

MATHEMATICAL AND COMPUTER MODELING OF NONLINEAR BIOSYSTEMS I

COMPUTER LABORATORY III: : Discrete logistic equation (solutions,
bifurcations), two dimensional discrete models

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Discrete logistic equation

Let us consider the logistic map (discrete logistic equation)

$$x_{n+1} = rx_n(1 - x_n)$$

where $r \in (0, 4]$ and $x_0 \in [0, 1]$.

Value x_n is the ratio of existing population to the maximum possible population at year n .

The above equation became famous because it exhibits chaotic dynamics. In the series of exercises we will try to understand what does it mean.

Exercise 1 - Implementation of logistic map

Implement the logistic map

$$x_{n+1} = rx_n(1 - x_n)$$

as a MATLAB function, with the following features:

- three arguments: r , initial value, number of iterations to perform;
- optional boolean argument stating if the whole trajectory x_0, x_1, \dots, x_n or only the last element x_n should be returned. By default the function should return only the last value;
- initial value can be a vector and solution to the logistic map will be calculated for each element of that vector.

Exercise 1 - solution

```
function a = logisticMapVectInit( r, a0, n, traj )
    if nargin<4 %check if argument is passed
        traj = false;
    end
    a0=a0(:);
    if traj %true: return whole trajectory
        a=zeros(length(a0),n+1); %predefine output
        a(:,1)=a0;
        for i=1:n
            a(:,i+1)=r*a(:,i).*(1-a(:,i));
        end
    else
        a=a0;
        for i=1:n
            a=r*a.*(1-a);
        end
    end
end
end
```

Exercise 1a - Implementation of logistic map

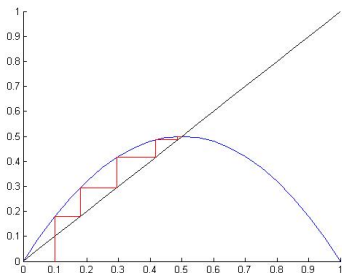
Implement the same function as in Exercise 1, but vectorized with respect to r instead of vectorization with respect to the initial condition.

Exercise 1 - solution

```
function a = logisticMapVectParam( r, a0, n, traj )
    if nargin<4 %check if argument is passed
        traj = false;
    end
    r=r(:); %make vertical vector
    if traj %true: return whole trajectory
        a=zeros(length(r),n+1); %predefine output
        a(:,1)=a0;
        for i=1:n
            a(:,i+1)=r.*a(:,i).*(1-a(:,i));
        end
    else
        a=a0*ones(length(r),1);
        for i=1:n
            a=r.*a.*(1-a);
        end
    end
end
end
```

Exercise 2 - solution illustration (cobweb plot)

Using the functions implemented in previous exercises write a MATLAB script which solves the logistic equation for specified parameters, initial conditions and number of iterations. Visualize solution using cobweb plot (Verhulst diagram).



cobweb plot

Exercise 2 - solution (part 1)

```
clear all
close all

%%%% Initial settings
r = 3.5;
a0 = 0.1;
n=10;
%%%%

sol=logisticMap(r,a0,n,true);
```


Exercise 2 - solution (part 2)

```
figure(1)
clf
hold on
%plotting RHS
plot(0:0.05:1,logisticMapVectInit(r,0:0.05:1,1));
%plotting auxiliary
plot(0:0.05:1,0:0.05:1,'k');
%plotting iterations
init = 0;
for i=2:n+1
    plot([sol(i-1) sol(i-1)],[init sol(i)],'r');
    plot([sol(i-1) sol(i)],[sol(i) sol(i)],'r');
    init = sol(i);
end
hold off
```

Exercise 3 - bifurcation diagram (Feigenbaum tree)

Plot the bifurcation diagram for the logistic map. Procedure description

- 1 define the uniform mesh for parameter r in the range $[1, 4]$;
- 2 for each mesh value of r solve the logistic equation starting from $x_0 = 0.5$ up to $n = 200$
- 3 plot the values of $x_{101}, x_{102}, \dots, x_n$ on the y-axis with the x-axis value equal to current r value.

