

Model

```
function dydt = model(t, y , params)

s = y(1); %sensitive bacteria
r = y(2); %resistant bacteria
b = y(3); %immune cells
a1 = y(4); % INH
a2 = y(5); % ZPA

% PARAMETERS
beta_S = 1.1; % Growth rate of sensitive bacteria
beta_R = 0.9; % Growth rate of resistant bacteria (with fitness cost)
eta = 0.6; % Immune killing rate
k = 0.4; % Immune cell proliferation rate

% For INH (antibiotic 1)
alpha1 = params.alpha1; % Mutation rate due to INH
d1 = params.d1; % Death rate due to INH

% For ZPA (antibiotic 2)
alpha2 = params.alpha2; % Mutation rate due to ZPA
d2 = params.d2; % Death rate due to ZPA

mu1 = 0.08; % Clearance rate for INH
mu2 = 0.04; % Clearance rate for ZPA

% DE
total_bac = s + r; % Total bacteria
```

% Sensitive bacteria

$$\% \frac{ds}{dt} = \beta_S * s * (1 - s - r) - \eta * s * b - s * \sum(\alpha_i + d_i) * a_i$$

$$\text{growth_s} = \text{beta_S} * s * (1 - \text{total_bac});$$

$$\text{immune_kill_s} = \text{eta} * s * b;$$

$$\text{antibiotic_effect_s} = s * ((\text{alpha1} + d1) * a1 + (\text{alpha2} + d2) * a2);$$

$$ds = \text{growth_s} - \text{immune_kill_s} - \text{antibiotic_effect_s};$$

% Resistant bacteria

$$\% \frac{dr}{dt} = \beta_R * r * (1 - s - r) - \eta * r * b + s * \sum(\alpha_i) * a_i$$

$$\text{growth_r} = \text{beta_R} * r * (1 - \text{total_bac});$$

$$\text{immune_kill_r} = \text{eta} * r * b;$$

$$\text{mutation_gain} = s * (\text{alpha1} * a1 + \text{alpha2} * a2);$$

$$dr = \text{growth_r} - \text{immune_kill_r} + \text{mutation_gain};$$

% Immune cells

$$\% \frac{db}{dt} = k * b * (1 - b/(s + r))$$

$$db = k * b * (1 - b / \text{total_bac});$$

% Antibiotic concentrations

$$\% \frac{da_i}{dt} = \mu_i * (1 - a_i)$$

$$da1 = \mu1 * (1 - a1);$$

$$da2 = \mu2 * (1 - a2);$$

```

% OUTPUT
dydt = [ds; dr; db; da1; da2];
end

```

All variables (A<B)

```

tspan = [0 90];

s = 0.8; %sensitive bacteria
r = 0.05; %resistant bacteria
b = 0.05; %immune cells
a1 = 0.05; % INH
a2 = 0.02; % ZPA

```

```

% For INH (antibiotic 1)
params.alpha1 = 0.02; % Mutation rate due to INH
params.d1 = 0.15; % Death rate due to INH

```

```

% For ZPA (antibiotic 2)
params.alpha2 = 0.06; % Mutation rate due to ZPA
params.d2 = 0.35; % Death rate due to ZPA

```

```

y0 = [s r b a1 a2];
[t,y] = ode45(@(t,y) model(t,y,params), tspan, y0);

s = y(:,1);
r = y(:,2);
b = y(:,3);

