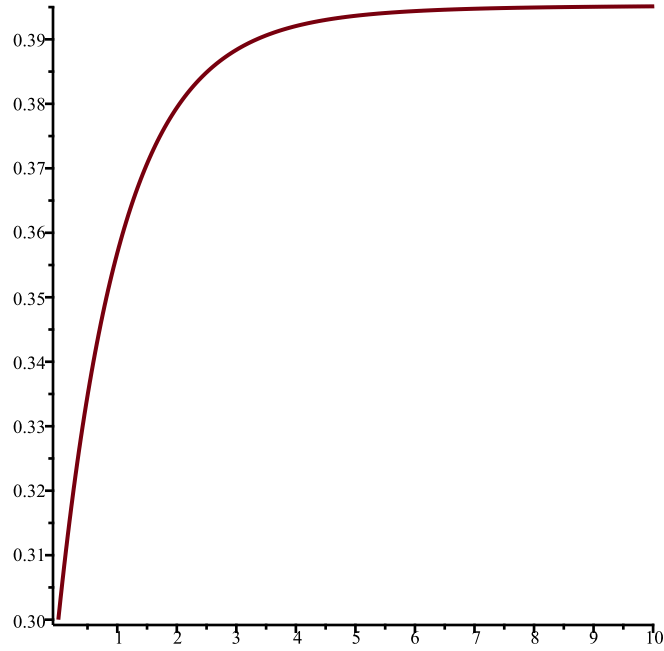
```
>
>
>
>
>
>
```

```
> TimeSeries(GeneNet(0.2, 0.2, 0.5, 4, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.4,
0.3, 0.1, 0.5, 0.5, 0.3], 0.01, 10, 1)
```

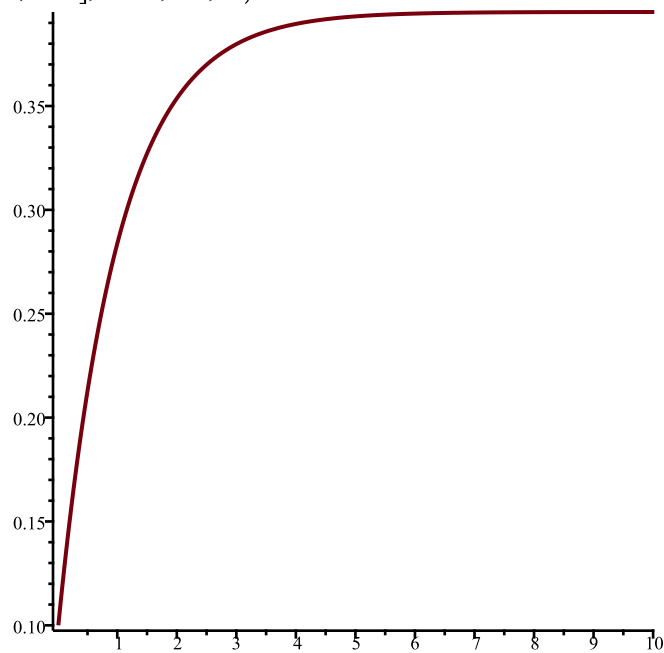


```
> TimeSeries(GeneNet(0.2, 0.2, 0.5, 4, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.4,
```

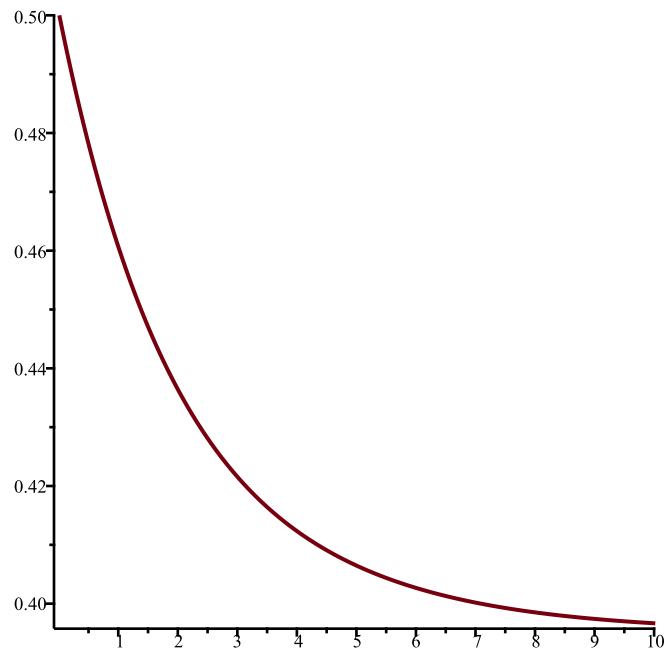
0.3, 0.1, 0.5, 0.5, 0.3], 0.01, 10, 2)



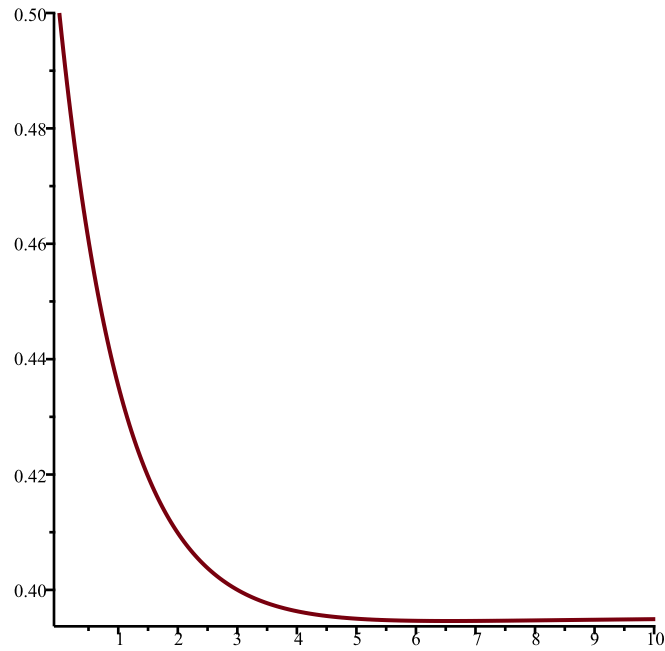
> TimeSeries(GeneNet(0.2, 0.2, 0.5, 4, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.4, 0.3, 0.1, 0.5, 0.5, 0.3], 0.01, 10, 3)



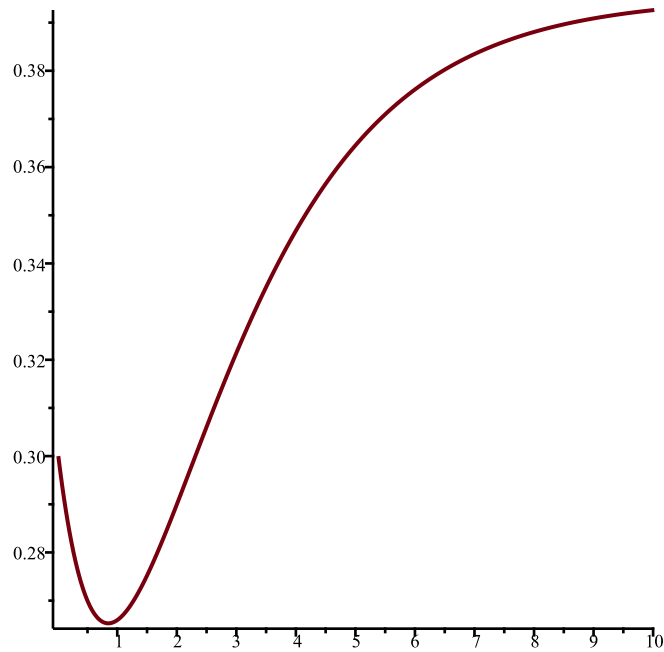
> TimeSeries(GeneNet(0.2, 0.2, 0.5, 4, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.4, 0.3, 0.1, 0.5, 0.5, 0.3], 0.01, 10, 4)