Exercise 3 - naive solution

```
function FeigenbaumTreeNaive
END=1500; %number of mesh points
hold on
for i=1:END,
r = 1 + i*(3/END); a = 0.5;
for j=1:100,
a = r*a*(1-a);
end
for j=1:100,
a = r*a*(1-a);
plot(r,a);
end;
end;
hold off
end
```

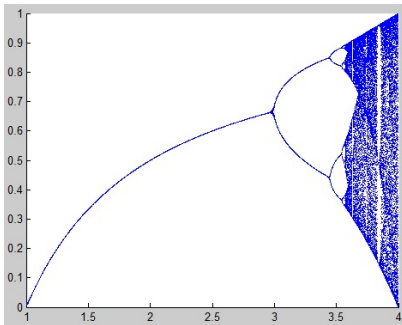
Exercise 3 - solution using previously implemented functions

```
function FeigenbaumTree(r, n)

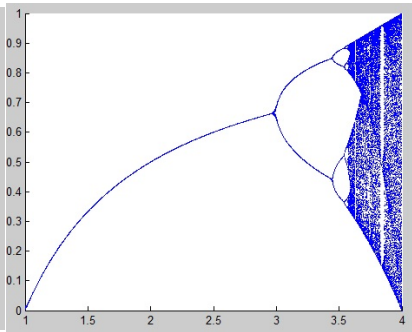
    Iter = logisticMapVectParam(r,0.5,2*n,true);

    h=scatter(repmat(r,1,n),reshape(Iter(:,n+2:end),1,[]),'.');
    hChildren = get(h, 'Children');
    set(hChildren, 'MarkerSize', 2)
end
```

Comparizon of solutions - 1500 mesh points for r



Naive: ≈ 76 seconds



More efficient: ≈ 1.6 seconds

Leslie model

Consider the population of individuals divided into ω age groups

$n_0, n_1, \dots, n_{\omega-1}$.

We may describe that population using the Leslie model

$$\begin{bmatrix} n_0 \\ n_1 \\ \vdots \\ n_{\omega-1} \end{bmatrix}_{t+1} = \begin{bmatrix} f_0 & f_1 & f_2 & f_3 & \dots & f_{\omega-1} \\ s_0 & 0 & 0 & 0 & \dots & 0 \\ 0 & s_1 & 0 & 0 & \dots & 0 \\ 0 & 0 & s_2 & 0 & \dots & 0 \\ 0 & 0 & 0 & \ddots & \dots & 0 \\ 0 & 0 & 0 & \dots & s_{\omega-2} & 0 \end{bmatrix} \begin{bmatrix} n_0 \\ n_1 \\ \vdots \\ n_{\omega-1} \end{bmatrix}_t$$

where parameter f_i describes the average number of offspring from the age class x and s_i is the fraction of individuals that survives from age class i to age class $i + 1$.

Exercise 4 - stable regions in Leslie model

Consider the Leslie model with 3 age groups described by the following transition matrix

$$M = \begin{bmatrix} a & b & 0.4 \\ 0.9 & 0 & 0 \\ 0 & 0.6 & 0 \end{bmatrix}$$

where a and b are within the range $[0, 1]$.

Plot the region of values of a and b for which the size of the population described by the above model doesn't go to infinity when time goes to infinity.

Exercise 4 - naive solution (simulations)

```
clear all;
close all;
%stabilizacja
M=[0 0 0.4; 0.9 0 0;0 0.6 0];

a=linspace(0,1,100);
b=linspace(0,1,100);

%1st version - pure simulations
tic
nIt=150;
dY=zeros(length(a),length(b));
tol=1e-4;
for i=1:length(a)
    for j=1:length(b)
        M(1,1)=a(i);
        M(1,2)=b(j);
```


Exercise 4 - naive solution (simulations)

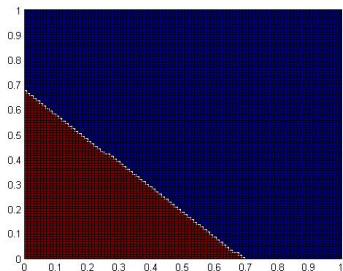
```
Y=[65; 70; 60];
for ii=1:nIt
    Yn=M*Y;
    diff=abs(sum(Y)-sum(Yn));
    Y=Yn;
    if Y>1e10
        break;
    end
end
dY(i,j)=diff;
end
figure(1)
clf
surface(a,b,double(dY<tol))
toc
```

Exercise 4 - solution using eigenvalues

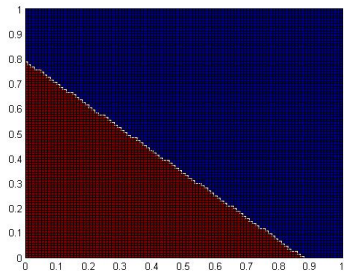
```
tic
mEig=zeros(length(a),length(b));
for i=1:length(a)
    for j=1:length(b)
        M(1,1)=a(i);
        M(1,2)=b(j);

        mEig(i,j)=max(real(eig(M)));
    end
end
figure(2)
clf
surface(a,b,double(mEig<=1));
toc
```

Comparison of both approaches



Naive: ≈ 16 seconds
Highly dependent on settings!!!



More efficient: ≈ 0.4 seconds