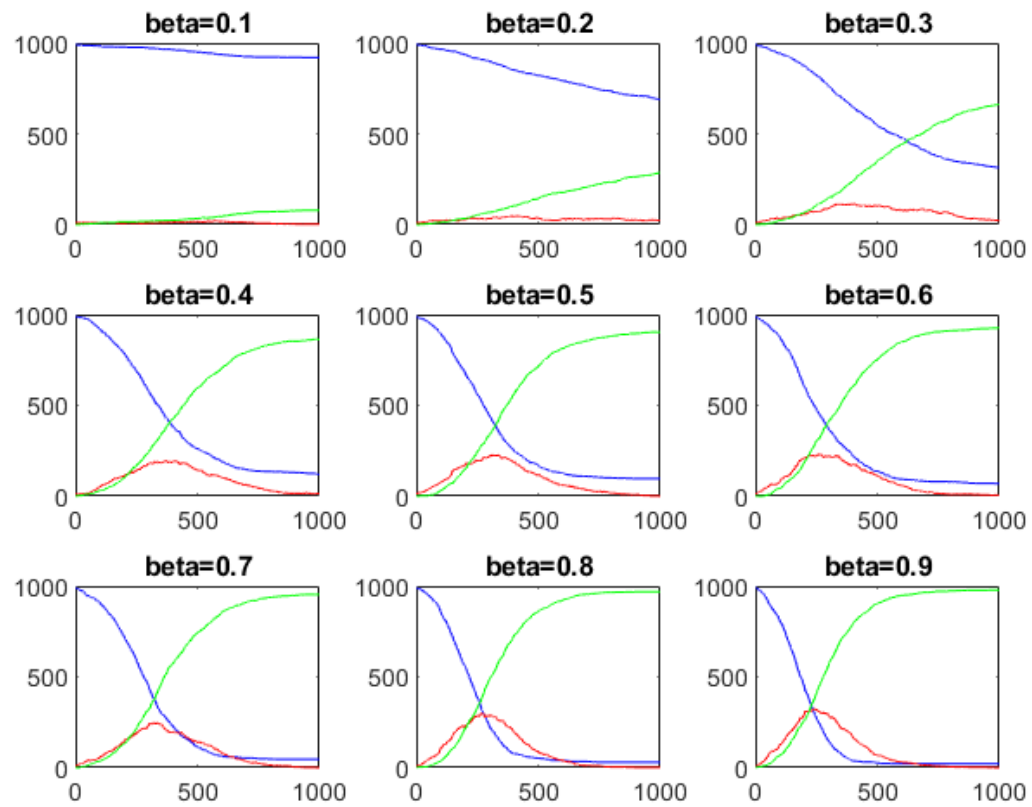


## 11.1

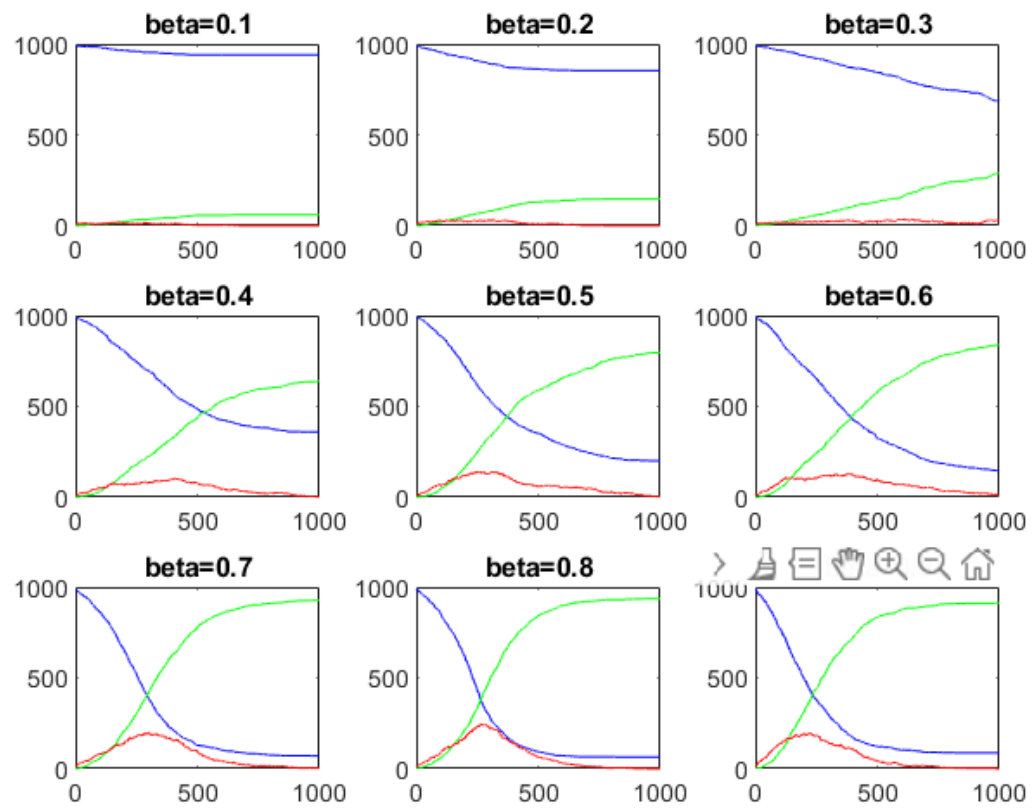
a)-c)

This is for  $\gamma=0.01$

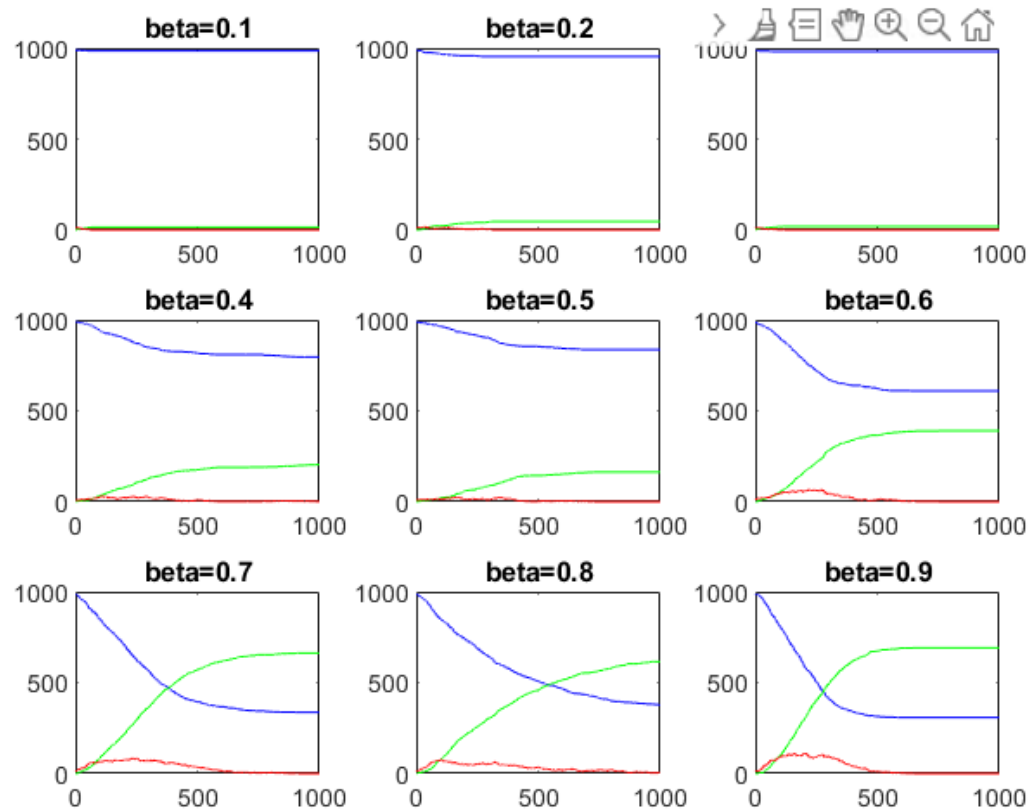


d)

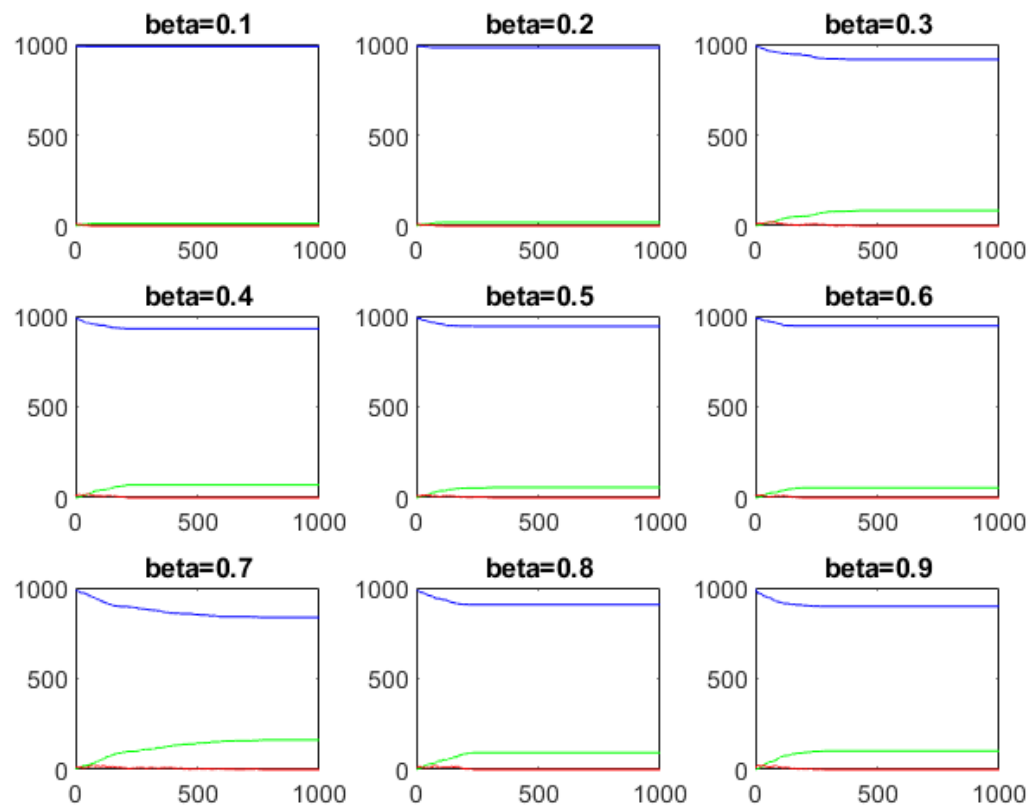
$\gamma=0.015$



Gamma=0.02:

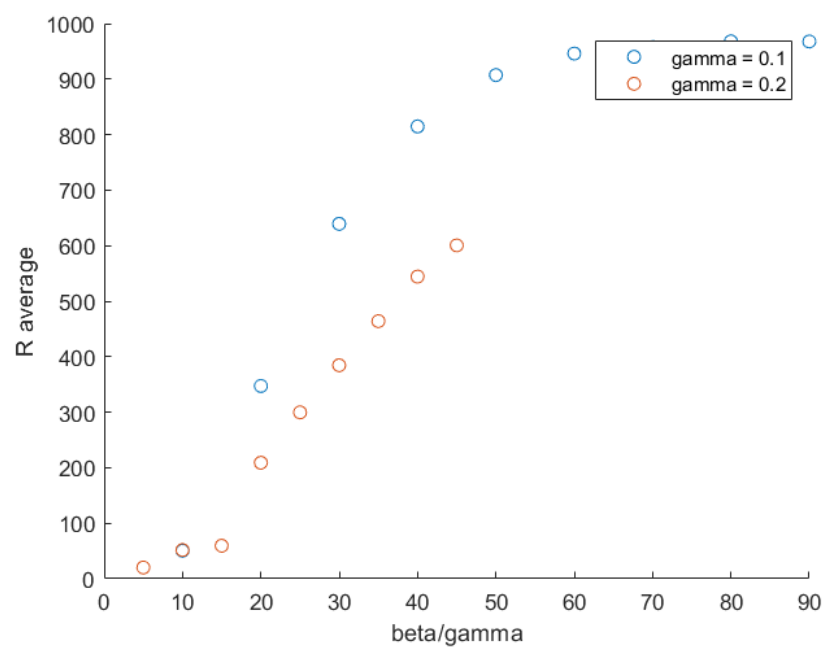
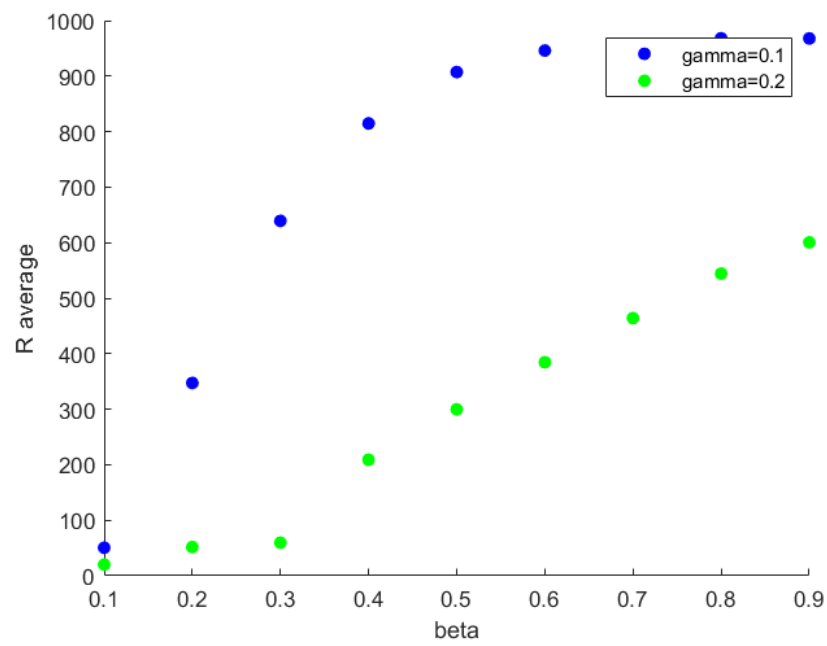


Gamma=0.03

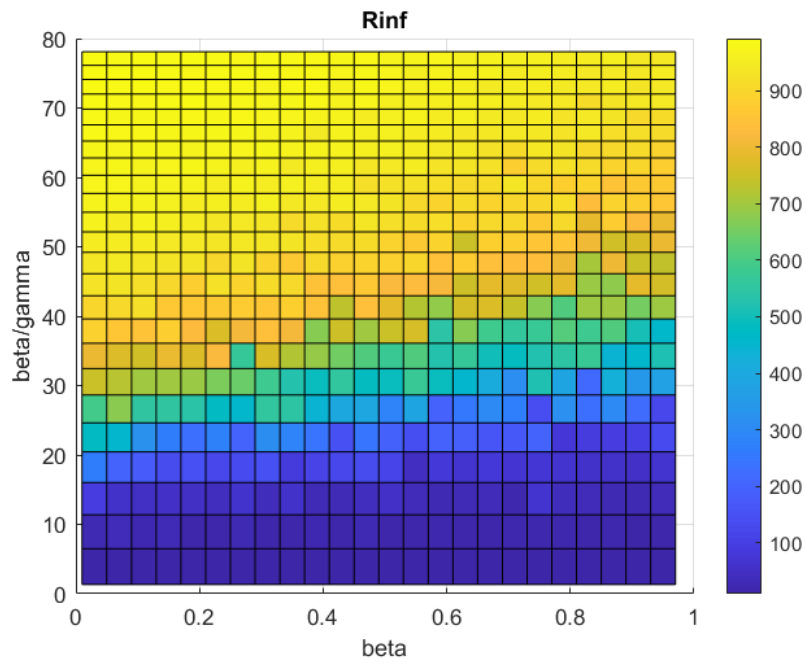


11.2

a)-b)



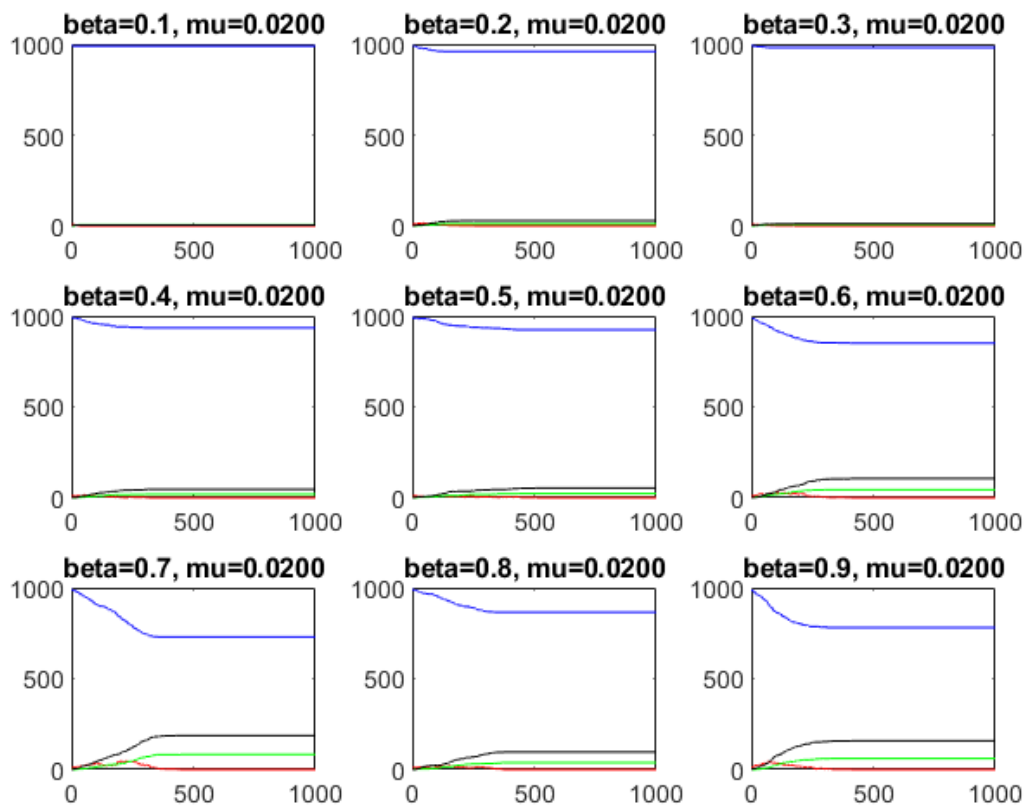
c)