> *TimeSeries*(*GeneNet*(0.2, 0.2, 0.5, 4, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [0.4, 0.3, 0.1, 0.5, 0.5, 0.3], 0.01, 10, 5)



> *TimeSeries*(*GeneNet*(0.2, 0.2, 0.5, 4, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*], [0.4, 0.3, 0.1, 0.5, 0.5, 0.3], 0.01, 10, 6)

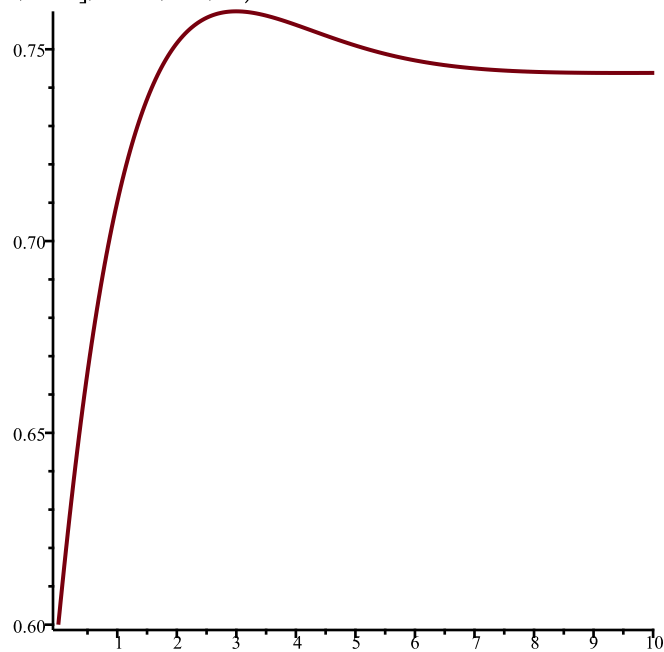


```
> SEquP(GeneNet(0.1, 0.2, 0.3, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3])
{[0.2849775630, 0.2849775630, 0.2849775630, 0.2849775630, 0.2849775630,
0.2849775630]}
```

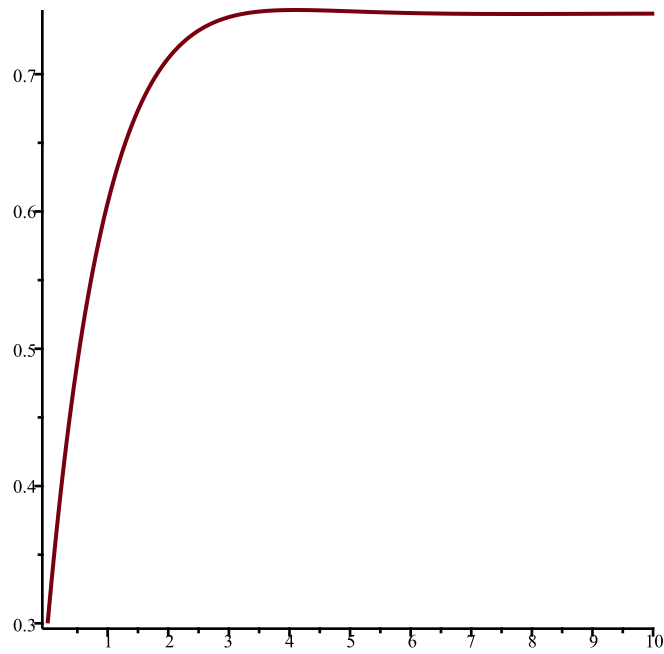
(6)

```
>
>
>
```

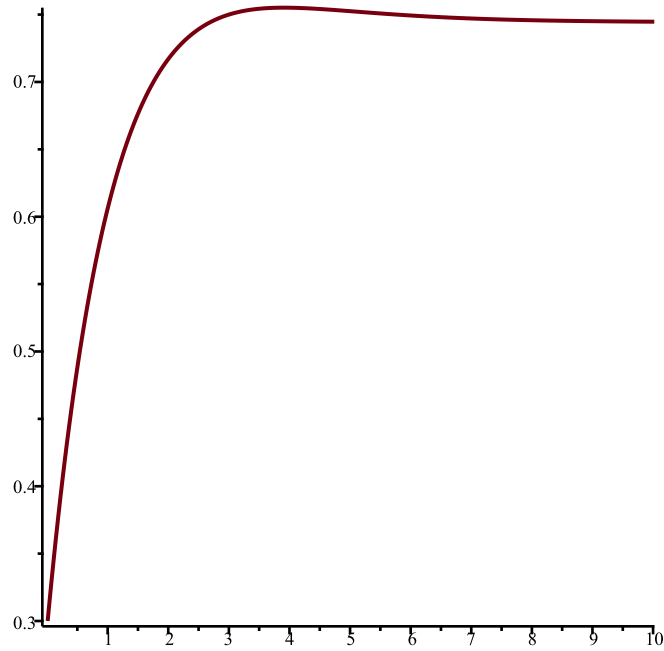
```
> TimeSeries(GeneNet(0.5, 0.3, 0.7, 5, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.6,
0.3, 0.3, 0.5, 0.6, 0.7], 0.01, 10, 1)
```



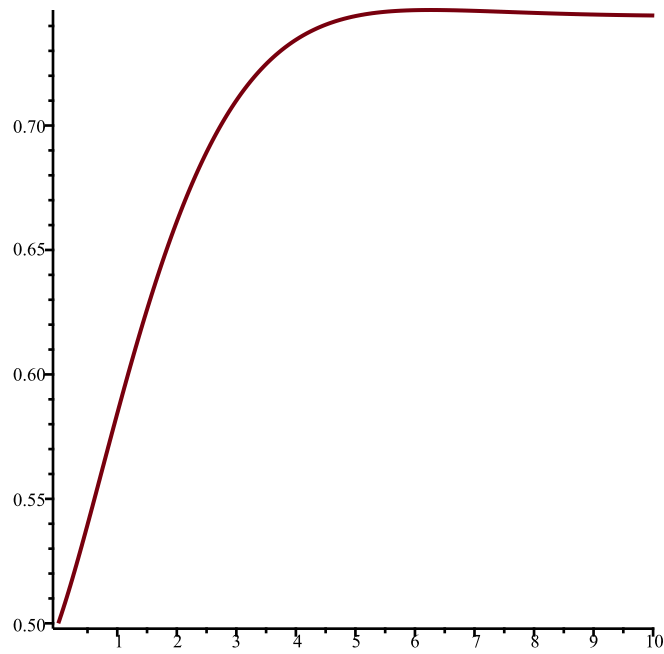
```
> TimeSeries(GeneNet(0.5, 0.3, 0.7, 5, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.6,
0.3, 0.3, 0.5, 0.6, 0.7], 0.01, 10, 2)
```



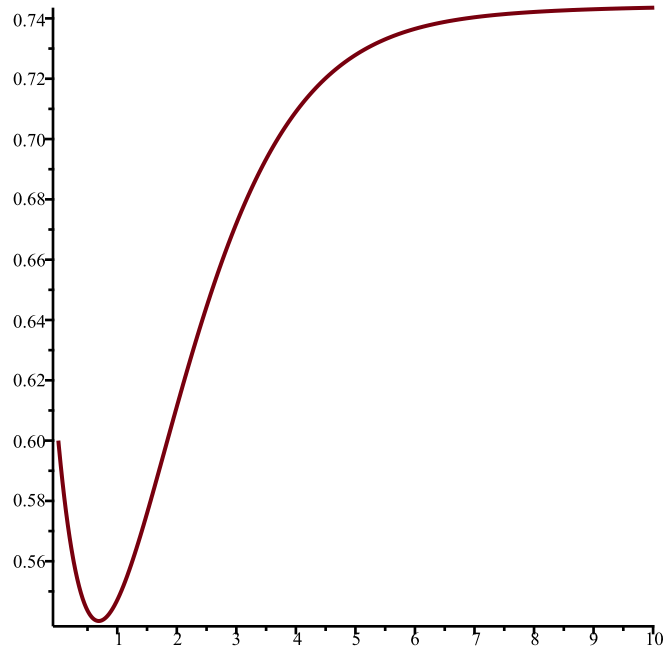
> *TimeSeries(GeneNet(0.5, 0.3, 0.7, 5, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.6, 0.3, 0.3, 0.5, 0.6, 0.7], 0.01, 10, 3)*