```

```

a1 = y(:,4);
a2 = y(:,5);

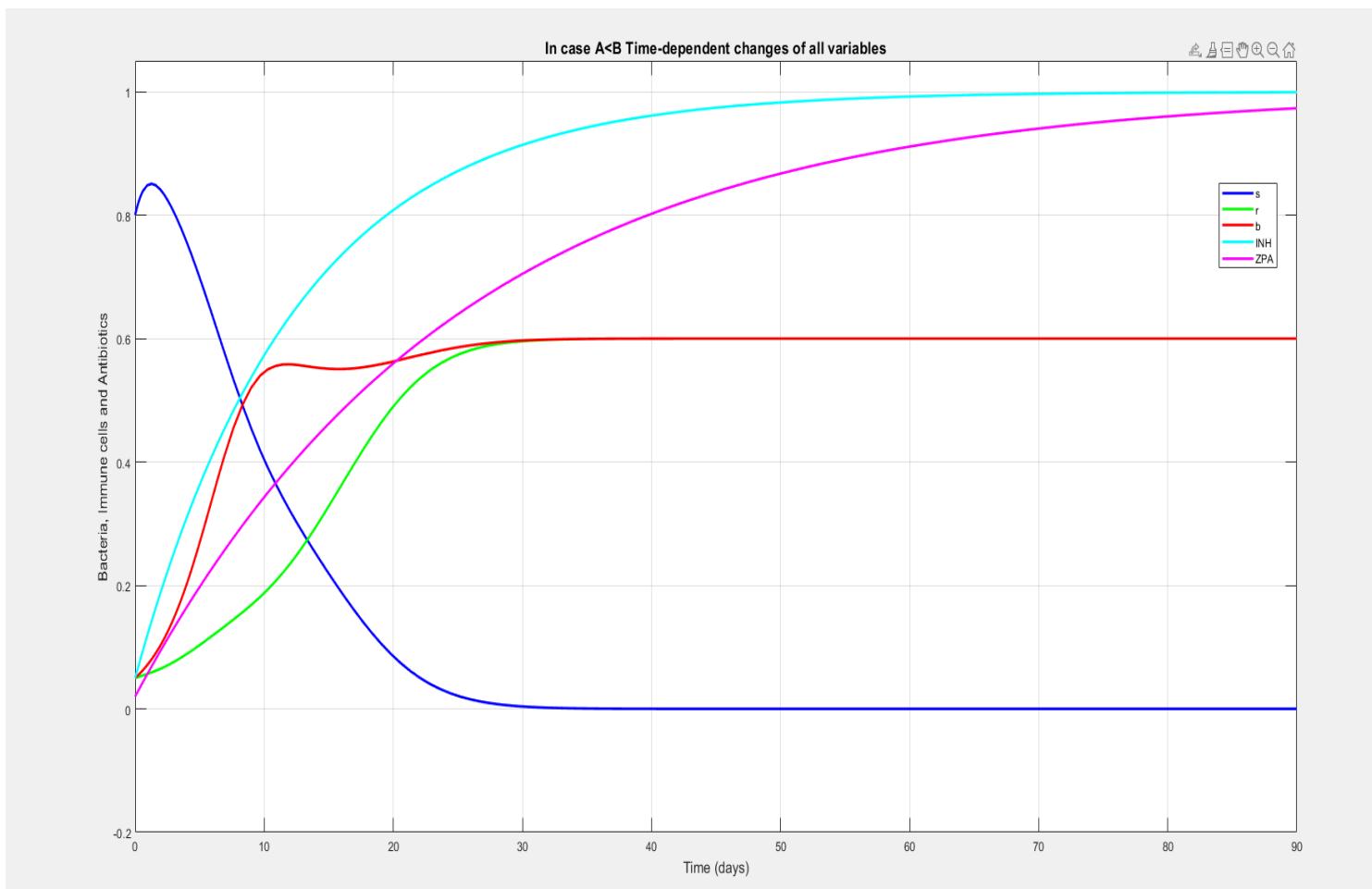
% Plot
figure;
hold on; grid on; box on;

plot(t,s,'b','LineWidth',2)
plot(t,r,'g','LineWidth',2)
plot(t,b,'r','LineWidth',2)
plot(t,a1,'c','LineWidth',2)
plot(t,a2,'m','LineWidth',2)

xlabel('Time (days)','FontSize',12)
ylabel('Bacteria, Immune cells and Antibiotics','FontSize',12)
title('In case A<B Time-dependent changes of all variables','FontSize',13)

legend('s','r','b','INH','ZPA','Location','best')
ylim([-0.2 1.05])
xlim([0 90])

