gut

August 4, 2025

## 1 Modeling Bacteria, Antibiotics and Immune System

This notebook is to accompany the Report file Computation and Code section. Figures and Simulations are generated here.

## 1.1 Parameter Selection

Our Invarient Set  $\mathcal{A}$  implied some restriction on parameters which alligned with our biological and mathematical intuition.

```
[1]: import numpy as np import matplotlib.pyplot as plt
```

## 1.2 Symbolic Calculation

Here is a Calculation of Jacoubian of the system without numerical computation. Using Python Symbolic library.

```
[2]: import sympy as sp
a, s, r, p = sp.symbols('a s r p')
mu, eta_r, eta_s, alpha, beta, gamma = sp.symbols('mu eta_r eta_s alpha beta_u egamma')

vars_params = [a, s, r, p, alpha, beta, gamma, eta_s, eta_r, mu]

def f(x):
    return x - x**2 + 3/4

n = s + r

f1 = mu * (1 - a)
f2 = eta_s*(1 - n)*s - alpha * a * s - (beta * s * r )/n - gamma * s * p
f3 = eta_r*(1 - n)*r + (beta * s * r )/n - gamma * r * p
f4 = p * (f(n) - p)

dyn = sp.Matrix([f1,f2,f3,f4])

J = dyn.jacobian([a,s,r,p])
```

```
[3]: ## for E1
     li = [1, 0, 0, f(0), 1, 0.1, 1, 1, 0.3, 3]
     dic = dict(zip(vars_params, li))
     res = J.subs(dic).evalf()
     res
[3]: _{\Gamma-3.0}
                    0
                           0
       0
           -0.75
                 -0.1
                  -0.35
       0
             0
                           0
       0
            0.75
                   0.75
                         -0.75
[4]: from scipy.integrate import odeint
     def evaluate_dyn(y , t, alpha, beta, gamma, eta_s, eta_r, mu) -> tuple:
         a, s, r, p = y
         state\_vars = [a, s, r, p, alpha, beta, gamma, eta\_s, eta\_r, mu]
         dic = dict(zip(vars_params, state_vars))
         res = dyn.subs(dic).evalf()
         res = sp.matrix2numpy(res, dtype=np.float64)
         return res.flatten()
     parset = (1, 0.1, 1, 1, 0.3, 3)
     y0 = np.repeat(0.5, 4)
     t = np.linspace(0, 10, 1000)
     sol = odeint(evaluate_dyn, y0, t, args=parset)
[5]: fig, ax = plt.subplots(figsize=(10, 6))
     variable_names = ['Antibiotic (a)', 'NARB (s)', 'ARB (r)', 'Immune System (p)']
     colors = ['#1f77b4', '#ff7f0e', '#2ca02c', '#d62728'] # Professional color_
      ⇔scheme
     for i in range(4):
         ax.plot(t, sol[:, i],
                 color=colors[i],
                 linewidth=2,
                 label=variable_names[i])
     ax.set_title('Approching E1', fontsize=14)
     ax.set xlabel('Time', fontsize=12)
     ax.set_ylabel('System Variables', fontsize=12)
     ax.grid(True, linestyle='--', alpha=0.7)
     ax.legend(loc='best')
```

plt.tight\_layout()
plt.show()