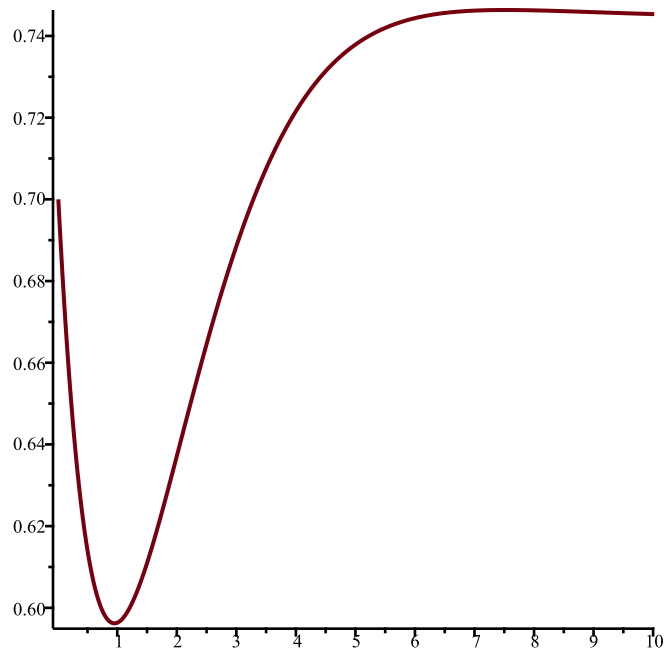
> *TimeSeries(GeneNet(0.5, 0.3, 0.7, 5, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.6, 0.3, 0.3, 0.5, 0.6, 0.7], 0.01, 10, 4)*



> *TimeSeries(GeneNet(0.5, 0.3, 0.7, 5, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.6, 0.3, 0.3, 0.5, 0.6, 0.7], 0.01, 10, 5)*



> *TimeSeries(GeneNet(0.5, 0.3, 0.7, 5, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.6, 0.3, 0.3, 0.5, 0.6, 0.7], 0.01, 10, 6)*



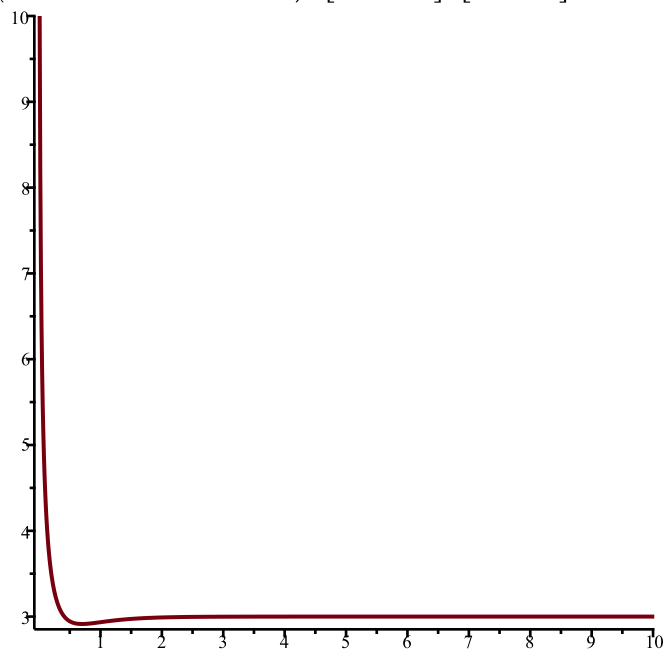
> *SEquP*(*GeneNet*(0.5, 0.3, 0.7, 5, *m1*, *m2*, *m3*, *p1*, *p2*, *p3*), [*m1*, *m2*, *m3*, *p1*, *p2*, *p3*])

∅

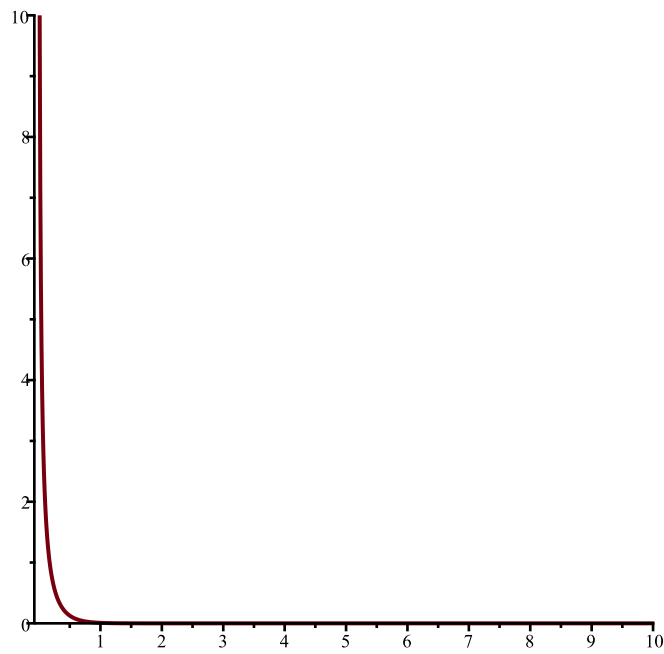
(7)

>
>
>
>
>
>

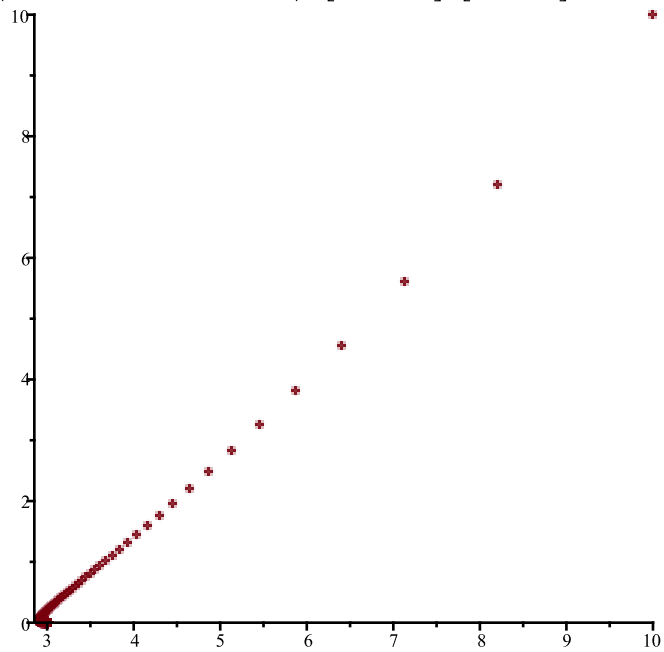
> *TimeSeries*(*Lotka*(2, 3, 2, 4, 2, 5, *N1*, *N2*), [*N1*, *N2*], [10, 10], 0.01, 10, 1)



> *TimeSeries*(*Lotka*(2, 3, 2, 4, 2, 5, *N1*, *N2*), [*N1*, *N2*], [10, 10], 0.01, 10, 2)



> PhaseDiag(Lotka(2, 3, 2, 4, 2, 5, N1, N2), [N1, N2], [10, 10], 0.01, 10)

