

# Various Type of HIV Infected Virus

P. Kalaiyarasi<sup>1</sup>, K. M. Manikandan<sup>2</sup>

<sup>1</sup>M.Sc Mathematics Department of Mathematics, Dr. SNS Rajalakshmi College of Arts and Science, Coimbatore, Tamil Nadu, India

<sup>2</sup>HOD, Assistant Professor, Department of Mathematics, Dr. SNS Rajalakshmi College of Arts and Science, Coimbatore, Tamil Nadu, India

## ABSTRACT

In this paper, we have reviewed the equation of human immune deficiency virus (HIV) delayed model with cure of infected cells in the eclipse phase described by ordinary differential equation (ODE). In addition we obtain the graph showing the stages of healthy cells, infected cells but not spread the virus, and infected cells spreading the virus and virus cells by using MATLAB and successive approximation method.

**Keywords :** HIV, MATLAB, Successive Approximation Method

## I. INTRODUCTION

The purpose of this work is to investigate the dynamical behavior of the following HIV infection model governed by the ordinary differential equation model which given by

$$\begin{aligned} \frac{dH}{dt} &= \mu - \lambda_H H(t) - f(H(t), R(t))R(t) + \rho L(t) \\ \frac{dL}{dt} &= f(H(t), R(t))R(t) - (\lambda_L + \rho + \gamma)L(t) \\ \frac{dN}{dt} &= \gamma L(t) - \lambda_N N(t) \\ \frac{dR}{dt} &= P e^{-m\tau} N(t-\tau) - \lambda_R R(t) \end{aligned} \quad (1)$$

The first equation of (1) describes the dynamics of the concentration of healthy cells (H).  $\mu$  is the recruitment rate and  $\lambda_H$  is the death rate of uninfected cells. During infection, healthy cells decreases proportionally to the product  $f(H, R)R$ . The second equation represents the dynamics of the concentration of the infected cells in eclipse stage. (i.e) infected that are not yet producing virus.  $\lambda_L$  is a die at rate,  $\rho$  is a productive rate infected cells. The third equation represents dynamics of the concentration productive infected cells N.  $\lambda_N$  is a dieing rate. The last equation represents dynamics of the concentration of viruses which are produced by

infected cells at the rate P,  $\lambda_R$  is a dieing a rate. The delay  $\tau$  describes the time needed for productive infected cells which are produced virions and  $e^{-m\tau}$  is the probability of surviving from time  $(t-\tau)$  to time t. The model is disease transmission process by hataf's incidence rate

$$F(N, R) = \frac{\alpha N}{1 + \delta_1 N + \delta_2 R + \delta_3 NR}$$

Where  $\delta_1, \delta_2, \delta_3 \leq 0$  are constants and  $\alpha$  is the infection rate.

## II. BASIC RESULTS

We can easily show that there exists a unique solution  $(H(t), L(t), N(t), R(t))$  for equation (1) initial condition  $(H_0, L_0, N_0, R_0)$  for a biological reason we assume this condition to satisfy

$$\begin{aligned} H_0(m) \leq 0, L_0(m) \leq 0, N_0(m) \leq 0, R_0(m) \leq 0 \\ \text{for } m \in (-\tau, 0) \end{aligned} \quad (2)$$

### THEOREM.2.1:

The solution of equation (1) satisfying condition (2) remains non negative and bounded for all  $t \leq 0$

**Proof:**

It is easy way to show that non negative of the solution of system (1) with initial condition satisfying (2).show that the boundedness of solution

We define

$$\begin{aligned}
 A(t) &= H(t) + L(t) + N(t) \\
 &= \mu - \lambda_H H(t) - f(H(t), R(t))R(t) + \rho L(t) + \lambda_N N(t) \\
 &= \mu - \lambda_H H(t) - \lambda_L L(t) - \rho L(t) - \gamma L(t) + \gamma L(t) - \lambda_N N(t) \\
 &= \mu - \lambda_H H(t) - \lambda_L L(t) - \lambda_N N(t) \\
 \frac{dA}{dt} &= \mu - \lambda_H H(t) - \lambda_L L(t) - \lambda_N N(t) \\
 &\geq \mu - \lambda A(t)
 \end{aligned}$$

Where  $\lambda = \max(\lambda_H, \lambda_L, \lambda_N)$

Here  $A(t) \geq \min\{A(0), \frac{\mu}{\lambda}\}$

Therefore  $H(t), L(t), N(t), R(t)$  are bounded.

