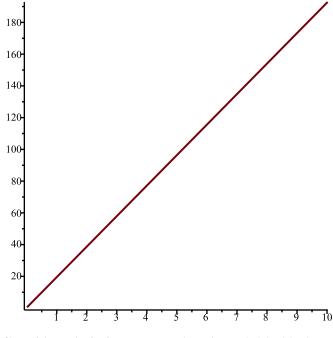
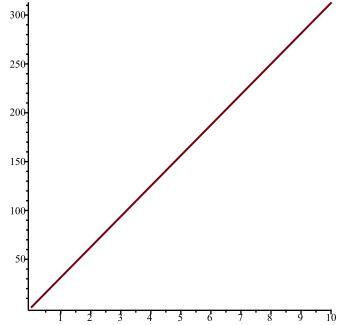
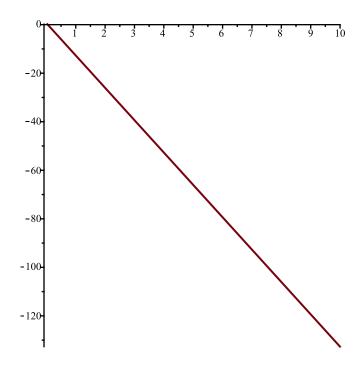
```
> #Mudassir Lodi
   #November 15, 2021
   #Homework 21
> read "C:/Users/mkl115/Documents/DMB.txt"
                                  First Written: Nov. 2021
This is DMB.txt, A Maple package to explore Dynamical models in Biology (both discrete and
    continuous)
accompanying the class Dynamical 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)
> Help(ChemoStat)
ChemoStat(N,C,a1,a2): The Chemostat continuous-time dynamical system with N=Bacterial
   population density, and C=nutient Concentration in growth chamber (see Table 4.1 of
    Edelstein-Keshet, p. 122)
with paramerts a1, a2, Equations (19a, (19b) in Edelestein-Keshet p. 127 (section 4.5, where
    they are called alpha1, alpha2). a1 and a2 can be symbolic or numeric. Try:
                                   ChemoStat(N,C,a1,a2);
                                    ChemoStat(N,C,2,3);
                                                                                                 (2)
\rightarrow TimeSeries (ChemoStat(20, 50, 2, 3), [x, y], [0.5, 0.5], 0.01, 10, 1)
```



> TimeSeries (ChemoStat(30, -50, 2, 3), [x, y], [0.5, 0.5], 0.01, 10, 1)



> TimeSeries (ChemoStat(-20, 5, 2, 3), [x, y], [0.5, 0.5], 0.01, 10, 1)



> *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 parameers a0 (called alpha_0 there), a (called alpha there), b (called beta there) and n. Try:

Gene
$$Net(0,0.5,0.2,2,m1,m2,m3,p1,p2,p3);$$
 (3)

> TimeSeries (GeneNet(0, 0.5, 0.2, 2, 10, 2, 3, 10, 2, 3), [x, y, z], [0.5, 0.5, 0.5], 0.01, 10, 1) bad input

(5)

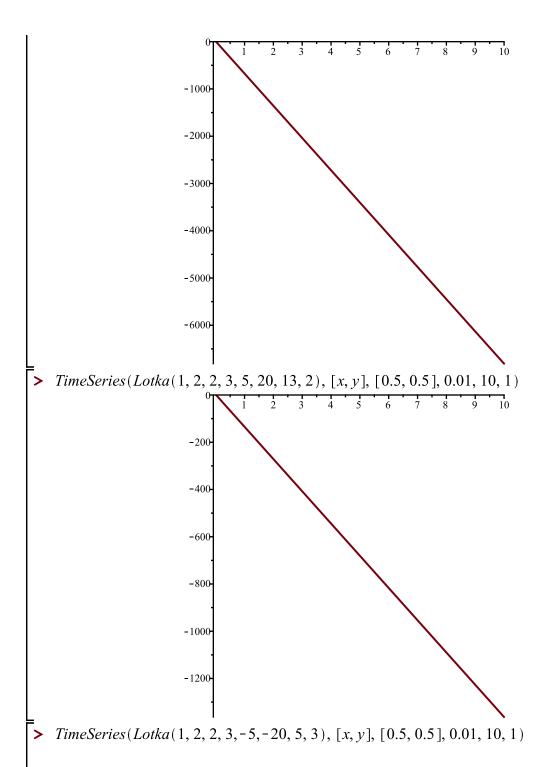
> Help(Lotka)