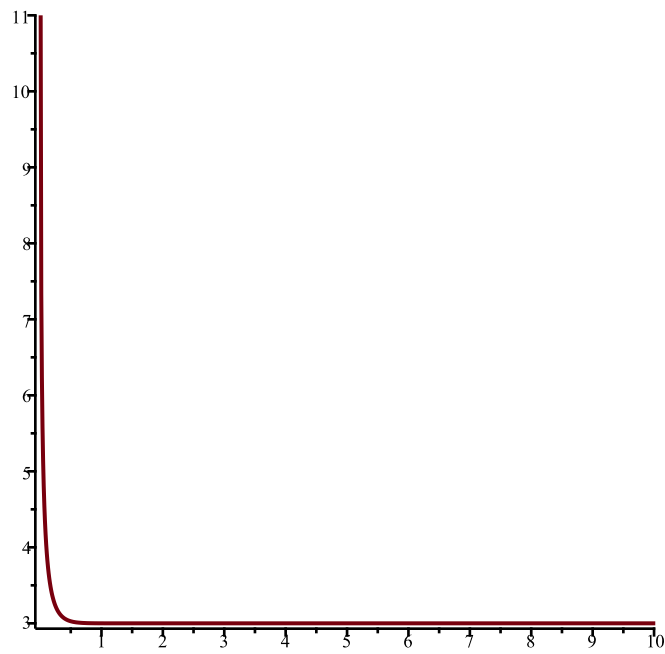


> SEquP(Lotka(2, 3, 2, 4, 2, 5, N1, N2), [N1, N2])
 { [0., 4.], [3., 0.] }

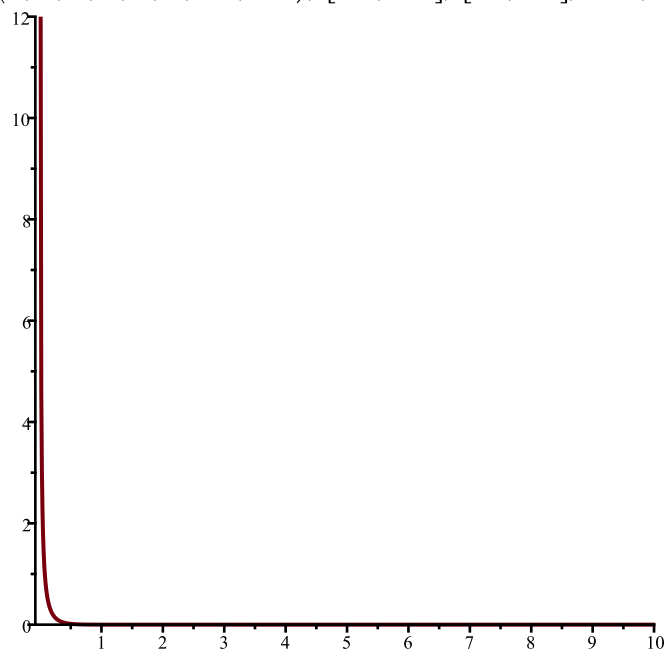
(8)

>
 >
 >
 >
 >

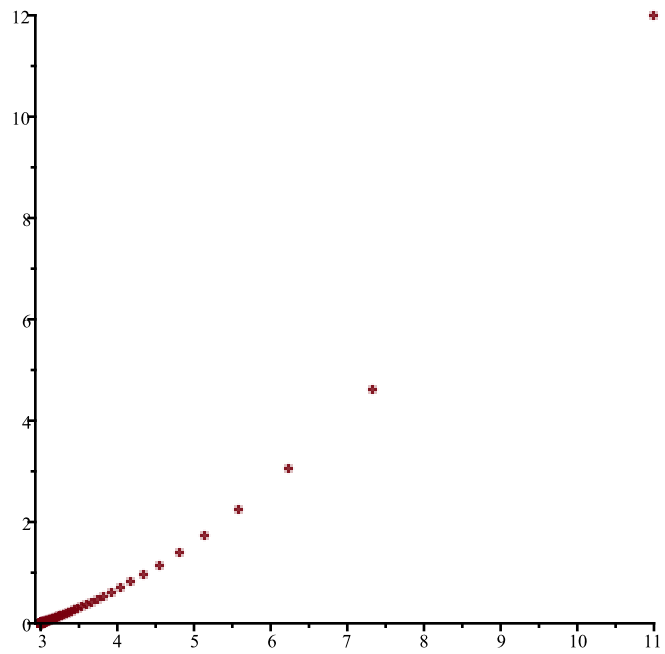
> TimeSeries(Lotka(5, 3, 6, 4, 1, 3, N1, N2), [N1, N2], [11, 12], 0.01, 10, 1)



> *TimeSeries*(Lotka(5, 3, 6, 4, 1, 3, $N1$, $N2$), [$N1$, $N2$], [11, 12], 0.01, 10, 2)



> *PhaseDiag*(Lotka(5, 3, 6, 4, 1, 3, $N1$, $N2$), [$N1$, $N2$], [11, 12], 0.01, 10)

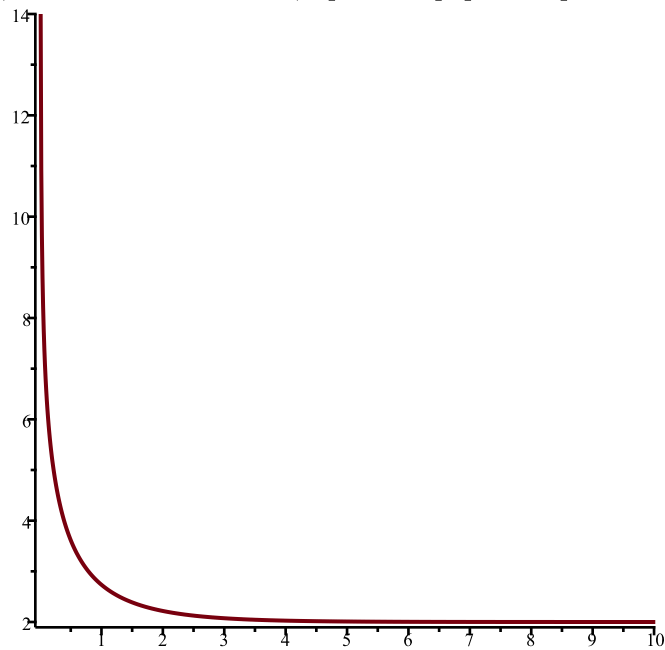


```
> SEquP(Lotka(5, 3, 6, 4, 1, 3, N1, N2), [N1, N2])
      { [0., 4.], [3., 0.] }
```

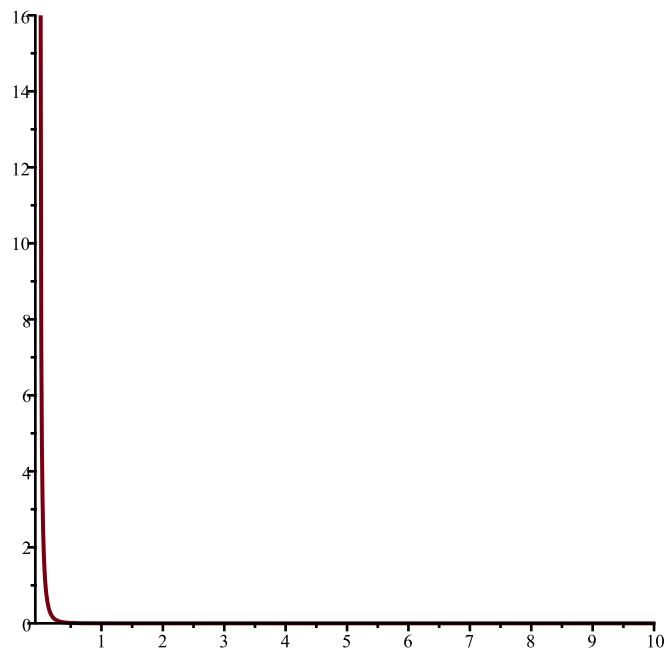
(9)

