

Bacteria Growth Model

Minerva University

CS166 - Modeling and Analysis of Complex Systems

March 2, 2025

1 Model Description

This paper focuses on the simulation and analysis of the bacteria growth model in closed 2 dimensional environment. In the environment, we track the dynamics between the amount of food and bacteria, which are influenced by factors including bacterial reproduction rate, and food growth rate, bacterial and food diffusion rate, bacterial consumption rate, food reseeding probability, and food reseed rate. We will run simulations on the model using cellular automata (CA), analyze the empirical results, and come up with mathematical relationship between variables.

1.1 Initializing the bacteria growth model

The simulation is run on a 100 x 100 grid, each grid consists of a tuple, recording the food count and the bacteria count. And the CA employs a Von Neumann neighborhood.

The initialization consists of three parameters — bacteria probability (bacteria_prob), food probability (food_prob), and the amount of bacteria for initialization (bacteria_amount). The bacteria population is initiated by choosing which grids of the population is to be populated using the bacteria probability. Afterwards, we put a fixed bacteria amount (bacteria_amount), which is set to 20, in those grids.

The food distribution is initiated by using food probability (food_prob) to decide the grids that contains food. Then, we randomly generate the amount of food between 0 and the maximum food amount (100 units) in them. There is no caps on the bacteria population. It is nonetheless constraint by the maximum food amount.

There are two reasons to make this arrangement. First, I wanted to observe the bacteria's growth, so I let them start from a small number. Second, I wanted to create different growth dynamics in the beginning of the simulation, thus I let food amount vary.

Name	Variable	Description
Food probability	food_prob	The probability that a grid contains certain amount of food when initialized.
Bacteria probability	bacteria_prob	The probability that a grid contains certain bacteria when initialized.
Bacteria amount	bacteria_amount	The fixed amount of bacteria in a grid if present.
Grid size	grid_size	The side length of the square grid.

Table 1: Parameter descriptions for initializing the bacteria growth CA simulation.

The simulation code for the bacterial growth model initialization shows the methods used to initialize the CA. The parameters are all predetermined.

```
1  def initialize_ca(self, food_prob=0.3, bacteria_prob=0.1, bacteria_amount=15):
2      """
3      Initialize the grid with random food and bacteria.
4      """
5      # Initialize food randomly by generating a temporary grid for deciding where to put
6      # food
7      # Initiate with randomly chosen food amount between 0 and 100
8      food_mask = np.random.random((self.grid_size, self.grid_size)) < food_prob
9      self.food = np.where(food_mask, (self.food_max * np.random.random((self.grid_size,
10     self.grid_size))), 0)
11
12     # Initialize bacteria randomly by generating a temporary grid for deciding where to
13     put bacteria
```

```

11     # Initiate with 15 bacteria for the grids with bacteria
12     bacteria_mask = np.random.random((self.grid_size, self.grid_size)) < bacteria_prob
13     self.bacteria = np.where(bacteria_mask, bacteria_amount, 0)
14
15     # Record initial state
16     self.food_history = [np.mean(self.food)]
17     self.bacteria_history = [np.mean(self.bacteria)]
18
19     if self.testing_mode == True:
20         np.set_printoptions(threshold=20) # Ensures all elements are printed
21         print("Initialized CA")
22         print("----- Food distribution -----")
23         print(self.food)
24         print("----- Bacteria population -----")
25         print(self.bacteria)

```

1.2 Updating the bacteria growth model

In every step of updates, the bacteria will first consume the amount of food available in its grid based on its consumption rate (`bacteria_consumption_rate`). Then, the bacteria reproduces based on its current amount by its reproduction rate (`bacteria_growth_rate`). The remaining food then grow by its growth rate (`food_growth_rate`) with a probability to reseed (`reseed_prob`). The food growth follows logistic growth formula, meaning that for the amount of food in a grid a time t (f_t). The food growth follows the formula:

$$f_{t+1} = f_t(1 + g_f(1 + \frac{f_t}{k_f})),$$

where g_f is the food growth rate and k_f is the maximum food amount (100 units).