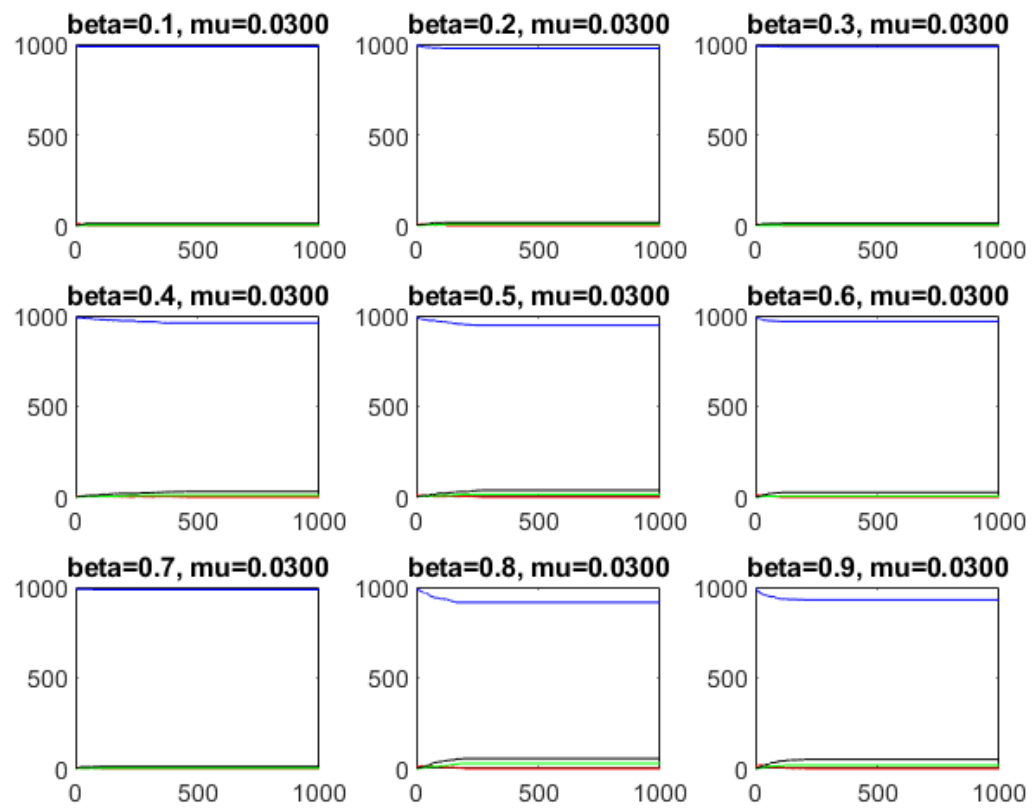
### 11.3)

$\Gamma=0.01$  for all

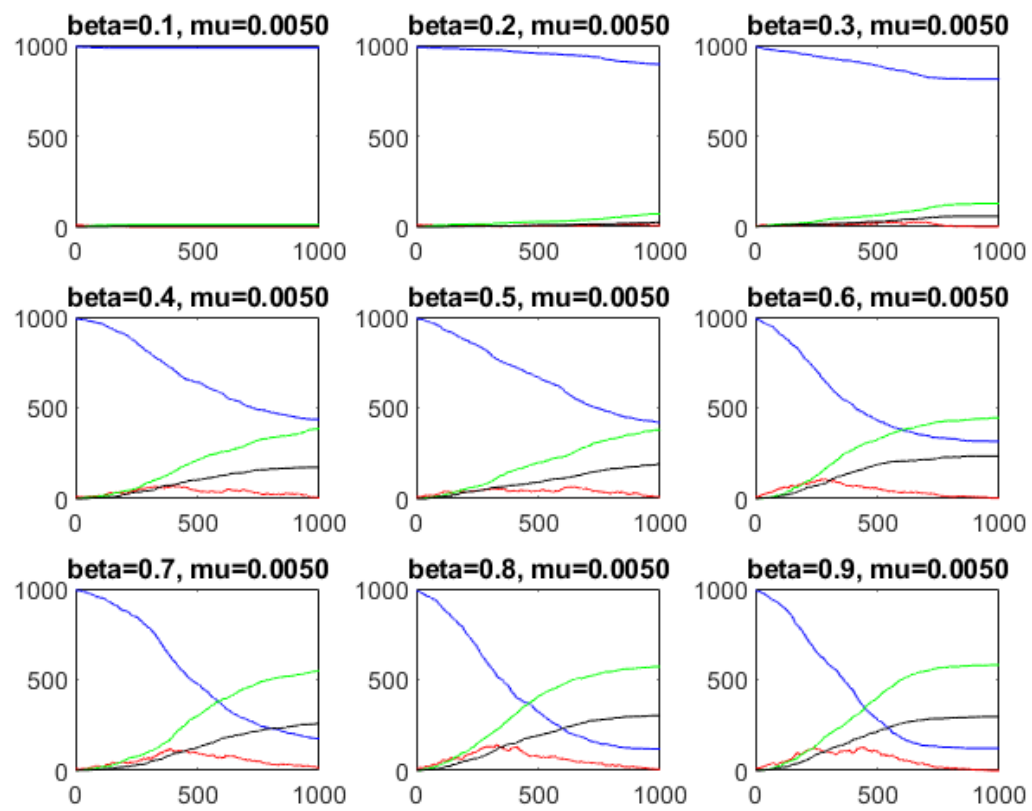
$\mu=0.02$



$\mu=0.03$

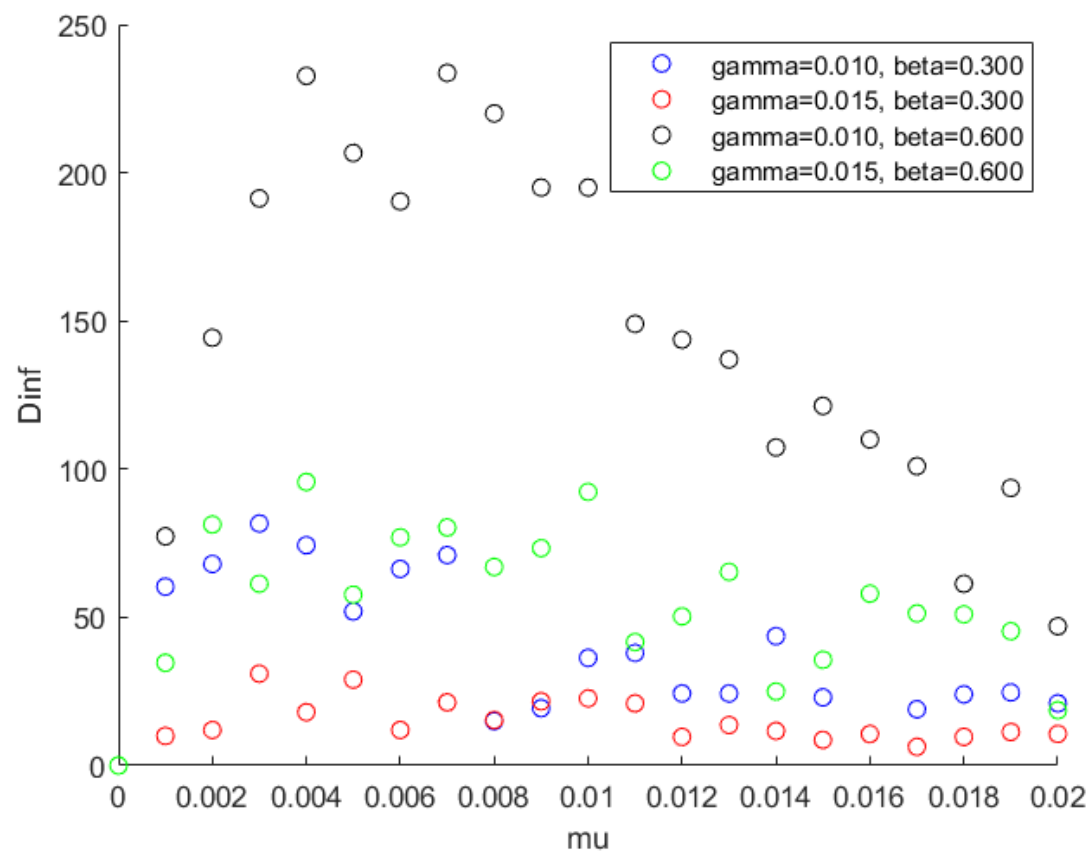
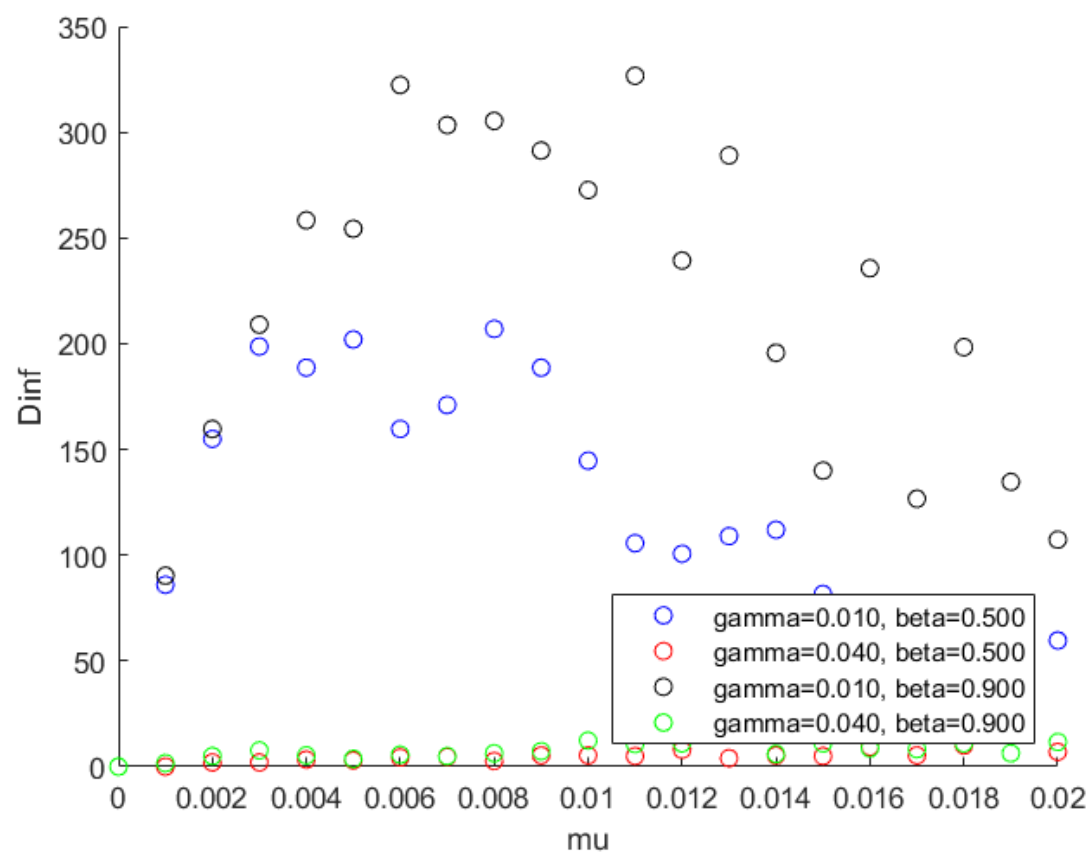