- >
- >
- >
- >
- >

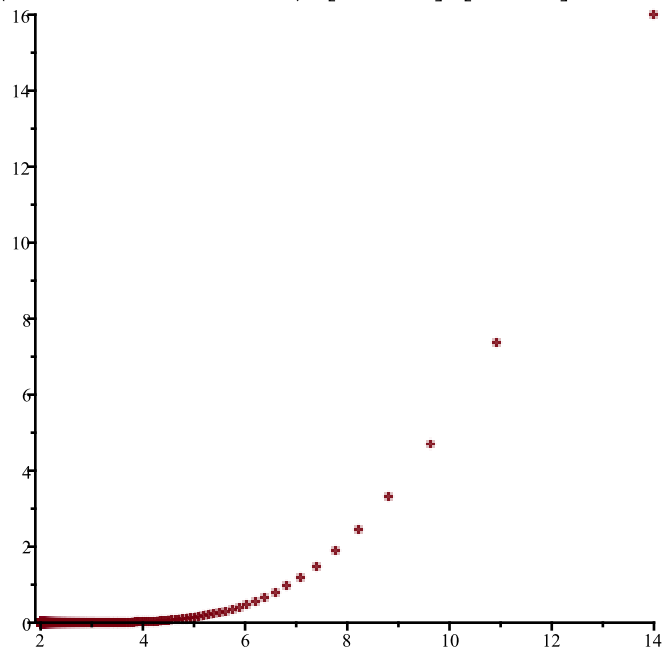
```
> TimeSeries(Lotka(1, 2, 4, 4, 2, 3, N1, N2), [N1, N2], [14, 16], 0.01, 10, 1)
```



```
> TimeSeries(Lotka(1, 2, 4, 4, 2, 3, N1, N2), [N1, N2], [14, 16], 0.01, 10, 2)
```



> *PhaseDiag*(*Lotka*(1, 2, 4, 4, 2, 3, *N1*, *N2*), [*N1*, *N2*], [14, 16], 0.01, 10)

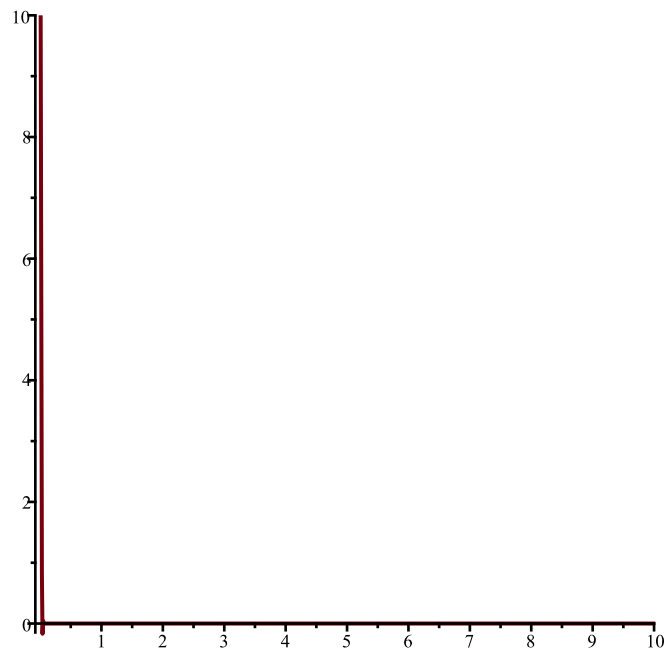


> *SEquP*(*Lotka*(1, 2, 4, 4, 2, 3, *N1*, *N2*), [*N1*, *N2*])
 { [0., 4.], [2., 0.] }

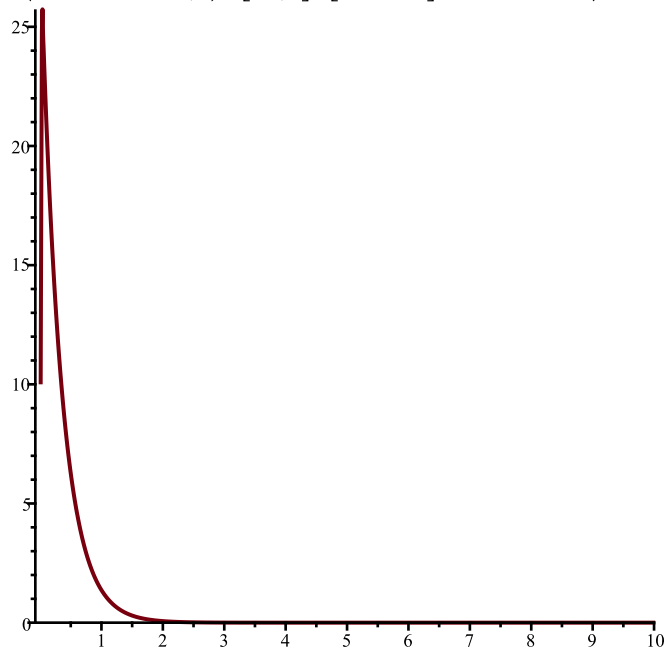
(10)

>
 >
 >
 >
 >

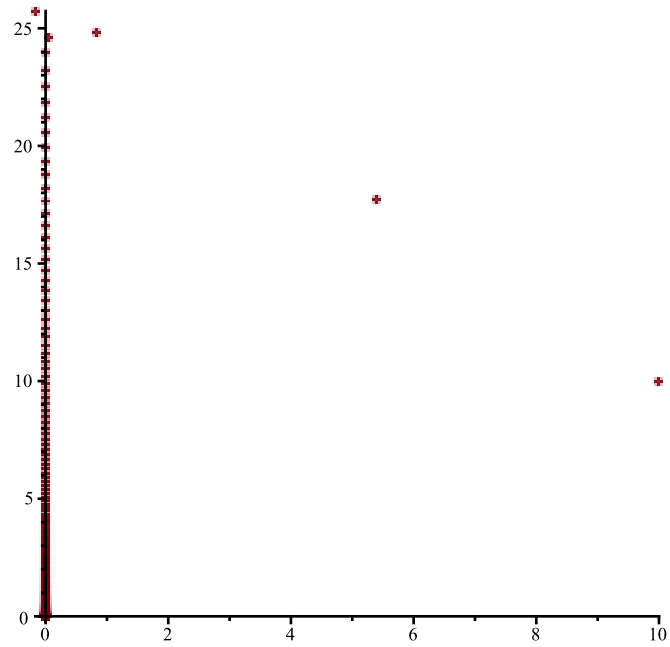
> *TimeSeries*(*Volterra*(4, 5, 3, 8, *x*, *y*), [*x*, *y*], [10, 10], 0.01, 10, 1)



> *TimeSeries*(*Volterra*(4, 5, 3, 8, x , y), [x , y], [10, 10], 0.01, 10, 2)



> *PhaseDiag*(*Volterra*(4, 5, 3, 8, x , y), [x , y], [10, 10], 0.01, 10, 1)



> *SEquP*(*Volterra*(4, 5, 3, 8, *x*, *y*), [*x*, *y*])

∅

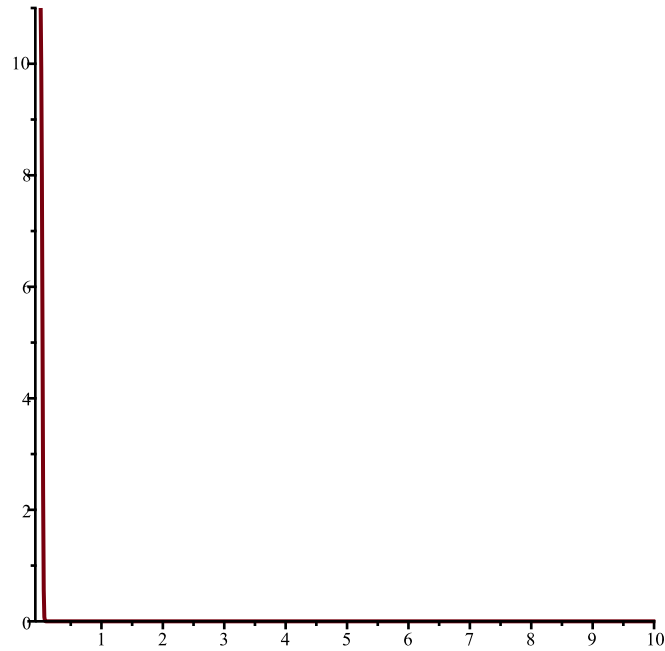
(11)