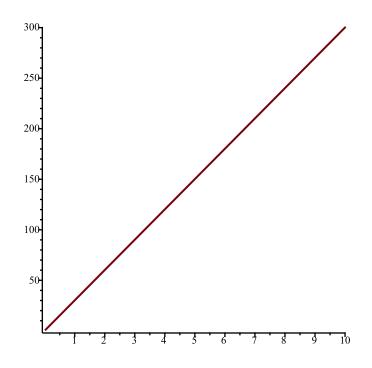
Lotka(r1,k1,r2,k2,b12,b21,N1,N2): The Lotka-Volterra continuous-time dynamical system, Eqs. (9a),(9b) (p. 224, section 6.3) of Edelstein-Keshet

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

Try:

> *TimeSeries*(*Lotka*(1, 2, 2, 3, 5, 20, 11, 23), [x, y], [0.5, 0.5], 0.01, 10, 1)





> Help(Volterra)

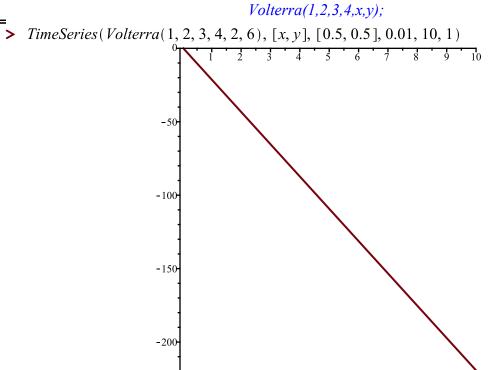
Volterra(a,b,c,d,x,y): The (simple, original) Volterra predator-prey continuous-time dynamical system with parameters a,b,c,d

Given by Eqs. (7a) (7b) in Edelstein-Keshet p. 219 (section 6.2). a,b,c,d may be symbolic or numeric

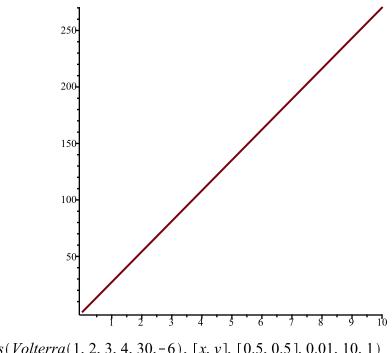
Try:

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

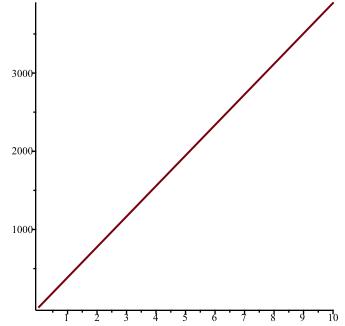
(6)



> *TimeSeries*(*Volterra*(1, 2, 3, 4, -3, 5), [x, y], [0.5, 0.5], 0.01, 10, 1)



> TimeSeries (Volterra (1, 2, 3, 4, 30, -6), [x, y], [0.5, 0.5], 0.01, 10, 1)



> 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

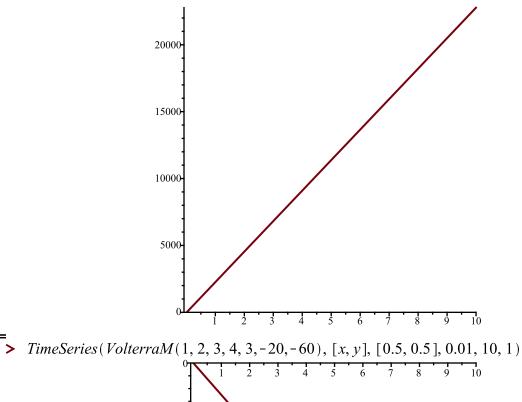
> Given by Eqs. (8a) (8b) in Edelstein-Keshet p. 220 (section 6.2). a,b,c,d,Kmay be symbolic or numeric

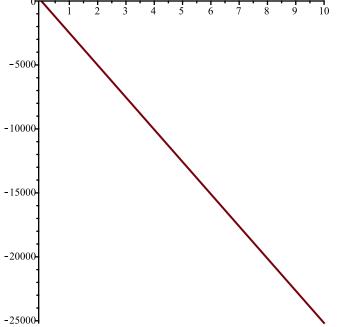
> > *Try:*

VolterraM(a,b,c,d,K,x,y);*VolterraM(1,2,3,4,3,x,y);*

 \rightarrow TimeSeries (VolterraM(1, 2, 3, 4, 3, -20, 60), [x, y], [0.5, 0.5], 0.01, 10, 1)

(7)





> *TimeSeries*(*VolterraM*(1, 2, 3, 4, 3, -2, 5), [x, y], [0.5, 0.5], 0.01, 10, 1)