Other hand

$$\begin{aligned}
 \frac{dR}{dt} &= Pe^{-m\tau}N(t-\tau) - \lambda_R R(t) \\
 &\geq Pe^{-m\tau}\|N\|_{\infty} - \lambda_R R(t)
 \end{aligned}$$

$$R(t) = \min\{R(0), \frac{Pe^{-m\tau}}{\lambda_R} \|N\|_{\infty}\}$$

We define  $R(t)$  is bounded

Graph for showing stages of healthy cells , infected cells but not spread the virus, and infected cells spreading the virus and virus cells by using MATLAB and successive approximation method.

**1.HEALTHY CELLS:**

$$H' = H, H(0) = 1$$

$$H' = f(s, H(s))$$

$$H_0 = 1 \text{ when } t_0 = 0$$

$$H_{n+1} = H_0 + \int_{t_0}^t f(s, H_n(s)) ds$$

When  $n=0$

$$\begin{aligned}
 H_1 &= H_0 + \int_{t_0}^t H_0 dt \\
 &= H_0 + [s]_{t_0}^t
 \end{aligned}$$

$$H_1(t) = 1 + t$$

$$H_2 = H_0 + \int_{t_0}^t f(s, H_1(s)) ds$$

$$= 1 + \int_{t_0}^t (1 + s) ds$$

$$= 1 + [t + \frac{t^2}{2}]$$

$$H_n(t) = 1 + t + \frac{t^2}{2}$$

**SUCCESSIVE APPROXIMATION METHOD BY USING MATLAB CODING**

```

1 %dsbydt=H=f(s,H(s)), H(0)=1
2 - t=0;%left end point of domain
3 - t0=0.5;%right end point of domain
4 - N=2;%number
5 - nosuccapr=2;%no of successive aprosimations
6 - sarray = linspace(t,t0,N);%array of domain from t to t0 consists of N points
7 - disp(sarray);
8 - Harray = sarray + 1; %initial f(s,L(s))
9 - for apros = 1:nosuccapr
10 -     yofs = trapzule(sarray,Harray,N)+1;%y(s)=y(a)+ integral a to s
11 -     Harray =sarray+ yofs;%new f(s,H(s));=s
12 -     disp(Harray);
13 -     plot(sarray,Harray,'green');%for plotting
14 -     hold on;
15 - end
    
```

**Figure 1**

```

1 - function newyarray = traprule(Harray,Sarray,N)
2     %str = sprintf('%8.6f integral',integral);
3     %disp(str);
4
5     h=Sarray(2)-Harray(1);
6     newyarray=Sarray;
7     for i = 1:length(Harray)
8         newyarray(i)=trapz(Harray(1:i),Sarray(1:i));
9     end