>

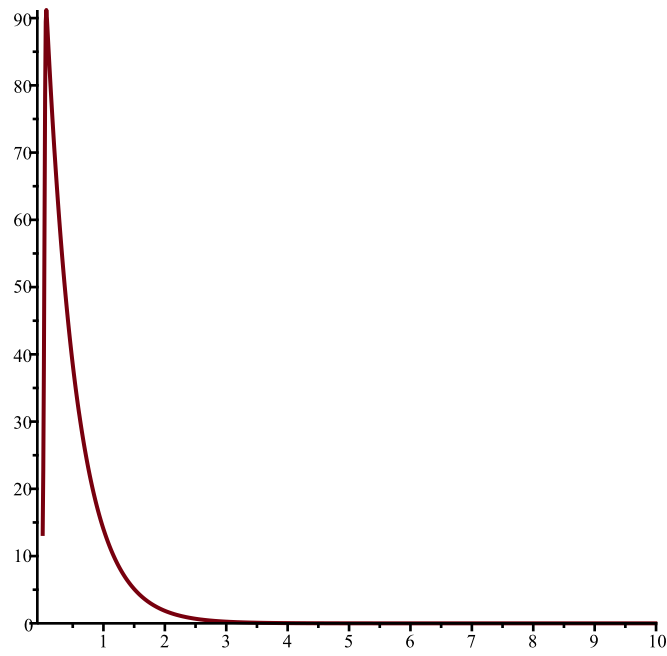
>

>

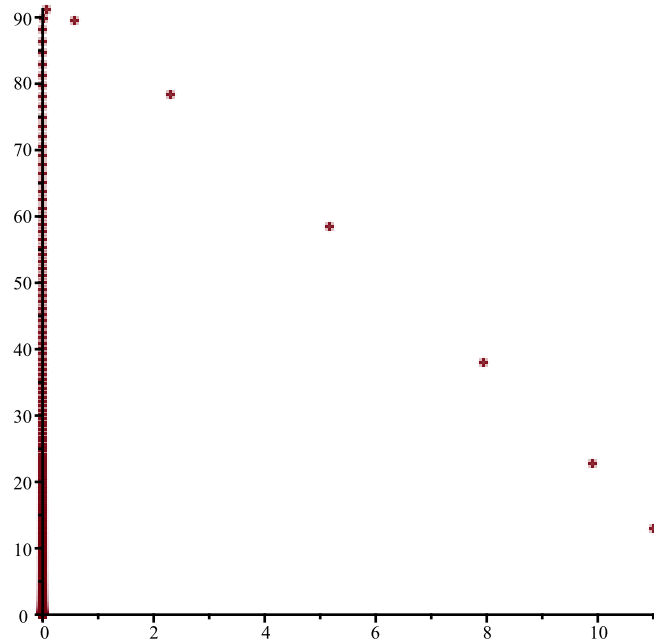
> *TimeSeries*(*Volterra*(3, 1, 2, 7, *x*, *y*), [*x*, *y*], [11, 13], 0.01, 10, 1)



> *TimeSeries*(*Volterra*(3, 1, 2, 7, *x*, *y*), [*x*, *y*], [11, 13], 0.01, 10, 2)



> *PhaseDiag*(*Volterra*(3, 1, 2, 7, x, y), [x, y], [11, 13], 0.01, 10)



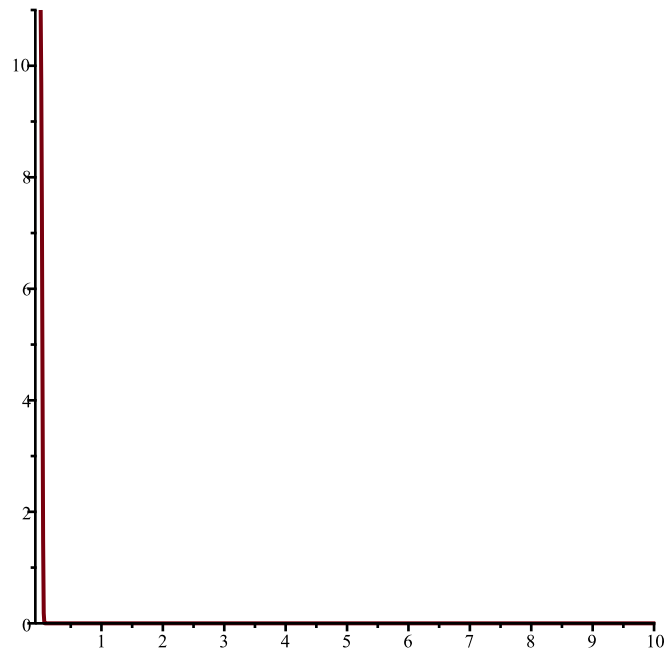
> *SEquP*(*Volterra*(3, 1, 2, 7, x, y), [x, y])

∅

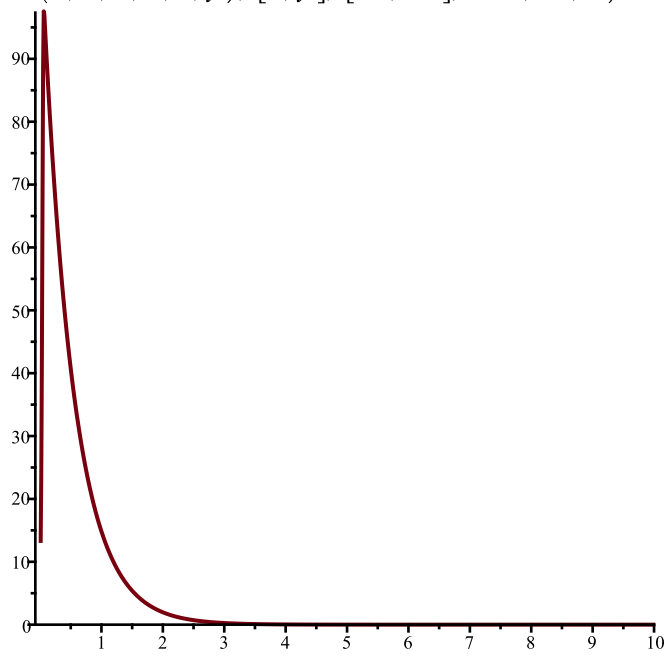
(12)

>
>
>
>

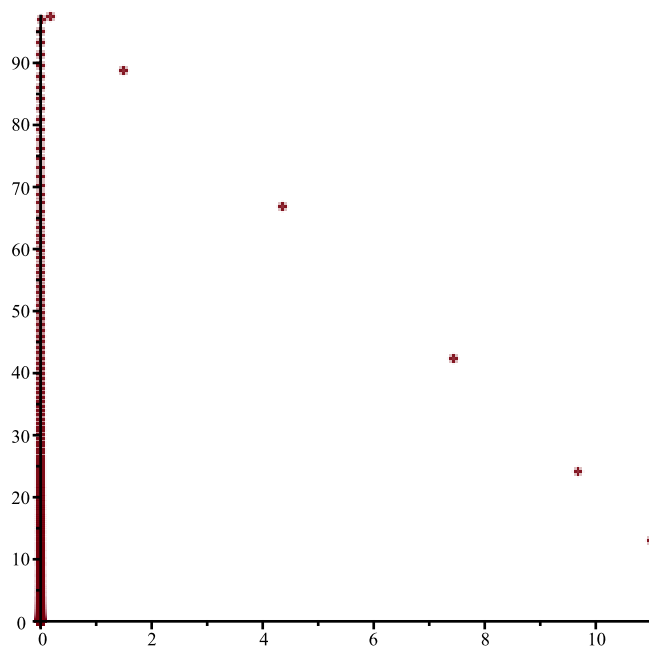
> *TimeSeries*(*Volterra*(1, 1, 2, 8, x, y), [x, y], [11, 13], 0.01, 10, 1)