$\mu=0.005$



b)

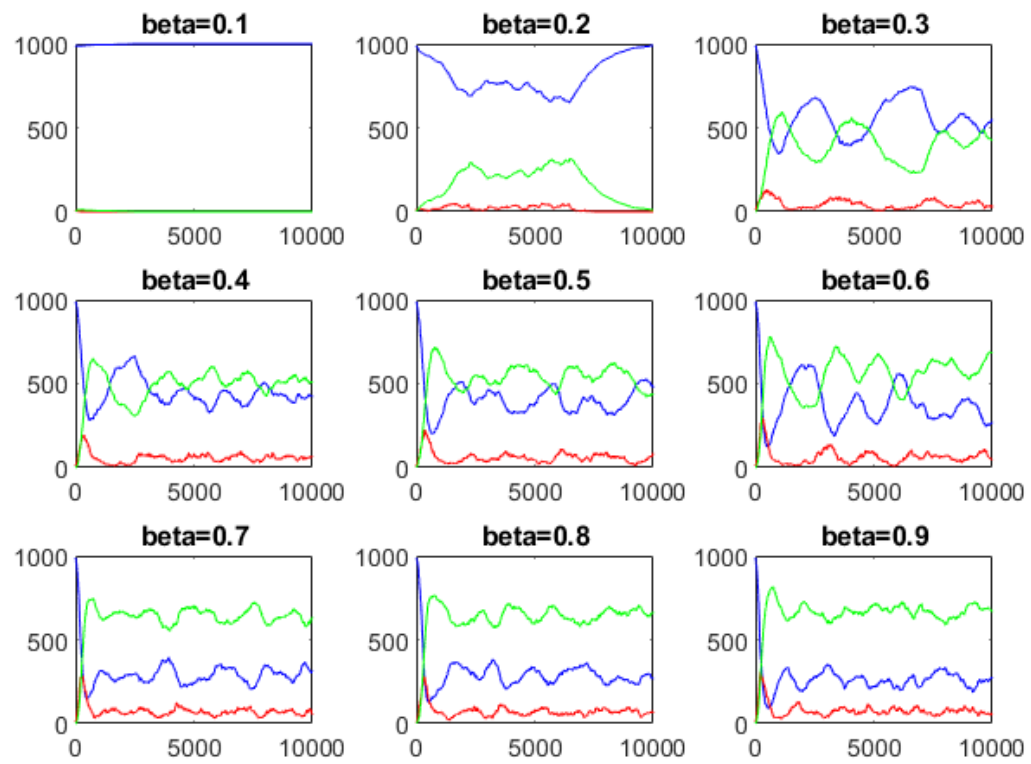




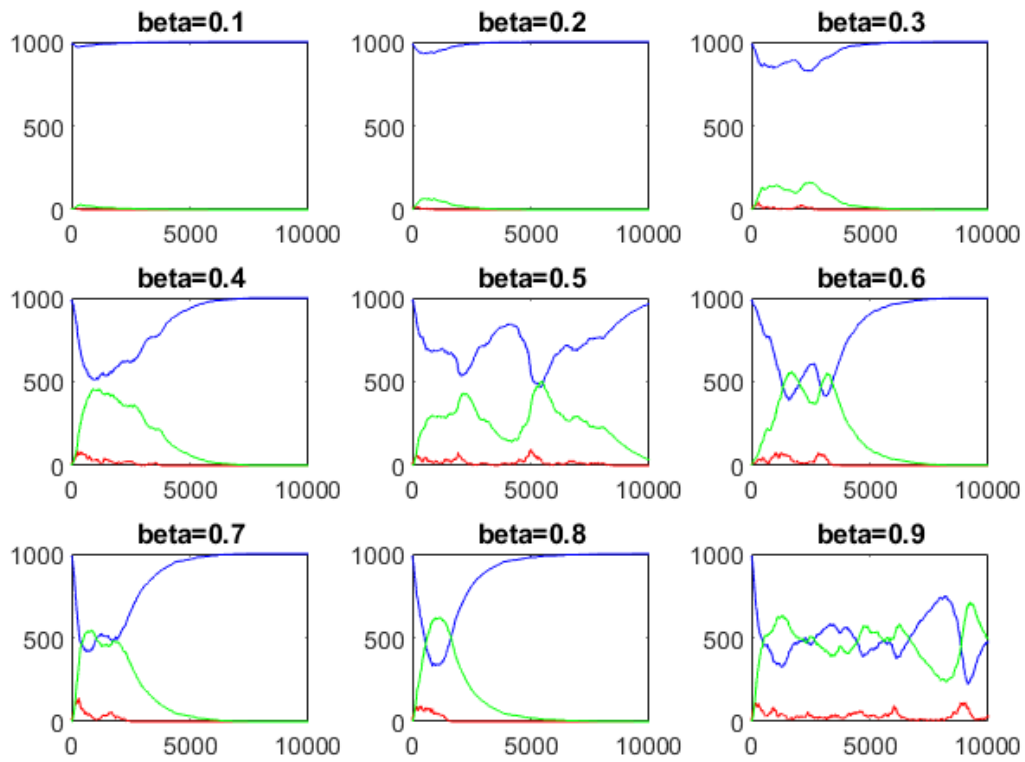
11.4

a)-b)