Then, the food reseeds with a predetermined probability of 0.01. Finally, the food and bacteria in a grid both diffuse to their's Von-Neumann neighbors by their respective diffusion rates (`food_diffusion_rate`, `bacteria_diffusion_rate`), and each of the neighbors gets $\frac{1}{4}$ of the amount that diffused.

Name	Variable	Description
Bacteria reproduction rate	<code>bacteria_growth_rate</code>	A fixed reproduction rate of bacteria population in each step.
Food growth rate	<code>food_growth_rate</code>	A logistic growth rate of food in each step.
Bacteria consumption rate	<code>bacteria_consumption_rate</code>	The amount of food the bacteria consume in proportion to the bacterial population each step.
Food diffusion rate	<code>food_diffusion_rate</code>	The proportion of food in a grid that equally diffuses to it's Von-Neumann neighborhood neighbors each step.
Bacteria diffusion rate	<code>bacteria_diffusion_rate</code>	The proportion of bacteria in a grid that equally diffuses to it's Von-Neumann neighborhood neighbors each step.
Food reseeding probability	<code>reseed_prob</code>	A predetermined probability that the food amount in a grid + 1

Table 2: Parameter descriptions for updating the bacteria growth CA.

The code for simulation also clearly demonstrated the sequence of interaction between food and bacteria, and their own population dynamics.

```

1 def update(self):
2     """
3     Perform one step of the simulation.
4     """
5
6     # Bacteria consume food, and die out if they don't have enough food
7     self.consumption_and_starvation()

```

```

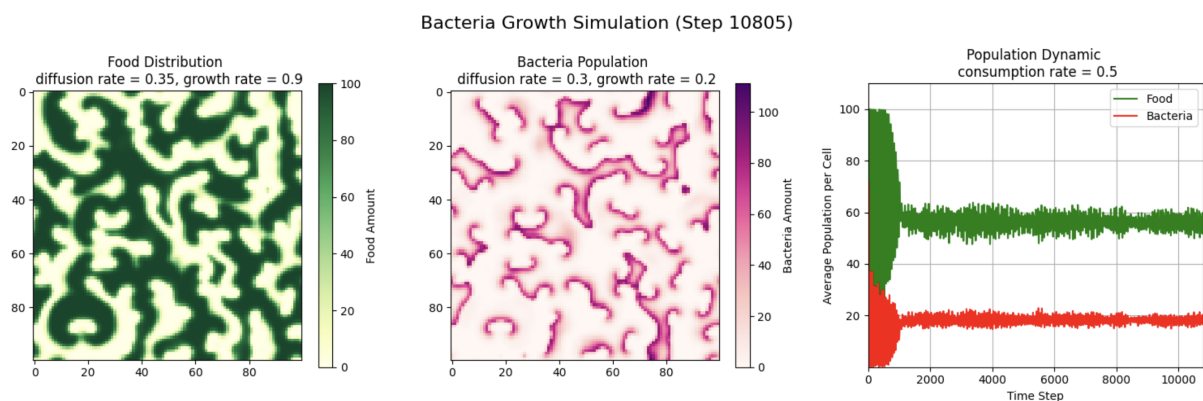
8
9  # The remaining bacteria reproduce
10 self.bacteria_reproduction()
11
12 # The remaining food grows
13 self.food_growth()
14
15 # Food reseeds with a probability
16 self.food_reseeding()
17
18 # Food diffuses
19 self.food_diffusion()
20
21 # Bacteria diffuses
22 self.bacteria_diffusion()
23
24 # Record history
25 self.food_history.append(np.mean(self.food))
26 self.bacteria_history.append(np.mean(self.bacteria))
27
28 if self.testing_mode == True:
29     print("One step done!")

```

2 Empirical analysis

2.1 Simulation

The CA simulation of the bacteria growth model were run for 12,000 steps each. There are generally two types of result — the ones that converges, and the ones that do not.