```



Bacteria Vs Immune

tspan = [0 90];

s = 0.8; %sensitive bacteria

r = 0.05; %resistant bacteria

b = 0.05; %immune cells

a1 = 0.05; % INH

a2 = 0.02; % ZPA

% For INH (antibiotic 1)

```

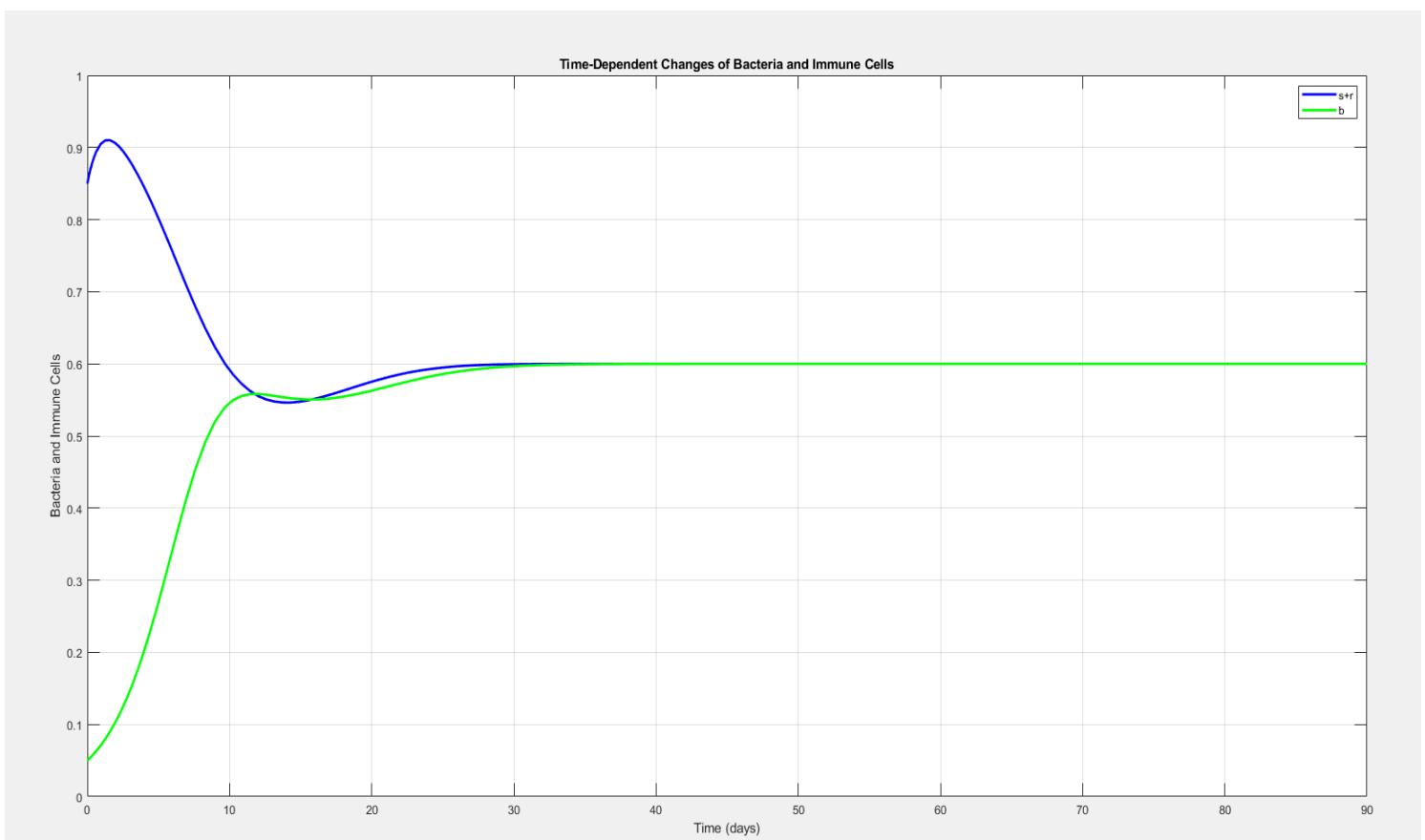
params.alpha1 = 0.02;    % Mutation rate due to INH
params.d1 = 0.15;        % Death rate due to INH

% For ZPA (antibiotic 2)
params.alpha2 = 0.06;    % Mutation rate due to ZPA
params.d2 = 0.35;        % Death rate due to ZPA

y0 = [s r b a1 a2];
[t,y] = ode45(@(t,y) model(t,y,params), tspan, y0);

figure
plot(t, y(:,1)+y(:,2),'b','LineWidth',2)    % s+r
hold on
plot(t, y(:,3),'g','LineWidth',2)              % b
xlabel('Time (days)')
ylabel('Bacteria and Immune Cells')
title('Time-Dependent Changes of Bacteria and Immune Cells')
legend('s+r','b')
grid on