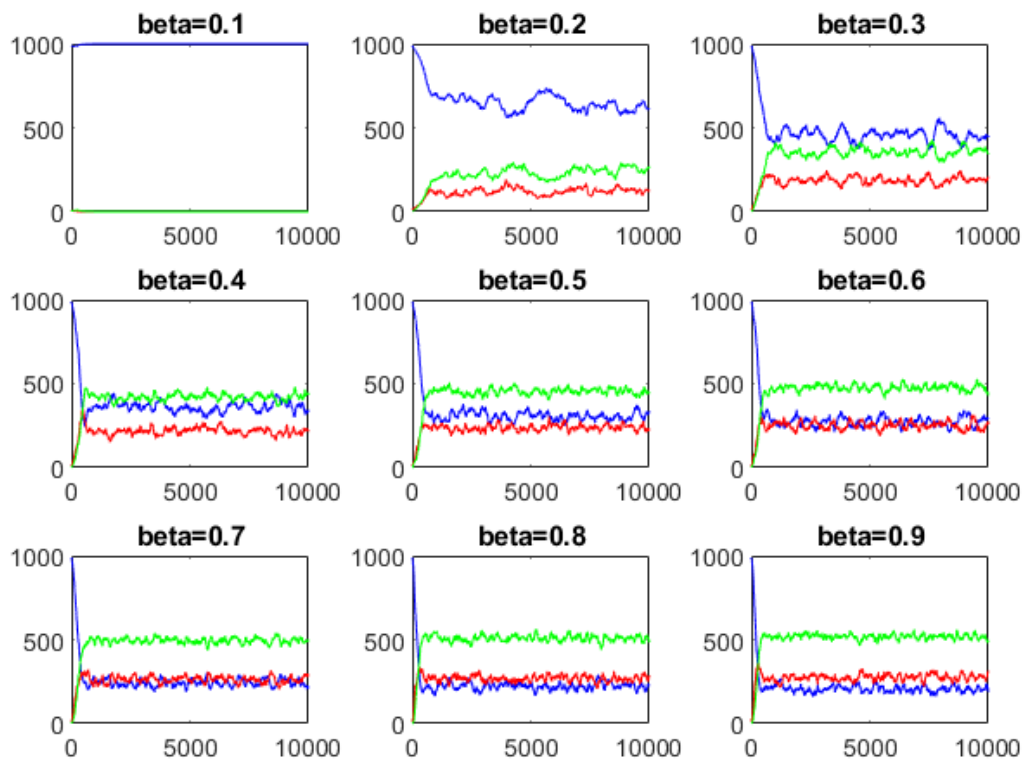
$\alpha=0.0010$ ,  $\gamma=0.010$



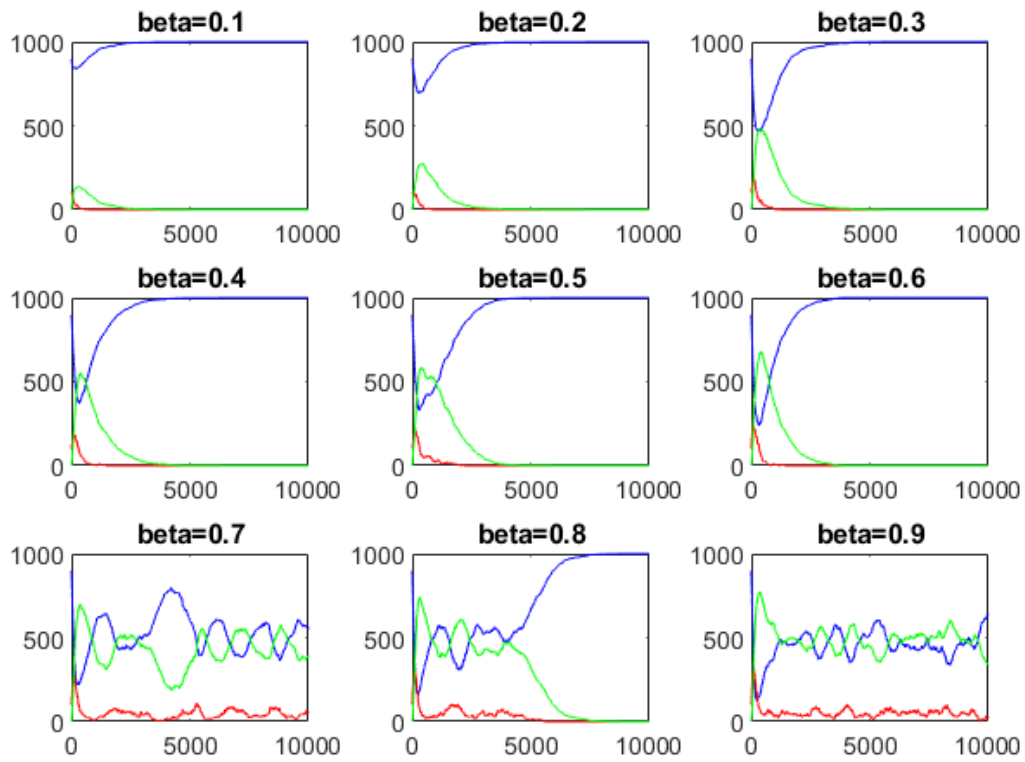
$\alpha=0.0010$ ,  $\gamma=0.015$



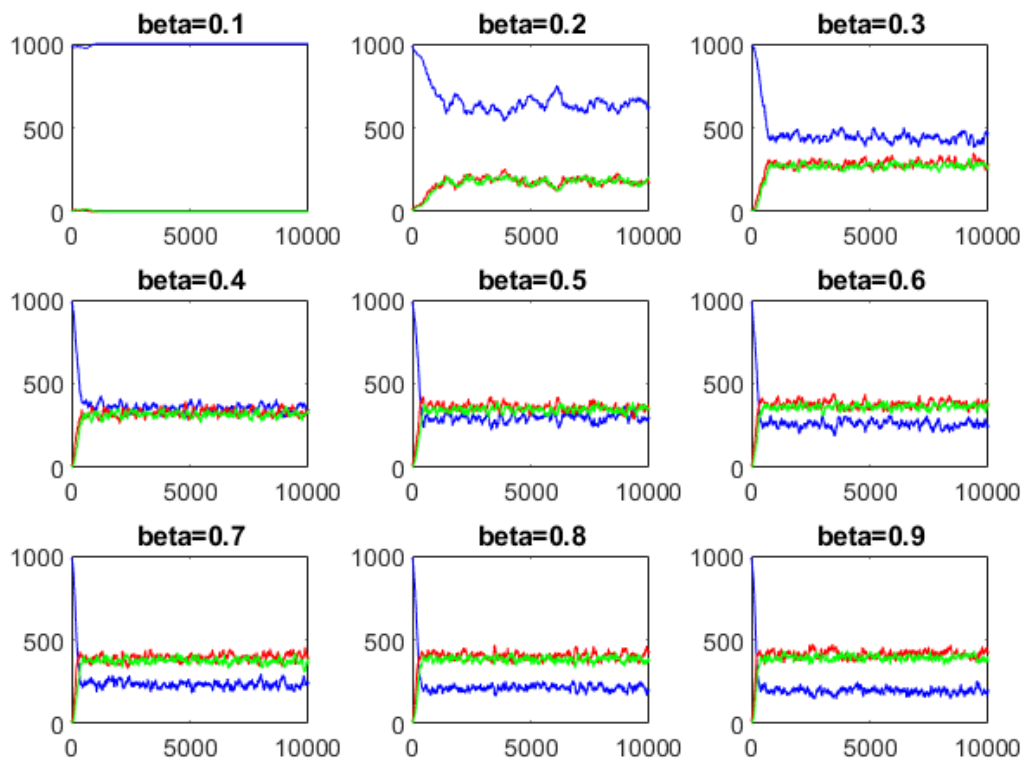
$\alpha=0.0050$ ,  $\gamma=0.010$



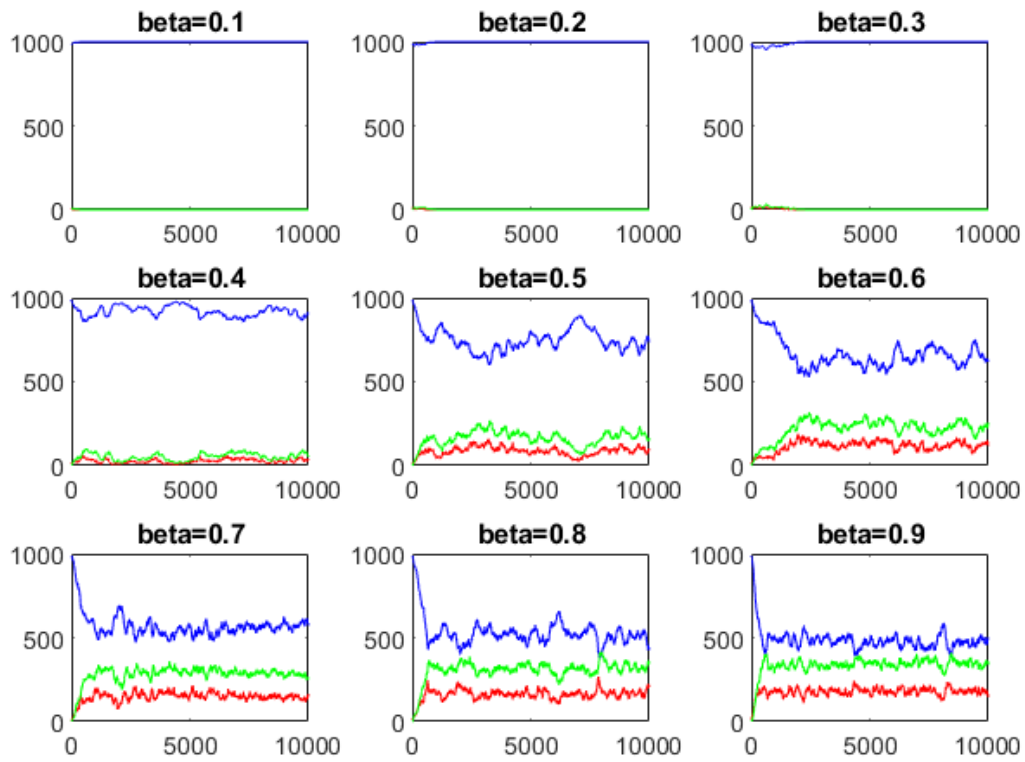
$\alpha=0.0015$ ,  $\gamma=0.015$



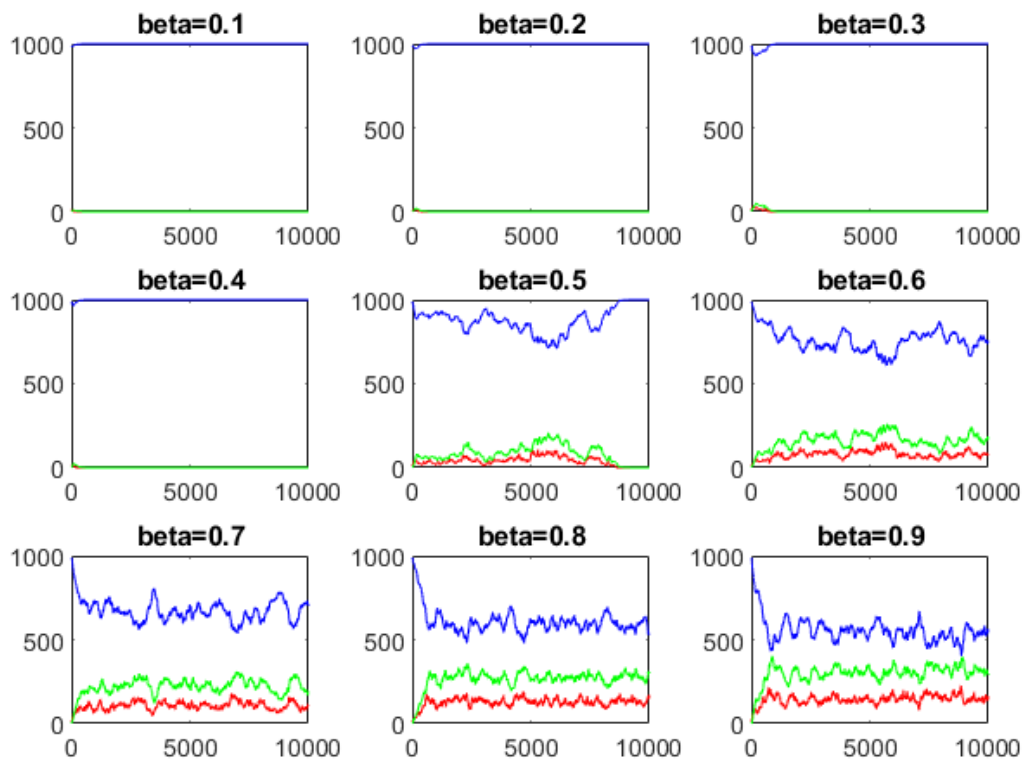
$\alpha=0.0100$ ,  $\gamma=0.010$



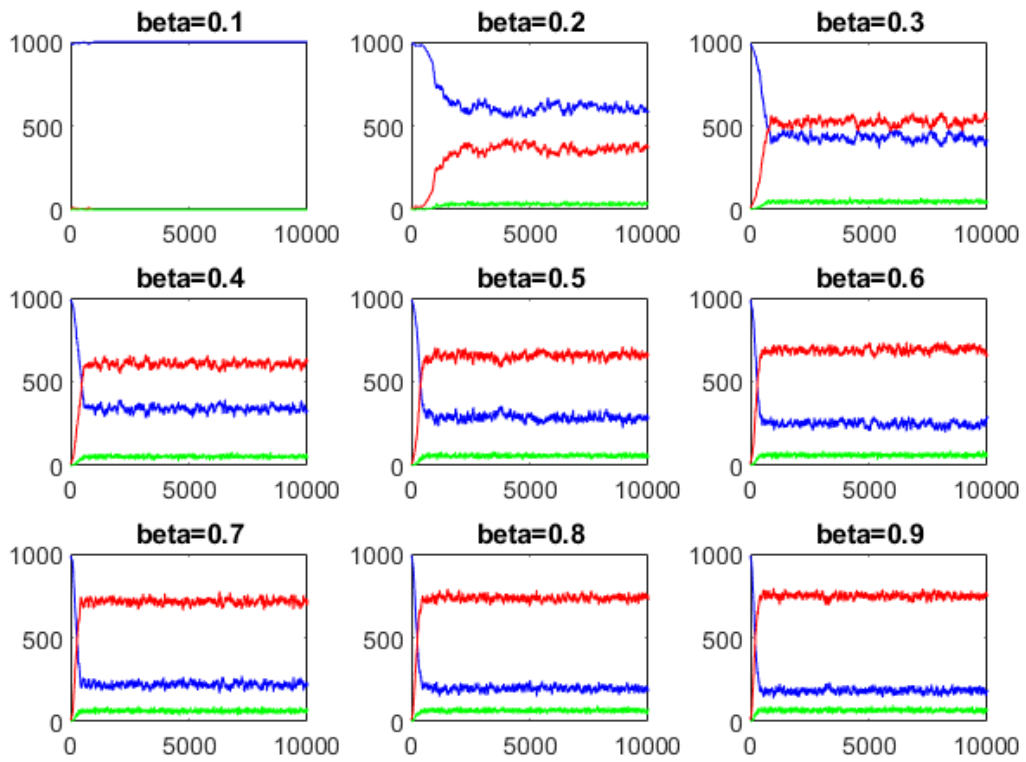
$\alpha=0.0100$ ,  $\gamma=0.020$



$\alpha=0.0100$ ,  $\gamma=0.022$



$\alpha=0.1000$ ,  $\gamma=0.010$



## Code

```
close all
clear all
clc
L = 100;
N = 1000;
infectedProb = 0.01;

% l = find(M==1);
% k = find(M==2);
% length(l)+length(k)
% %
% [x1, y1] = find(M==1);
% [x2, y2] = find(M==2);
% scatter(x1, y1, 10, "filled", MarkerFaceColor='r')
% hold on
% scatter(x2, y2, 10, "filled", MarkerFaceColor='g')

% Initialize lattice
initLattice = InitLatticeWithAgents(L,N,infectedProb);
initData = Data(initLattice);
nrOfInitInfected = initData(2);

beta = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9]; % infected probability
gamma = 0.03; % recover probability
d = 0.8;
for b = beta
    time = [];
    t = 0;
    nSuceptibleList = [];
    nInfectedList = [];
    nRecoveredList = [];
    lattice = initLattice;
    % while(FindInfected(lattice))
    for h = 1:1000
        data = Data(lattice);
        nSuceptibleList(end+1) = data(1);
        nInfectedList(end+1) = data(2);
        nRecoveredList(end+1) = data(3);
        % clf
        lattice = Move(lattice, d);
        lattice = infection(lattice, b);
        lattice = Recover(lattice, gamma);
        % PlotModel(lattice)
```

```
%    drawnow
t = t + 1;
time(end+1) = t;
%    if (nSuceptibleList(t) + nInfectedList(t) + nRecoveredList(t)) ~= N
%        disp('something wrong')
%        break
%    end
end
if b == 0.1
    b1Suceptible = nSuceptibleList;
    b1Infected = nInfectedList;
    b1Recovered = nRecoveredList;
elseif b== 0.2
    b2Suceptible = nSuceptibleList;
    b2Infected = nInfectedList;
    b2Recovered = nRecoveredList;
elseif b== 0.3
    b3Suceptible = nSuceptibleList;
    b3Infected = nInfectedList;
    b3Recovered = nRecoveredList;
elseif b== 0.4
    b4Suceptible = nSuceptibleList;
    b4Infected = nInfectedList;
    b4Recovered = nRecoveredList;
elseif b== 0.5
    b5Suceptible = nSuceptibleList;
    b5Infected = nInfectedList;
    b5Recovered = nRecoveredList;
elseif b== 0.6
    b6Suceptible = nSuceptibleList;
    b6Infected = nInfectedList;
    b6Recovered = nRecoveredList;
elseif b== 0.7
    b7Suceptible = nSuceptibleList;
    b7Infected = nInfectedList;
    b7Recovered = nRecoveredList;
elseif b== 0.8
    b8Suceptible = nSuceptibleList;
    b8Infected = nInfectedList;
    b8Recovered = nRecoveredList;
elseif b== 0.9
    b9Suceptible = nSuceptibleList;
    b9Infected = nInfectedList;
    b9Recovered = nRecoveredList;
end
end
```



```
subplot(3,3,1)
plot(time, b1Suceptible,'b')
hold on
plot(time, b1Recovered,'g')
plot(time, b1Infected,'r')
title('beta=0.1')
```

```
subplot(3,3,2)