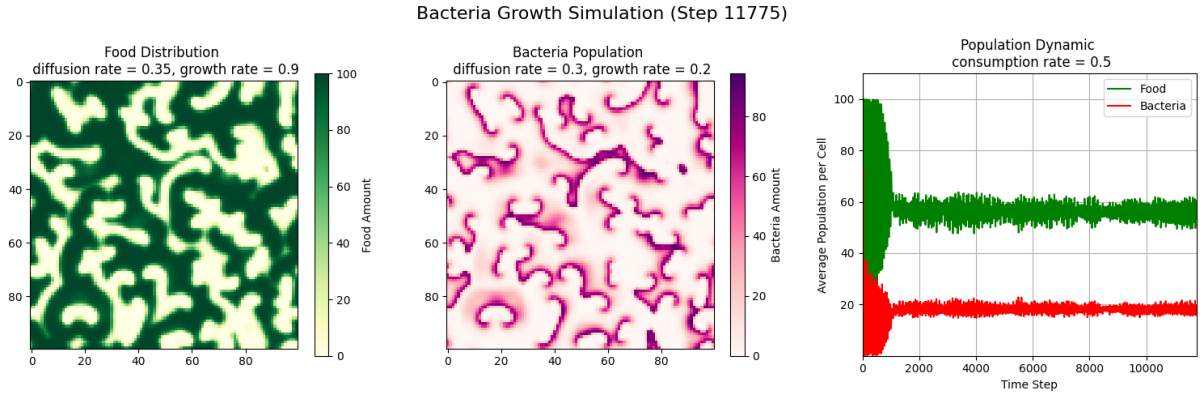


Figure 1: A simulation result that converges. Captures of step 10805 and 11775

When the population of bacteria and food converges, the animation of the grid dynamics shows that the density of bacteria and food both stays relatively constant (See figure 1). The oscillation remains relatively small and the pattern of the CA looks more scattered.