```

Figure 2

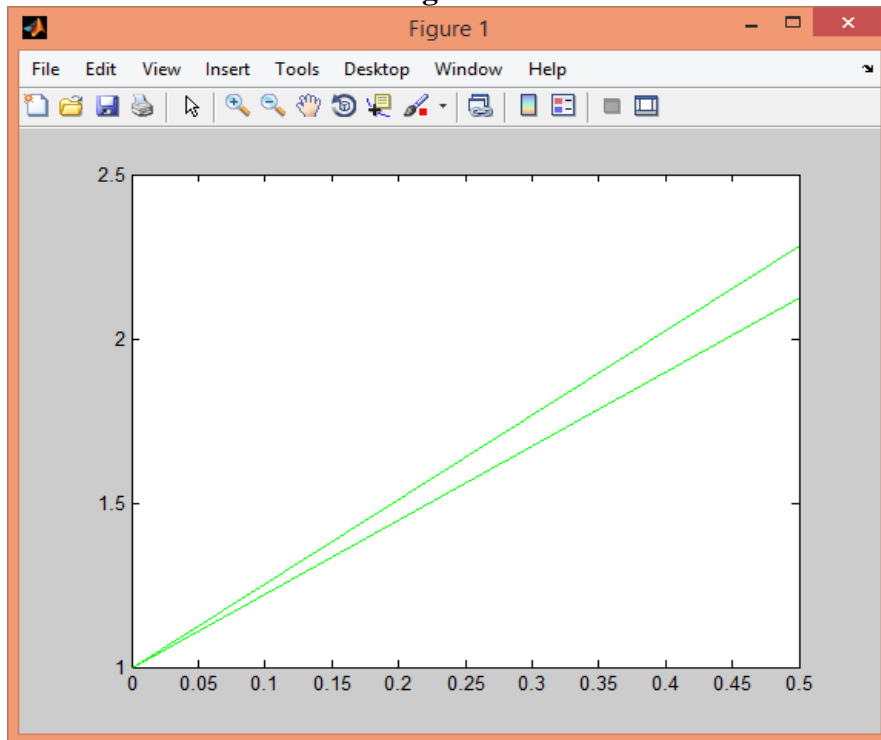


Figure 3

**2.INFECTED CELLS BUT NOT YET PRODUCING VIRUS:**

$$L' = L, L(0) = 1$$

$$L' = f(s, L(s))$$

$$L_0 = 2 \text{ when } t_0 = 1$$

$$L_{n+1} = L_0 + \int_{t_0}^t f(s, L_n(s)) ds$$

When n=1

$$L_1 = L_0 + \int_{t_0}^t L_0 dt$$

$$= 2 + [s]_1^t$$

$$= 2 + [t - 1]$$

$$L_1(t) = 1 + t$$

$$L_2 = L_0 + \int_{t_0}^t f(s, L_1(s)) ds$$

$$= 2 + \int_{t_0}^t (1 + s) ds$$

$$= 2 + [s + \frac{s^2}{2}]_{t_0}^t$$

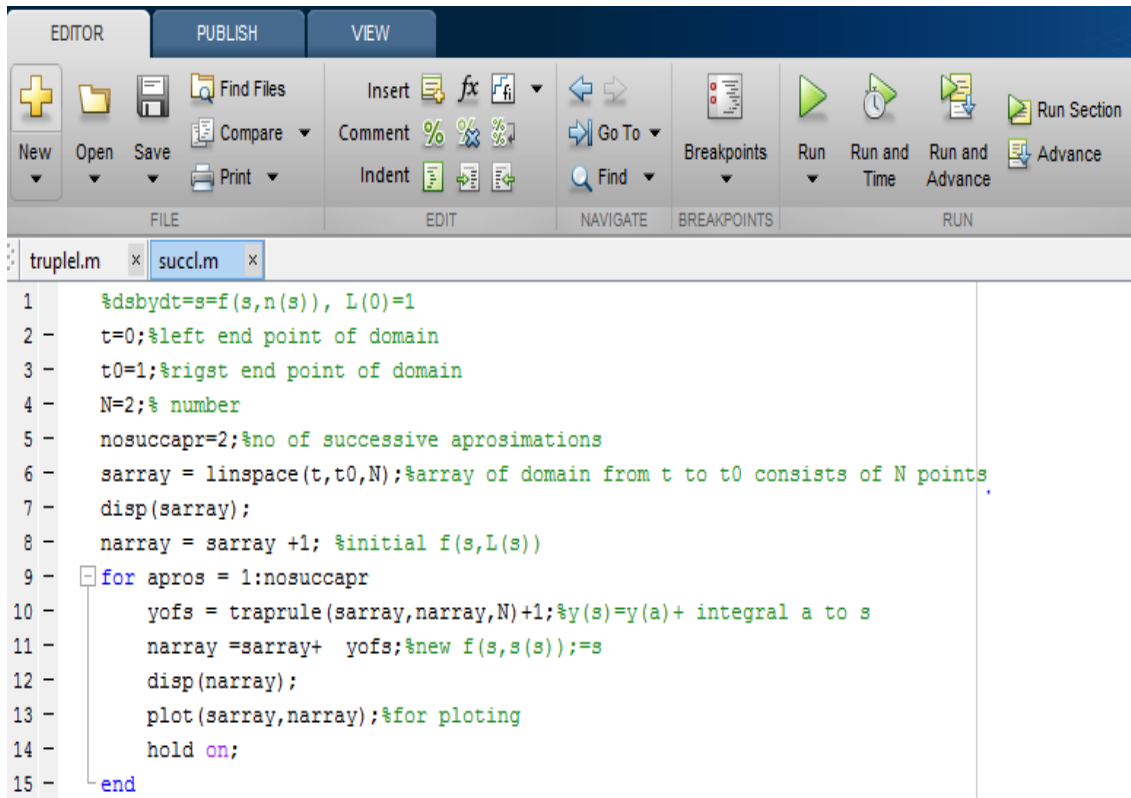
$$= 2 + [s + \frac{s^2}{2}]_1^t$$

$$= 2 + [(t + \frac{t^2}{2}) - (1 + \frac{1}{2})]$$

$$=2+\left[t+\frac{t^2}{2}+\frac{3}{2}\right]$$