```



All Variables (B<A)

tspan = [0 90];

s = 0.8; %sensitive bacteria

r = 0.07; %resistant bacteria

b = 0.05; %immune cells

a1 = 0.05; % INH

```
a2 = 0.02; % ZPA
```

```
% For INH (antibiotic 1)
```

```
params.alpha1 = 0.06; % Mutation rate due to INH
```

```
params.d1 = 0.35; % Death rate due to INH
```

```
% For ZPA (antibiotic 2)
```

```
params.alpha2 = 0.02; % Mutation rate due to ZPA
```

```
params.d2 = 0.15; % Death rate due to ZPA
```

```
y0 = [s r b a1 a2];
```

```
[t,y] = ode45(@(t,y) model(t,y,params), tspan, y0);
```

```
s = y(:,1);
```

```
r = y(:,2);
```

```
b = y(:,3);
```

```
a1 = y(:,4);
```

```
a2 = y(:,5);
```

```
% Plot
```

```
figure;
```

```
hold on; grid on; box on;
```

```
plot(t,s,'b','LineWidth',2)
```

```
plot(t,r,'g','LineWidth',2)
```

```
plot(t,b,'r','LineWidth',2)
```

```
plot(t,a1,'c','LineWidth',2)
```

```
plot(t,a2,'m','LineWidth',2)
```

```

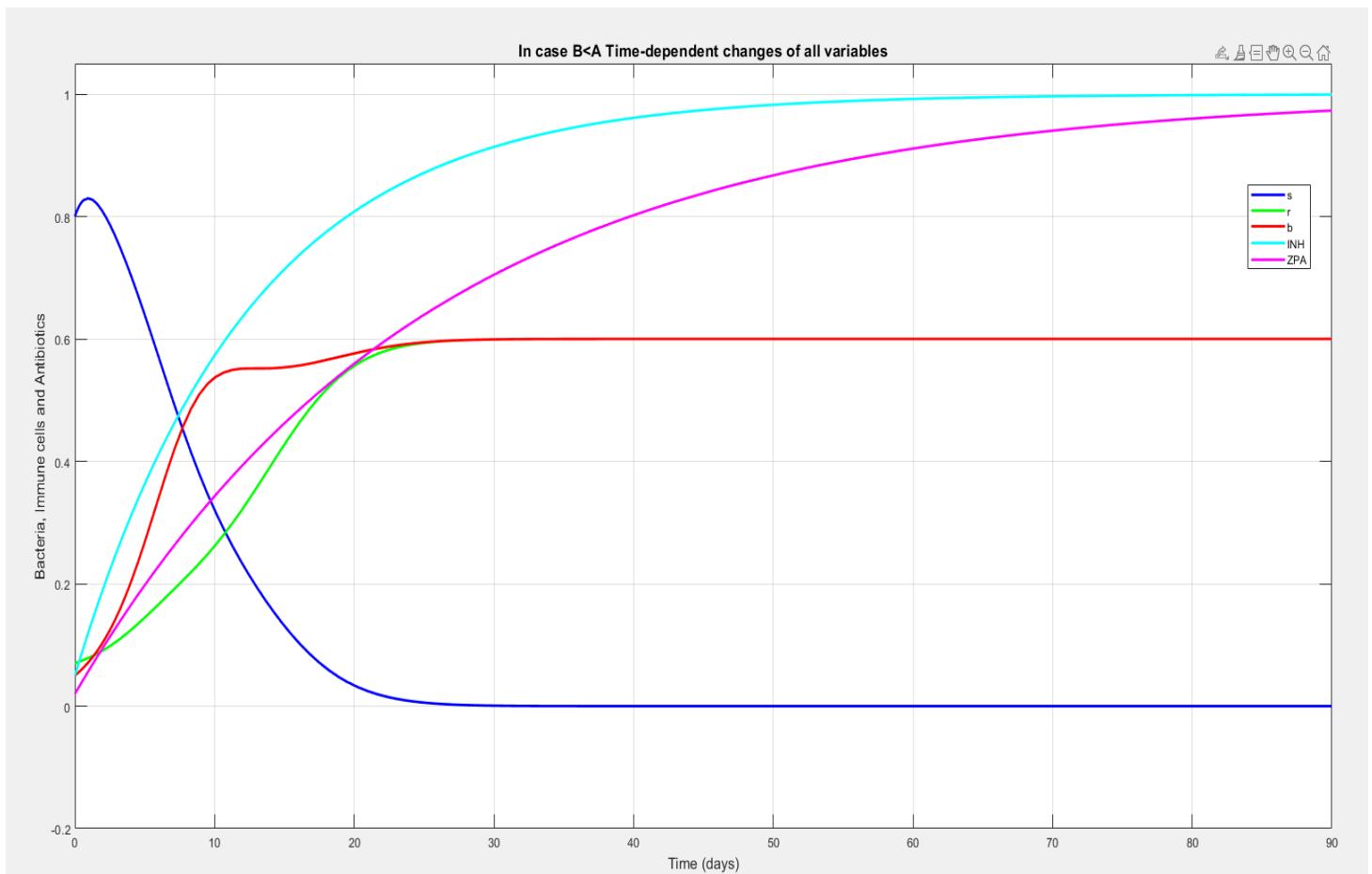
xlabel('Time (days)', 'FontSize', 12)