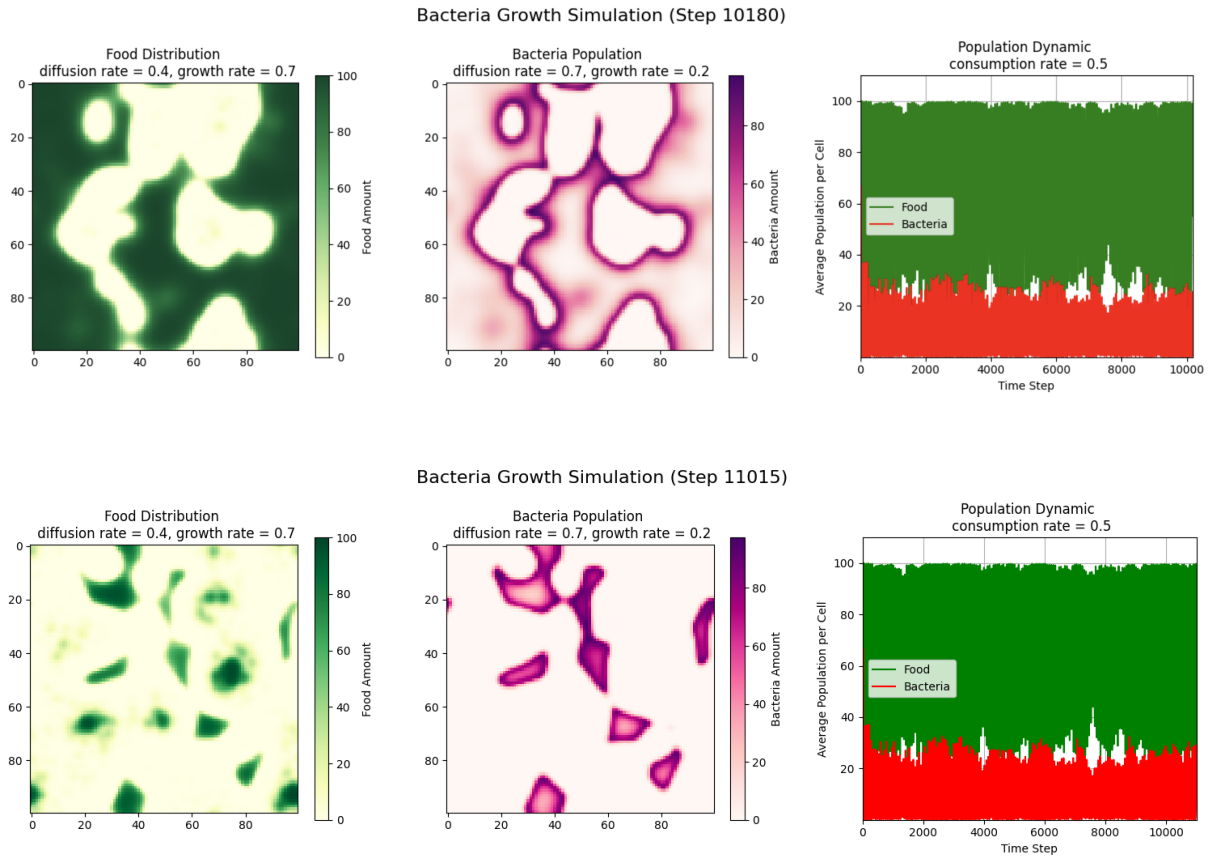


Figure 2: A simulation result does not converge. Captures of step 10180 and 11015

When the population of bacteria and food does not converge, the animation of the grid dynamics shows that the density of bacteria and food oscillates largely (See figure 2). From the animation we can observe that the food keeps blooming and dying out.

In both cases, we can see that bacteria are usually the most dense at the borders of food clusters. Then, the food regrows from the areas that are empty (no bacteria or food), then bacteria returns.

2.2 Relationship between variables

After simulation, I plotted the relationship between the five variables with the outputs — the food and bacteria population. All of the 5 parameters are set to 0.5 in the default mode.

Name	Variable	Value
Bacteria reproduction rate	bacteria_growth_rate	0.5
Food growth rate	food_growth_rate	0.5
Bacteria consumption rate	bacteria_consumption_rate	0.5
Food diffusion rate	food_diffusion_rate	0.5
Bacteria diffusion rate	bacteria_diffusion_rate	0.5

Table 3: Default parameters used for simulation.

I ran the simulation 100 times and each simulation went for 5000 steps (I wanted to run more steps but it ended up taking hours to run). The output of the last 500 steps were recorded and averaged to show the data after the system converges. The results are shown below.