$$L_n(t)=t^2 + 2t - \frac{1}{2}$$

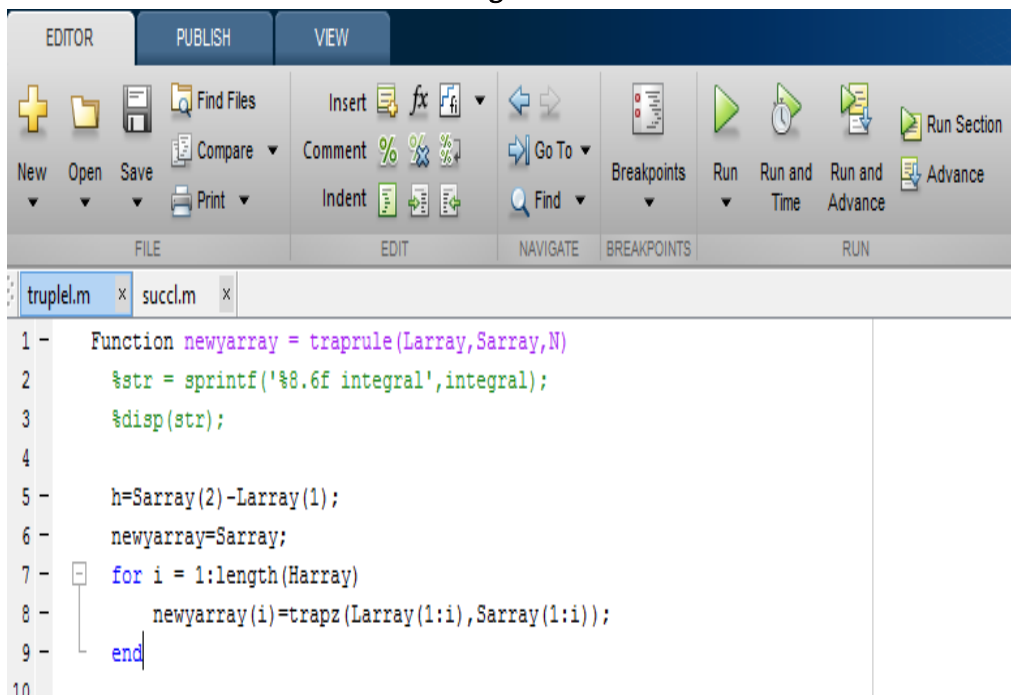
### SUCCESSIVE APPROXIMATION METHOD BY USING MATLAB CODING



```

1 %dsbydt=s=f(s,n(s)), L(0)=1
2 t=0;%left end point of domain
3 t0=1;%right end point of domain
4 N=2;% number
5 nosuccapr=2;%no of successive aprosimations
6 sarray = linspace(t,t0,N);%array of domain from t to t0 consists of N points
7 disp(sarray);
8 narray = sarray +1; %initial f(s,L(s))
9 for apros = 1:nosuccapr
10     yofs = traprule(sarray,narray,N)+1;%y(s)=y(a)+ integral a to s
11     narray =sarray+ yofs;%new f(s,s(s));=s
12     disp(narray);
13     plot(sarray,narray);%for plotting
14     hold on;
15 end
    
```