plot(time, b2Suceptible,'b')
hold on
plot(time, b2Recovered,'g')
plot(time, b2Infected,'r')
title('beta=0.2')
```

```
subplot(3,3,3)
plot(time, b3Suceptible,'b')
hold on
plot(time, b3Recovered,'g')
plot(time, b3Infected,'r')
title('beta=0.3')
```

```
subplot(3,3,4)
plot(time, b4Suceptible,'b')
hold on
plot(time, b4Recovered,'g')
plot(time, b4Infected,'r')
title('beta=0.4')
```

```
subplot(3,3,5)
plot(time, b5Suceptible,'b')
hold on
plot(time, b5Recovered,'g')
plot(time, b5Infected,'r')
title('beta=0.5')
```

```
subplot(3,3,6)
plot(time, b6Suceptible,'b')
hold on
plot(time, b6Recovered,'g')
plot(time, b6Infected,'r')
title('beta=0.6')
```

```
subplot(3,3,7)
plot(time, b7Suceptible,'b')
hold on
plot(time, b7Recovered,'g')
plot(time, b7Infected,'r')
title('beta=0.7')
```

```
subplot(3,3,8)
plot(time, b8Suceptible,ld on
plot(time, b8Recovered,'g')
plot(time, b8Infected,'r')
title('beta=0.8')

subplot(3,3,9)
plot(time, b9Suceptible,'b')
hold on
plot(time, b9Recovered,'g')
plot(time, b9Infected,'r')
title('beta=0.9')

%% check to see that we initialize N agents in the grid
nrOfA = 0;
for j= 1:length(lattice)
    for i= 1:length(lattice)
        a = lattice{j, i};
        nrElements = length(a);
        nrOfA = nrOfA + nrElements;
    end
end

close all
clear all
clc
L = 100;
N = 1000;
infectedProb = 0.01;

% l = find(M==1);
% k = find(M==2);
% length(l)+length(k)
% %
% [x1, y1] = find(M==1);
% [x2, y2] = find(M==2);
% scatter(x1, y1, 10, "filled", MarkerFaceColor='r')
% hold on
% scatter(x2, y2, 10, "filled", MarkerFaceColor='g')

% Initialize lattice
initLattice = InitLatticeWithAgents(L,N,infectedProb);
initData = Data(initLattice);
nrOfInitInfected = initData(2);
```

```

betas = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9]; % infected probability
gammas = [0.01, 0.02]; % recover probability
d = 0.8;
trials = 4;

RList1 = zeros(trials, length(betas));
RList2 = zeros(trials, length(betas));
for trial = 1:trials
    for g = 1:length(gammas)
        gamma = gammas(g);
        for l = 1:length(betas)
            beta = betas(l);
            time = [];
            t = 0;
            lattice = initLattice;
            % while(FindInfected(lattice))
            for h = 1:1000
                data = Data(lattice);
                % clf
                lattice = Move(lattice, d);
                lattice = infection(lattice, beta);
                lattice = Recover(lattice, gamma);
                % PlotModel(lattice)
                % drawnow
                t = t + 1;
                time(end+1) = t;

                % if (nSuceptibleList(t) + nInfectedList(t) + nRecoveredList(t)) ~= N
                % disp('something wrong')
                % break
            % end
            end
            data = Data(lattice);
            if g == 1
                RList1(trial, l) = data(3);
            else
                RList2(trial, l) = data(3);
            end
        end
    end
end

averageR1 = sum(RList1,1)/trials;
averageR2 = sum(RList2,1)/trials;

scatter(betas, averageR1,'filled','b')