> *TimeSeries*(*Volterra*(1, 1, 2, 8, x, y), [x, y], [11, 13], 0.01, 10, 2)



> *PhaseDiag*(*Volterra*(1, 1, 2, 8, x, y), [x, y], [11, 13], 0.01, 10)



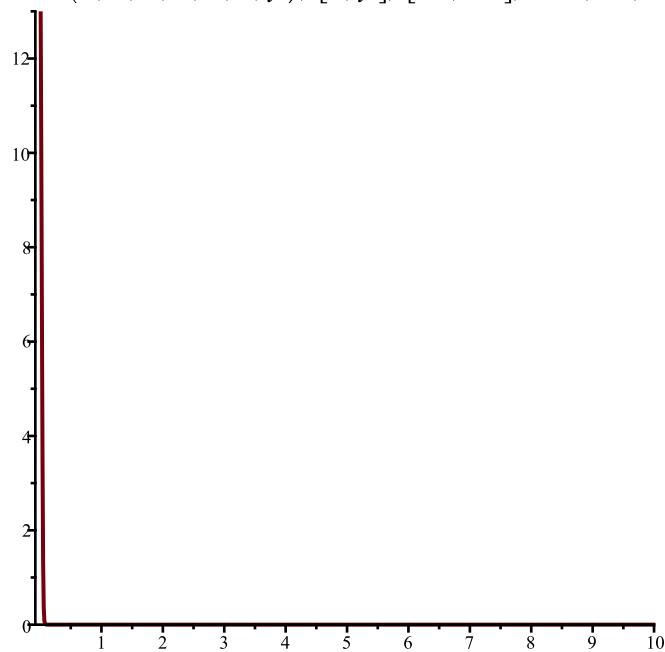
> *SEquP*(*Volterra*(1, 1, 2, 8, *x*, *y*), [*x*, *y*])

∅

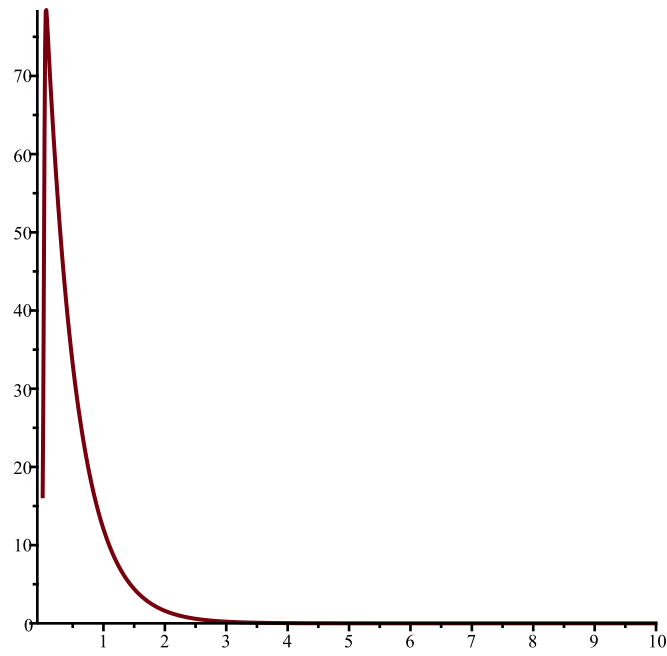
(13)

>
>
>
>
>
>
>
>

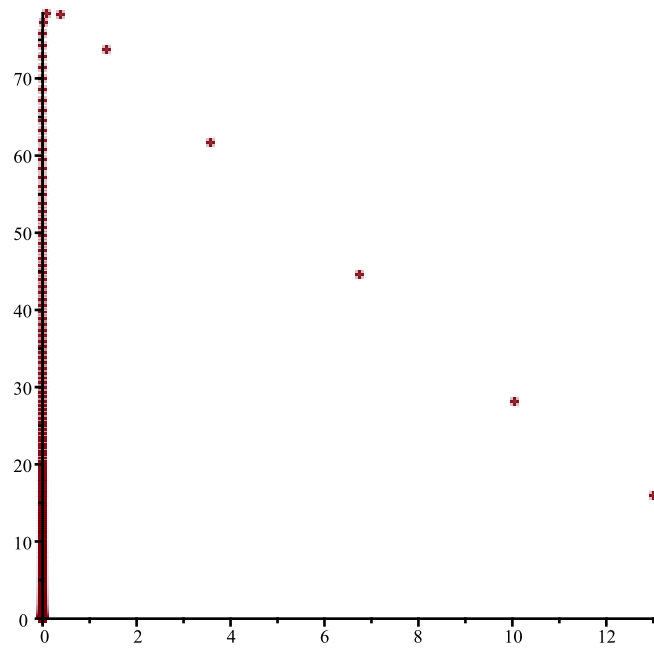
> *TimeSeries*(*VolterraM*(2, 1, 2, 3, 6, *x*, *y*), [*x*, *y*], [13, 16], 0.01, 10, 1)



> *TimeSeries*(*VolterraM*(2, 1, 2, 3, 6, *x*, *y*), [*x*, *y*], [13, 16], 0.01, 10, 2)



> *PhaseDiag*(*VolterraM*(2, 1, 2, 3, 6, *x*, *y*), [*x*, *y*], [13, 16], 0.01, 10)

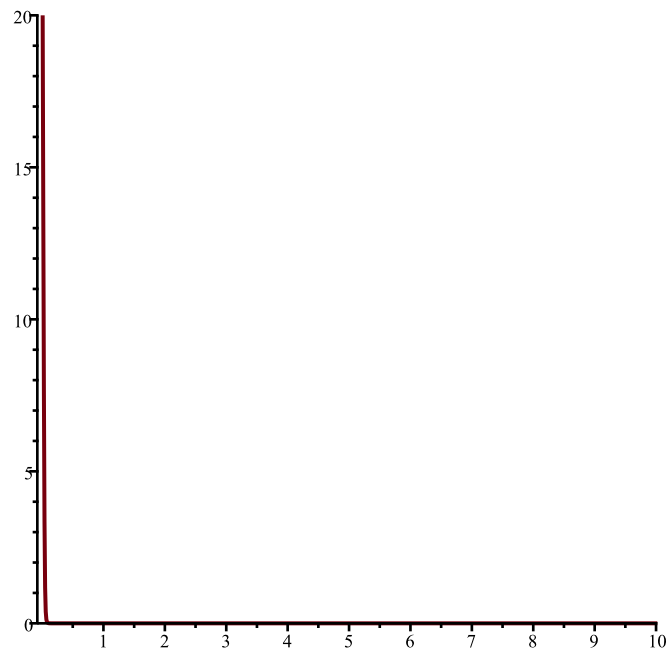


> *SEquP*(*VolterraM*(2, 1, 2, 3, 6, *x*, *y*), [*x*, *y*])
 { [0.3333333333, 1.777777778] }