Figure 4



```

1 Function newyarray = traprule(Larray,Sarray,N)
2 %str = sprintf('%0.6f integral',integral);
3 %disp(str);
4
5 h=Sarray(2)-Larray(1);
6 newyarray=Sarray;
7 for i = 1:length(Harray)
8     newyarray(i)=trapz(Larray(1:i),Sarray(1:i));
9 end
10
    
```

Figure 5

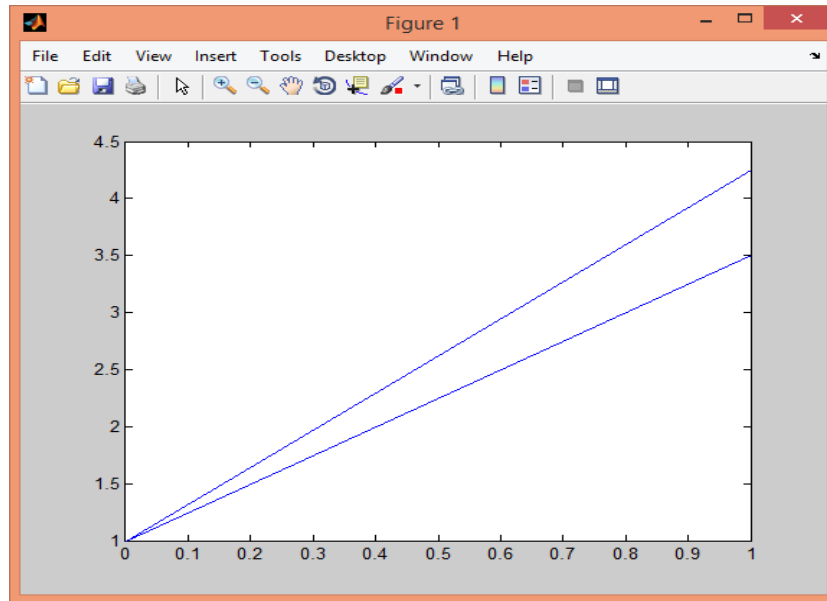


Figure 6

### 3. INFECTED CELLS SPREADS THE VIRUS:

$$N' = N, N(0) = 1$$

$$N' = f(s, N(s))$$

$$N_0 = 3 \text{ when } t_0 = 2$$

$$N_{n+1} = L_0 + \int_{t_0}^t f(s, N_n(s)) ds$$

When  $n=1$

$$N_1 = N_0 + \int_{t_0}^t N_0 dt$$

$$= 3 + [s]_2^t$$

$$= 3 + [t - 2]$$

$$N_1(t) = 1 + t$$

$$N_2 = N_0 + \int_{t_0}^t f(s, N_1(s)) ds$$

$$= 3 + \int_{t_0}^t (1 + s) ds$$

$$= 3 + [s + \frac{s^2}{2}]_{t_0}^t$$

$$= 3 + [s + \frac{s^2}{2}]_2^t$$

$$= 3 + [(t + \frac{t^2}{2}) - (2 + \frac{4}{2})]$$

$$= 3 + [\frac{2t + t^2 - 8}{2}]$$

$$N_n(t) = [\frac{t^2 + 2t - 2}{2}]$$

### SUCCESSIVE APPROXIMATION METHOD BY USING MATLAB CODING

```

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succN.m  x  truplen.m  x
1  %dsbydt=n=f(s,n(s)), n(0)=1
2  t=0;%left end point of domain
3  t0=2;%right end point of domain
4  N=3;% number
5  nosuccapr=2;%no of successive aprosimations
6  sarray = linspace(t,t0,N);%array of domain from t to t0 consists of N points
7  disp(sarray);
8  narray = sarray +1; %initial f(s,L(s))
9  for apros = 1:nosuccapr
10     yofs = traprule(sarray,narray,N)+1;%y(s)=y(a)+ integral a to s
11     narray =sarray+ yofs;%new f(s,s(s));=s
12     disp(narray);
13     plot(sarray,narray);%for plotting
14     hold on;
15  end
16
    
```