```

```

hold on
scatter(betas, averageR2,'filled','g')
legend('gamma=0.1', 'gamma=0.2')

%% store
save('R1average',"averageR1")
save('Raverage2','averageR2')
save('betas',"betas")
save('gammas',"gammas")
betaGamma1 = betas/gammas(1);
betaGamma2 = betas/gammas(2);

figure(1)
scatter(betaGamma1, averageR1)
hold on
scatter(betaGamma2, averageR2)
xlabel('beta/gamma')
ylabel('R average')
legend('gamma = 0.1', 'gamma = 0.2')
hold off

figure(2)
scatter(betas, averageR1,'filled','b')
hold on
scatter(betas, averageR2,'filled','g')
legend('gamma=0.1', 'gamma=0.2')
xlabel('beta')
ylabel('R average')
hold off

figure(3)
imagesc(betas, betaGamma1, averageR1)
colorbar

%% check to see that we initialize N agents in the grid
nrOfA = 0;
for j= 1:length(lattice)
    for i= 1:length(lattice)
        a = lattice{j, i};
        nrElements = length(a);
        nrOfA = nrOfA + nrElements;
    end
end

function PlotModel(lattice)

```

```

positions = zeros(length(lattice), length(lattice));

for i = 1:length(lattice)
    for j = 1:length(lattice)
        mostFrequentAgent = mode(lattice{i,j});
        if (mostFrequentAgent == 1)
            positions(i,j) = 1;
        elseif (mostFrequentAgent == 2)
            positions(i,j) = 2;
        elseif (mostFrequentAgent == 3)
            positions(i,j) = 3;
        else
            positions(i,j) = 0;
        end
    end
end

[x1,y1] = find(positions == 1);
[x2, y2] = find(positions == 2);
[x3, y3] = find(positions == 3);
scatter(x1, y1, 10, "filled", MarkerFaceColor='b') % suceptible
hold on
scatter(x2, y2, 10, "filled", MarkerFaceColor='r') % infected
scatter(x3, y3, 10, "filled", MarkerFaceColor='g') % recovered
end

function lattice = Move(lattice, d)
newLattice = lattice;
for i = 1:length(lattice)
    for j = 1:length(lattice)
        numberOfAgents = size(lattice{i,j}, 2);
        for n = 1:numberOfAgents
            r = rand();
            if (d > r)
                savedAgent = lattice{i,j}(1);
                lattice{i,j}(1) = [];
                newLattice{i, j}(1) = [];
                dir = randi(4);
                if (dir == 1)
                    if (j == 1)
                        newLattice{i, end} = [newLattice{i, end}, savedAgent];
                    else
                        newLattice{i, j-1} = [newLattice{i, j-1}, savedAgent];
                    end
                elseif (dir == 2)
                    if (j == length(lattice))
                        newLattice{i, 1} = [newLattice{i, 1}, savedAgent];
                    else
                        newLattice{i, j+1} = [newLattice{i, j+1}, savedAgent];
                    end
                end
            end
        end
    end
end

```

```

        end
    elseif (dir == 3)
        if (i == 1)
            newLattice{end, j} = [newLattice{end, j}, savedAgent];
        else
            newLattice{i-1, j} = [newLattice{i-1, j}, savedAgent];
        end
    elseif (dir == 4)
        if (i == length(lattice))
            newLattice{1, j} = [newLattice{1, j}, savedAgent];
        else
            newLattice{i+1, j} = [newLattice{i+1, j}, savedAgent];
        end
    end
    newLattice{x, y} = [newLattice{x, y}, savedAgent];
end
end
end
lattice = newLattice;
end

```

```
function lattice = InitLatticeWithAgents(L, N, infectedProb)
```

```

lattice = cell(L);
for n = 1:(N - N*infectedProb)
    coord = randi(L,1,2);
    x = coord(1);
    y = coord(2);
    lattice{x, y} = [lattice{x, y}, 1];
    if (n > (N - 2*N*infectedProb))
        lattice{x, y} = [lattice{x, y}, 2];
    end
end
end

```

```
end
```

```

function newLattice = infection(lattice, beta)
latticeLength = length(lattice);
for i = 1:latticeLength
    for j = 1:latticeLength
        indexInfected = find(lattice{i, j} == 2);
        if isempty(indexInfected)
            continue
        end
        indexSuceptible = find(lattice{i, j} == 1);
        if isempty(indexSuceptible)

```

```

        continue
    end
    for index=indexInfected
        randomNr = rand();
        if randomNr < beta
            for indexS = indexSuceptible
                lattice{i, j}(indexS) = 2; % suceptible gets infected
            end
        end
    end
end
end
end

newLattice = lattice;

end

function data = Data(lattice)

    nSuceptible = 0;
    nInfected = 0;
    nRecovered = 0;
    latticeLength = length(lattice);
    for i = 1:latticeLength
        for j = 1:latticeLength
            site = lattice{i, j};
            for agent = site
                if agent == 1
                    nSuceptible = nSuceptible + 1;
                elseif agent == 2
                    nInfected = nInfected + 1;
                elseif agent == 3
                    nRecovered = nRecovered + 1;
                end
            end
        end
    end
end

data = [nSuceptible, nInfected, nRecovered];

end

```

I Found out that my code was ineffective for the rest of the assignments so I rewrote to a much more effective code that is shown below

```

clear all
close all
clc
L = 100;

```

```

N = 1000;
nrOfInfected = 10;
d = 0.8;
beta = 0.6;
gamma = 0.01;
time = 1000;
betas = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9];
initializeAgents = InitializeAgentsWithPos(L, N, nrOfInfected);

```

```

dataArray = zeros(3*length(betas),time);
for i = 1:length(betas)
    beta = betas(i);
    agents = initializeAgents;
    for t = 1:time
        dataArray(1+3*(i-1):3+3*(i-1),t) = Data(agents);
        agents = Move(agents, d, L);
        agents = Recover(agents,gamma);
        agents = Infected(agents, beta);
    end
end
end

```

```

% timeVector = [1:time];
% length(timeVector)
% plot(timeVector, dataArray(1,:))
% hold on
% plot(timeVector, dataArray(2,:))
% plot(timeVector,dataArray(3,:))

```

```

for i = 1:length(betas)
    beta = betas(i);
    subplot(3,3,i)
    plot(dataArray(1+3*(i-1),:),'b')
    hold on
    plot(dataArray(2+3*(i-1),:),'r')
    plot(dataArray(3+3*(i-1),:),'g')
    title(sprintf('beta=%0.1f',beta))
end

```

```

clear all
close all
clc
L = 100;
N = 1000;
nrOfInfected = 10;
d = 0.8;
beta = 0.6;
gamma = 0.01;

```



```

time = 1000;
betas = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9];
gammas = [0.01, 0.02];
initializeAgents = InitializeAgentsWithPos(L, N, nrOfInfected);

trials = 4;
RArray1 = zeros(trials, length(betas));
RArray2 = zeros(trials, length(betas));
for j = 1:length(gammas)
    gamma = gammas(j);
    for trial = 1:trials
        for i = 1:length(betas)
            beta = betas(i);
            agents = initializeAgents;
            for t = 1:time
                agents = Move(agents, d, L);
                agents = Recover(agents, gamma);
                agents = Infected(agents, beta);
                data = Data(agents);
            end
            if j == 1
                RArray1(trial, i) = data(3);
            elseif j == 2
                RArray2(trial, i) = data(3);
            end
        end
    end
end

R1Averages = sum(RArray1, 1) / trials;
R2Averages = sum(RArray2, 1) / trials;
scatter(betas, R1Averages, 'filled', 'b')
hold on
scatter(betas, R2Averages, 'filled', 'g')
legend(sprintf('gamma=%0.3f', gammas(1)), sprintf('gamma=%0.3f', gammas(2)), 'Location', 'southeast')

clear all
close all
clc
L = 100;
N = 1000;
nrOfInfected = 10;
d = 0.8;
beta = 0.6;
gamma = 0.01;
time = 1000;
betas = [0.01:0.04:1];
gammas = [0.0075:0.000205:0.0125];
initializeAgents = InitializeAgentsWithPos(L, N, nrOfInfected);