ylabel('Bacteria, Immune cells and Antibiotics', 'FontSize', 12)
title('In case B<A Time-dependent changes of all variables', 'FontSize', 13)

```

```

legend('s','r','b','INH','ZPA','Location','best')
ylim([-0.2 1.05])
xlim([0 90])

```



Resistance VS Sensitive Bacteria

```

params.alpha1 = 0.06; % Mutation rate due to INH
params.d1 = 0.35; % Death rate due to INH
params.alpha2 = 0.02; % Mutation rate due to ZPA
params.d2 = 0.15; % Death rate due to ZPA

s = 0.8; %sensitive bacteria
r = 0.1; %resistant bacteria
b = 0.5; %immune cells
a1 = 1; % INH
a2 = 1; % ZPA

y0 = [s, r, b, a1, a2];

tspan = [0 90];

[t, y] = ode45(@(t,y) model(t, y, params), tspan, y0);

figure;
plot(y(:,1), y(:,2), 'b', 'LineWidth', 2); % y(:,1)=s, y(:,2)=r
xlabel('Sensitive Bacteria');
ylabel('Resistant Bacteria');
title('Time-Dependent Changes of Bacteria ');
grid on;

```

```
hold on;  
plot(y(1,1), y(1,2), 'ks', 'MarkerFaceColor', 'k'); % initial point  
plot(y(end,1), y(end,2), 'ks', 'MarkerFaceColor', 'r'); % final point
```