Figure 7

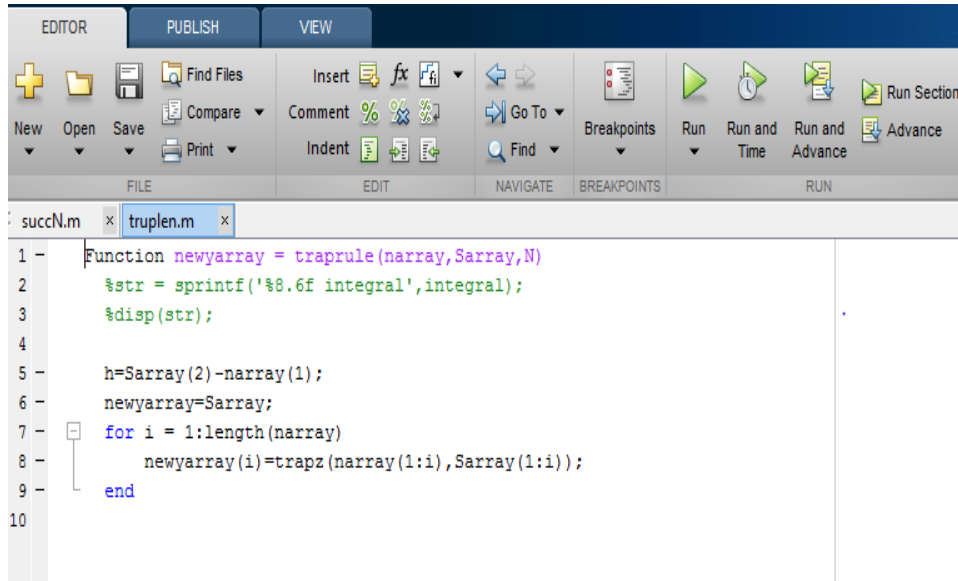


Figure 8

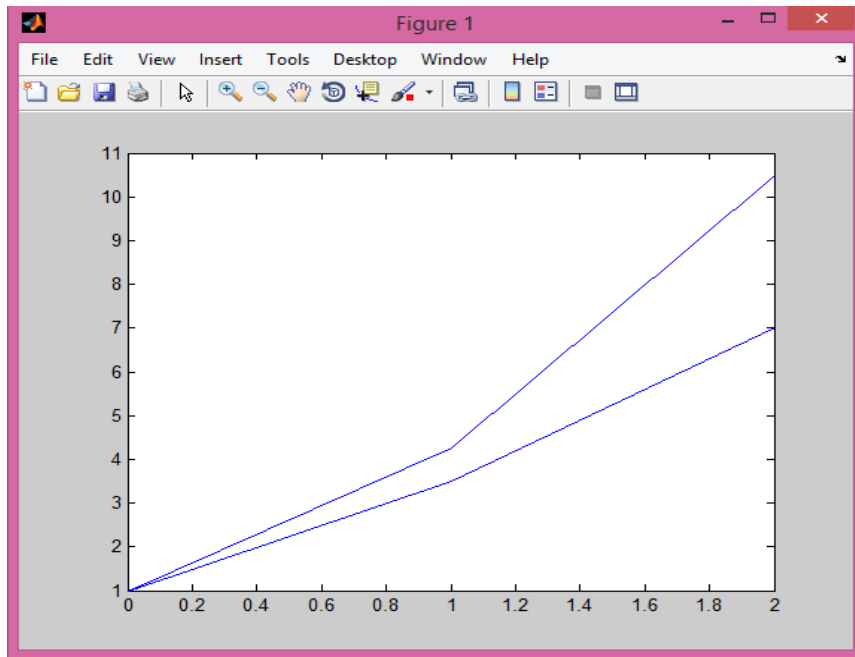


Figure 9

**4.VIRUS CELLS;**

$$R'=R,R(0)=1$$

$$R'=f(s, R(s))$$

$$R_0 = 4 \text{ when } t_0=3$$

$$N_{n+1}=L_0+\int_{t_0}^t f(s, N_n(s))ds$$

When n=1

$$R_1 = R_0 + \int_{t_0}^t R_0 dt$$

$$=4+[s]_3^t$$

$$=4+[t-3]$$

$$R_1(t)=1+t$$

$$R_2=R_0+\int_{t_0}^t f(s, R_1(s))ds$$

$$=4+\int_{t_0}^t (1+s)ds$$

$$=4+[s + \frac{s^2}{2}]_{t_0}^t$$

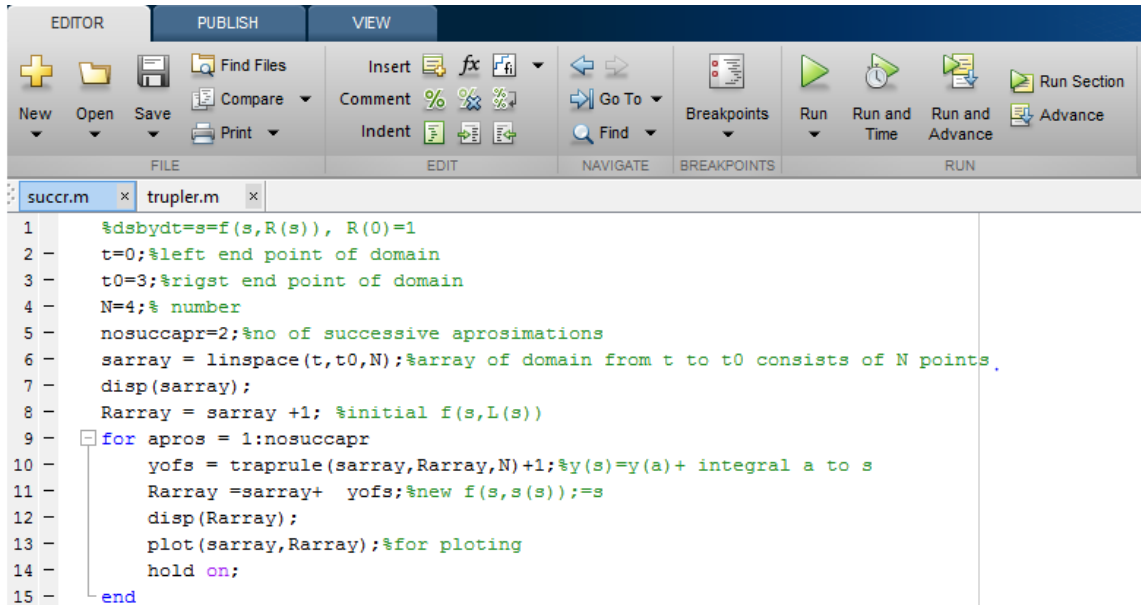
$$=4+[s + \frac{s^2}{2}]_3^t$$

$$=4+[(t+\frac{t^2}{2})-(3+\frac{9}{2})]$$

$$=4+[\frac{t^2+2t-15+8}{2}]$$

$$R_n(t)= [\frac{t^2+2t-7}{2}]$$