```

```

trials = 4;
% RArray1 = zeros(trials, length(betas));
% RArray2 = zeros(trials, length(betas));
RArray = zeros(length(gammas), length(betas));
for j = 1:length(gammas)
    gamma = gammas(j);
    R = zeros(trials, length(betas));
    for trial = 1:trials
        for i = 1:length(betas)
            beta = betas(i);
            agents = initializeAgents;
            for t = 1:time
                agents = Move(agents, d, L);
                agents = Recover(agents, gamma);
                agents = Infected(agents, beta);
                data = Data(agents);
            end
            R(trial, i) = data(3);
        end
    end
    RAverages = sum(R,1) / trials;
    RArray(j, :) = RAverages;
end

% R1Averages = sum(RArray1,1) / trials;
% R2Averages = sum(RArray2,1) / trials;
% scatter(betas, R1Averages, 'filled', 'b')
% hold on
% scatter(betas, R2Averages, 'filled', 'g')
% legend(sprintf('gamma=%0.3f', gammas(1)), sprintf('gamma=%0.3f', gammas(2)), 'Location', 'southeast')

%% Saving RArray and plot phase diagram

save('RAverages', 'RArray')
gammaBetas = betas ./ gammas;
[x, y] = meshgrid(betas, gammaBetas);
surf(x, y, RArray)
xlabel('beta')
ylabel('beta/gamma')
title('Rinf')
view(2)
colorbar

clear all
close all
clc
L = 100;

```

```

N = 1000;
nrOfInfected = 10;
d = 0.8;
beta = 0.6;
gamma = 0.01;
mu = 0.00007;
time = 1000;
betas = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9];
initializeAgents = InitializeAgentsWithPos(L, N, nrOfInfected);

% dataArray = zeros(4*length(betas),time);
% for i = 1:length(betas)
%     beta = betas(i);
%     agents = initializeAgents;
%     for t = 1:time
%         dataArray(1+4*(i-1):4+4*(i-1),t) = DataMortality(agents);
%         agents = Move(agents, d, L);
%         agents = Recover(agents,gamma);
%         agents = Infected(agents, beta);
%         agents = Death(agents, mu);
%     end
% end
%
%
%
%
% for i = 1:length(betas)
%     beta = betas(i);
%     subplot(3,3,i)
%     plot(dataArray(1+4*(i-1),:),'b')
%     hold on
%     plot(dataArray(2+4*(i-1),:),'r')
%     plot(dataArray(3+4*(i-1),:),'g')
%     plot(dataArray(4+4*(i-1),:),'k')
%     title(sprintf('beta=%0.1f, mu=%0.4f',beta,mu))
% end

%% 11.3b)
betas = [0.3, 0.6];
gammas = [0.01, 0.015];
trials = 3;
mus = [0:0.001:0.02];
DArray1 = zeros(trials, length(mus));
DArray2 = zeros(trials, length(mus));
DArray3 = zeros(trials, length(mus));
DArray4 = zeros(trials, length(mus));
DArray5 = zeros(trials, length(mus));

```

```

DArray6 = zeros(trials, length(mus));
DArray7 = zeros(trials, length(mus));
DArray8 = zeros(trials, length(mus));
for m = 1:length(mus)
    mu = mus(m);
    for j = 1:length(gammas)
        gamma = gammas(j);
        for i = 1:length(betas)
            beta = betas(i);
            for trial = 1:trials
                agents = initializeAgents;
                for t = 1:time
                    agents = Move(agents, d, L);
                    agents = Recover(agents, gamma);
                    agents = Infected(agents, beta);
                    agents = Death(agents, mu);
                    data = DataMortality(agents);
                end
                if i==1 && j==1
                    DArray1(trial,m) = data(4); % beta1 & gamma1
                elseif i==1 && j==2
                    DArray2(trial,m) = data(4); %beta1 & gamma2
                elseif i==2 && j==1
                    DArray3(trial,m) = data(4); %beta2 & gamma1
                elseif i==2 && j==2
                    DArray4(trial,m) = data(4); %beta2 & gamma2
                end
            end
        end
    end
end
end
end

D1Averages = sum(DArray1,1) / trials;
D2Averages = sum(DArray2,1)/ trials;
D3Averages = sum(DArray3,1)/ trials;
D4Averages = sum(DArray4,1)/ trials;
scatter(mus, D1Averages, 'b')
hold on
scatter(mus, D2Averages, 'r')
scatter(mus, D3Averages, 'k')
scatter(mus, D4Averages, 'g')
prints1 = sprintf('gamma=%0.3f, beta=%0.3f', gammas(1), betas(1));
prints2 = sprintf('gamma=%0.3f, beta=%0.3f', gammas(2), betas(1));
prints3 = sprintf('gamma=%0.3f, beta=%0.3f', gammas(1), betas(2));
prints4 = sprintf('gamma=%0.3f, beta=%0.3f', gammas(2), betas(2));
legend(prints1, prints2, prints3, prints4, 'Location', 'best');
xlabel('mu')

```

```
ylabel('Dinf')
```

```
%
%           if j == 1 && i==1
%               DArray1(trial,i) = data(4);
%           elseif j==1 && i==2
%               DArray3(trial,i) = data(4);
%           elseif j==1 && i==3
%               DArray4(trial,i) = data(4)
%
%           elseif j == 2
%               DArray2(trial,i) = data(4);
%           end
%       end
%   end
% end
%
% R1Averages = sum(DArray1,1) / trials;
% R2Averages = sum(DArray2,1)/ trials;
% scatter(betas, R1Averages,'filled','b')
% hold on
% scatter(betas, R2Averages,'filled','g')
% legend(sprintf('gamma=%0.3f',gammas(1)), sprintf('gamma=%0.3f', gammas(2)),'Location','southeas

function agents = Recover(agents, gamma)

indexOfInfected = find(agents(:, 3)==2);
randomNumbers = rand(length(indexOfInfected),1);

indexGettingRecovered = indexOfInfected(find(randomNumbers<gamma));
if ~isempty(indexGettingRecovered)
    agents(indexGettingRecovered,3) = 3;
end

end

function agents = Move(agents, d, L)
N = length(agents);    %number of agents
randomNumbers = rand(N,1);
indexForMove = find(randomNumbers < d);

nMovingAgents = length(indexForMove);    %nr of agents that will move
```

```

randomMoveIndex = randi(4, nMovingAgents, 1);
Movements = [0, 1; 0, -1; 1, 0; -1, 0];

move = Movements(randomMoveIndex, :);
agents(indexForMove, 1:2) = agents(indexForMove, 1:2) + move;

% coordinates that gets outside grid gets to other side (PBC)
tooSmallCoordinates = find(agents(:, 1:2) < 1);
tooBigCoordinates = find(agents(:, 1:2) > L);
if ~isempty(tooBigCoordinates)
    agents(tooBigCoordinates) = agents(tooBigCoordinates) - L;
end
if ~isempty(tooSmallCoordinates)
    agents(tooSmallCoordinates) = agents(tooSmallCoordinates) + L;
end
end
end

function agents = InitializeAgentsWithPos(L, N, nrOfInfected)
agents = zeros(N,3);
for n = 1:N
    coord = randi(L,1,2);
    x = coord(1);
    y = coord(2);
    agents(n,1) = x;
    agents(n,2) = y;
    agents(n,3) = 1;
    if (n > (N - nrOfInfected))
        agents(n,3) = 2;
    end
end
end

function agents = Infected(agents, beta)

% Find index of suceptible and infected
indexOfInfected = find(agents(:, 3) == 2);
coordsOfInfected = agents(indexOfInfected, 1:2);

% use index to find pos of suceptible and infected
indexOfSuceptible = find(agents(:,3)==1);
coordsOfSuceptible = agents(indexOfSuceptible,1:2);

% find sites where there is suceptible and infected
InfectedWithSuceptibleIndex = find(ismember(coordsOfSuceptible, coordsOfInfected, 'rows'));
if isempty(InfectedWithSuceptibleIndex)
    return
else

```

```

% Suceptible gets infected with probability beta
randomNumbers = rand(length(InfectedWithSuceptibleIndex),1);
suceptibleGetInfectedIndex = InfectedWithSuceptibleIndex(find(randomNumbers<beta));
if ~isempty(suceptibleGetInfectedIndex)
    coordsOfGettingInfected = coordsOfSuceptible(suceptibleGetInfectedIndex,:);
    gettingInfectedIndex = find(ismember(agents(:,1:2),coordsOfGettingInfected,"rows"));
    agents(gettingInfectedIndex,3) = 2;

end
end
end

function agents = Immunity(agents, alpha)

indexRecovered = find(agents(:,3)==3);
randomNumbers = rand(length(indexRecovered),1);

indexGettingSuceptible = indexRecovered(find(randomNumbers<alpha));
if ~isempty(indexGettingSuceptible)
    agents(indexGettingSuceptible,3) = 1;
end

end

function agents = Death(agents, mu)

indexOfInfected = find(agents(:, 3) == 2);
randomNr = rand(length(indexOfInfected),1);
indexOfGettingDead = indexOfInfected(find(randomNr<mu));
if ~isempty(indexOfGettingDead)
    agents(indexOfGettingDead,3) = 0;
end

end

function data = DataMortality(agents)

% Find index of the Three different types of agents
indexOfSuceptible = find(agents(:,3)==1);
indexOfInfected = find(agents(:,3)==2);
indexOfRecovered = find(agents(:,3)==3);
indexOfDead = find(agents(:,3)==0);

nrOfSuceptible = length(indexOfSuceptible);
nrOfInfected = length(indexOfInfected);
nrOfRecovered = length(indexOfRecovered);
nrOfDead = length(indexOfDead);
data = [nrOfSuceptible; nrOfInfected; nrOfRecovered; nrOfDead];

```

```
function data = Data(agents)

% Find index of the Three different types of agents
indexOfSuceptible = find(agents(:,3)==1);
indexOfInfected = find(agents(:,3)==2);
indexOfRecovered = find(agents(:,3)==3);

nrOfSuceptible = length(indexOfSuceptible);
nrOfInfected = length(indexOfInfected);
nrOfRecovered = length(indexOfRecovered);

data = [nrOfSuceptible; nrOfInfected; nrOfRecovered];
```