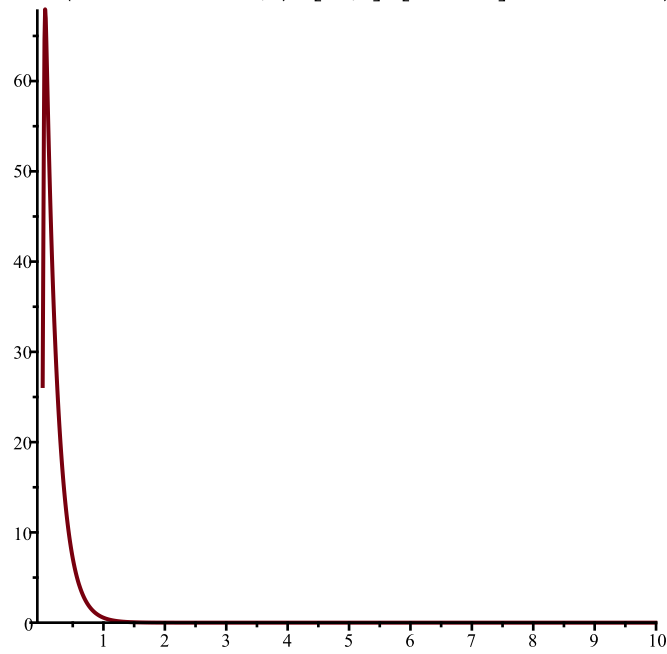
(14)

>
 >
 >
 >
 >

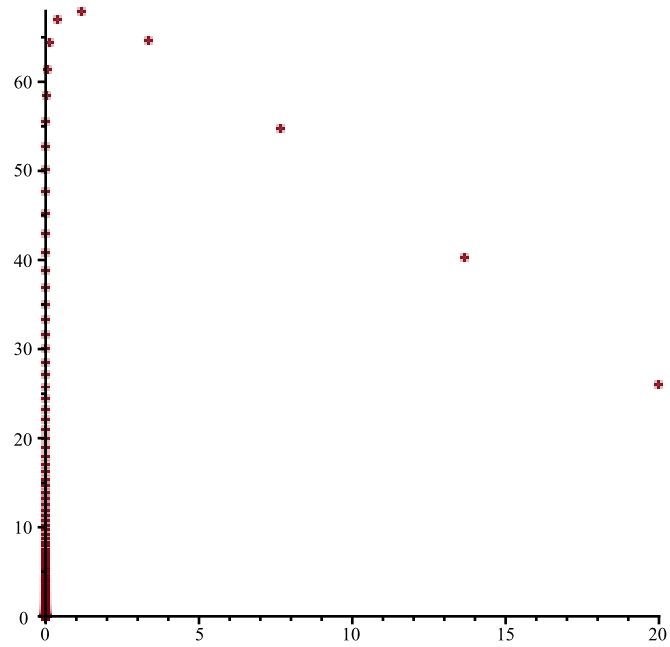
> *TimeSeries*(*VolterraM*(1, 1, 5, 3, 3, *x*, *y*), [*x*, *y*], [20, 26], 0.01, 10, 1)



> *TimeSeries*(*VolterraM*(1, 1, 5, 3, 3, x, y), [x, y], [20, 26], 0.01, 10, 2)



> *PhaseDiag*(*VolterraM*(1, 1, 5, 3, 3, x, y), [x, y], [20, 26], 0.01, 10)

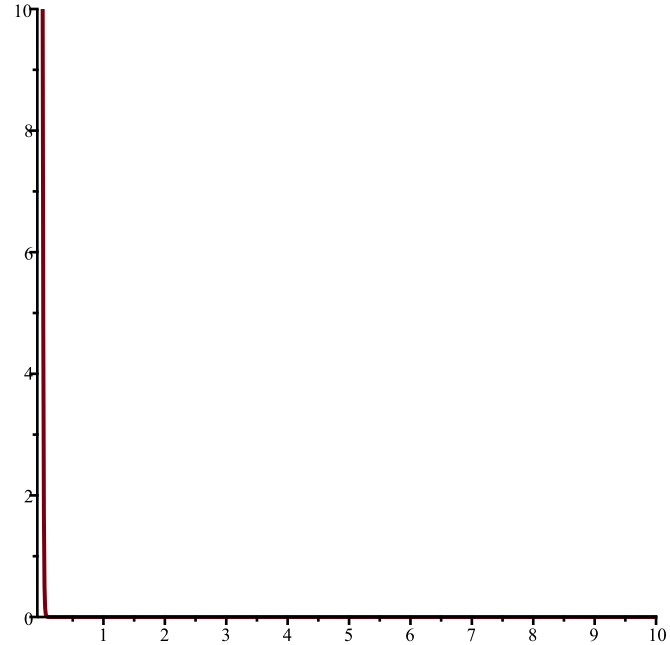


```
> SEquP(VolterraM(1, 1, 5, 3, 3, x, y), [x, y])
      { [1.666666667, 0.4444444444]}
```

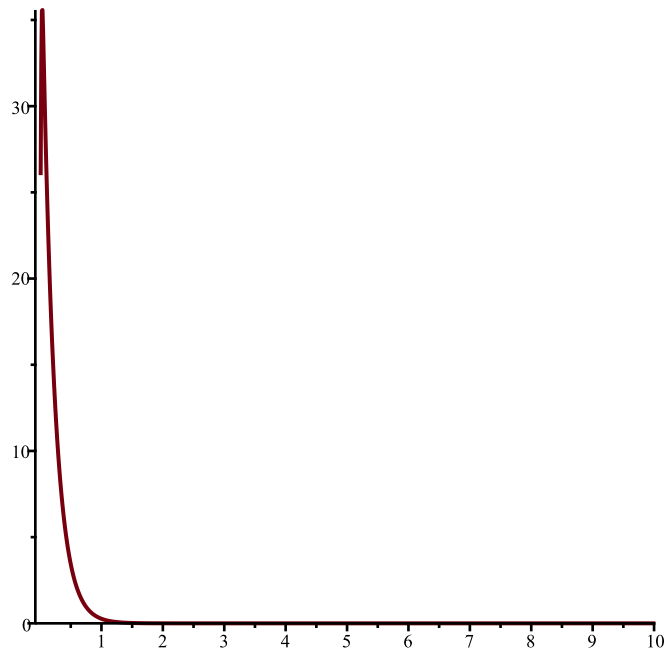
(15)

```
>
>
>
>
>
```

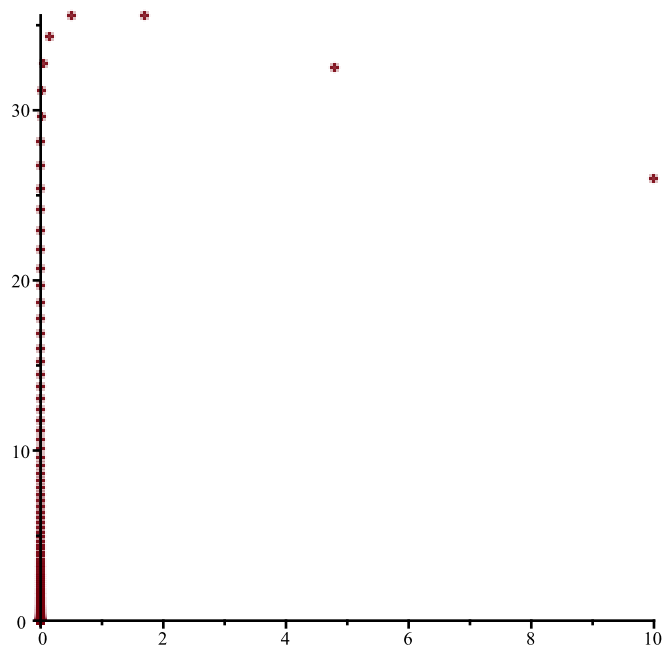
```
> TimeSeries(VolterraM(1, 2, 5, 9, 3, x, y), [x, y], [10, 26], 0.01, 10, 1)
```



```
> TimeSeries(VolterraM(1, 2, 5, 9, 3, x, y), [x, y], [10, 26], 0.01, 10, 2)
```



> PhaseDiag(VolterraM(1, 2, 5, 9, 3, x, y), [x, y], [10, 26], 0.01, 10)



> SEquP(VolterraM(1, 2, 5, 9, 3, x, y), [x, y])
 { [1.666666667, 0.4074074074] }