## SUCCESSIVE APPROXIMATION METHOD BY USING MATLAB CODING

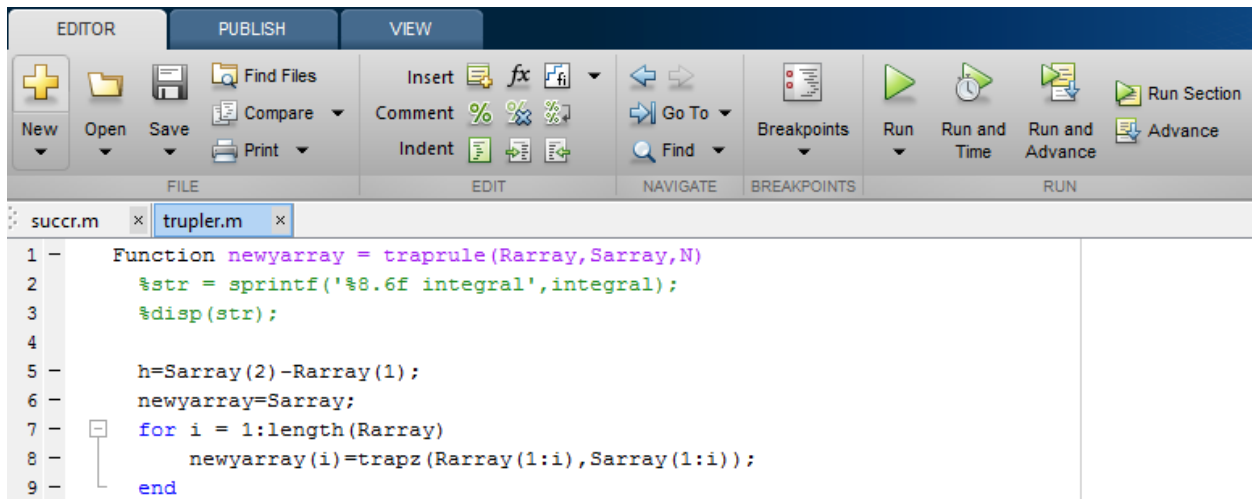


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succr.m x trupler.m x
1 %sdydt=s=f(s,R(s)), R(0)=1
2 t=0;%left end point of domain
3 t0=3;%right end point of domain
4 N=4;% number
5 nosuccapr=2;%no of successive aprosimations
6 sarray = linspace(t,t0,N);%array of domain from t to t0 consists of N points
7 disp(sarray);
8 Rarray = sarray +1; %initial f(s,L(s))
9 for apros = 1:nosuccapr
10     yofs = traprule(sarray,Rarray,N)+1;%y(s)=y(a) + integral a to s
11     Rarray =sarray+ yofs;%new f(s,s(s));=s
12     disp(Rarray);
13     plot(sarray,Rarray);%for plotting
14     hold on;
15 end
    
```

Figure 10

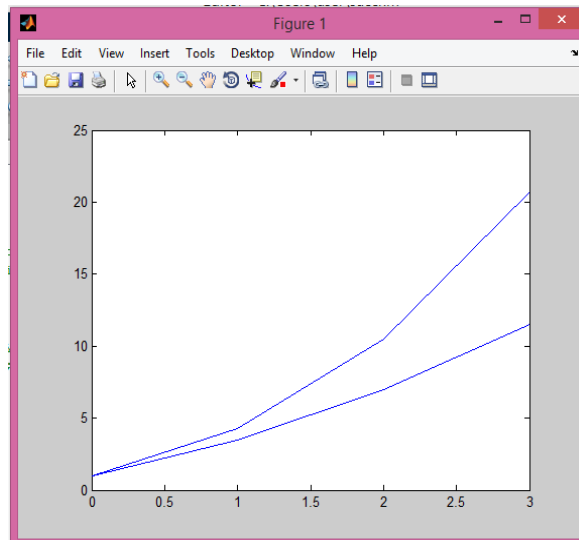


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succr.m x trupler.m x
1 Function newyarray = traprule(Rarray,Sarray,N)
2 %str = sprintf('%8.6f integral',integral);
3 %disp(str);
4
5 h=Sarray(2)-Rarray(1);
6 newyarray=Sarray;
7 for i = 1:length(Rarray)
8     newyarray(i)=trapz (Rarray(1:i),Sarray(1:i));
9 end
    
```

Figure 11



**Figure 12**

**III. CONCLUSION**

In this paper we showed the graph of various types of infected cells that produce virus. In this we used the MATLAB Program and successive approximation method and hence we formulated graph.

**IV. REFERENCES**

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