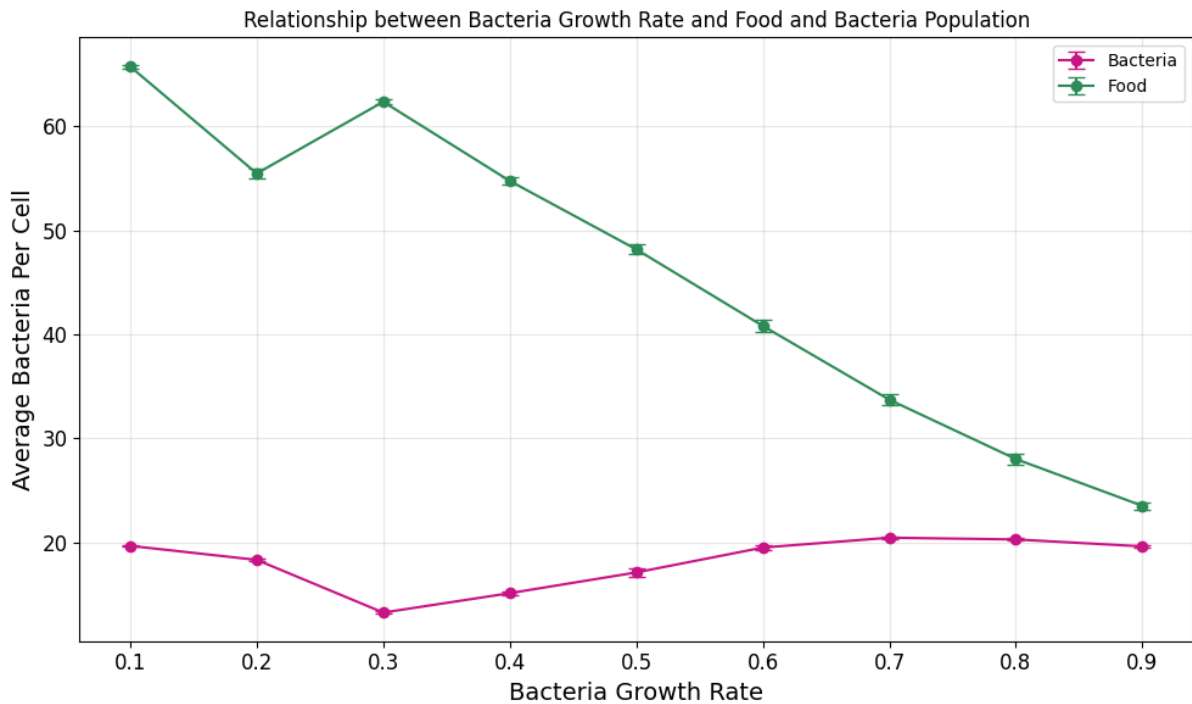


Figure 3: Relationship between bacteria growth rate and food and bacteria population

Figure 3 shows the bacteria growth rate does not actually influence the bacteria population that much — when bacteria growth rate is 0.1 the average bacteria per cell is around 20, which is the same as when it is 0.9. However, it influenced the average food amount per cell. I assume that the bacteria population is still constraint by settings like its diffusion rate and maximum food amount. However, when they have a higher growth rate, the bacteria population can grow too much and each out most of the food rapidly.

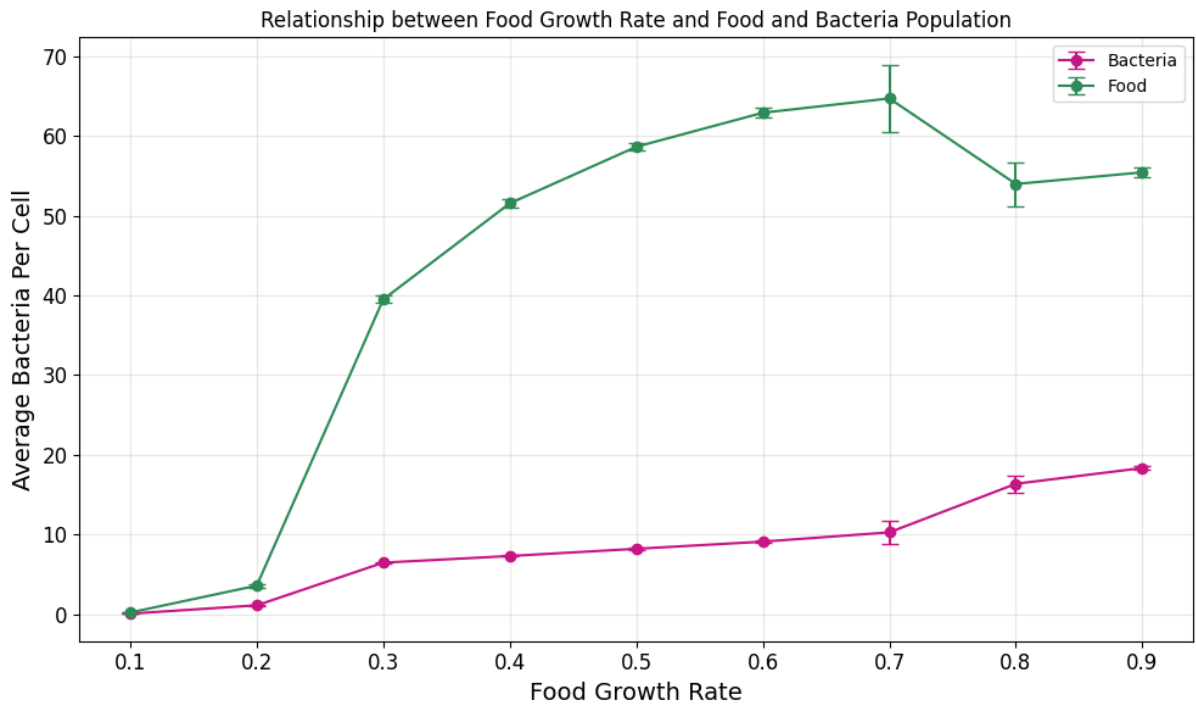


Figure 4: Relationship between food growth rate and food and bacteria population

Figure 4 shows the food growth rate influences the food population largely. The higher the food growth rate is, the higher the food population until it reaches around 0.7. More abundant food also higher the bacteria population, and perhaps this causes higher consumption rate that eventually lowers the food population after food growth rate reaches 0.7.

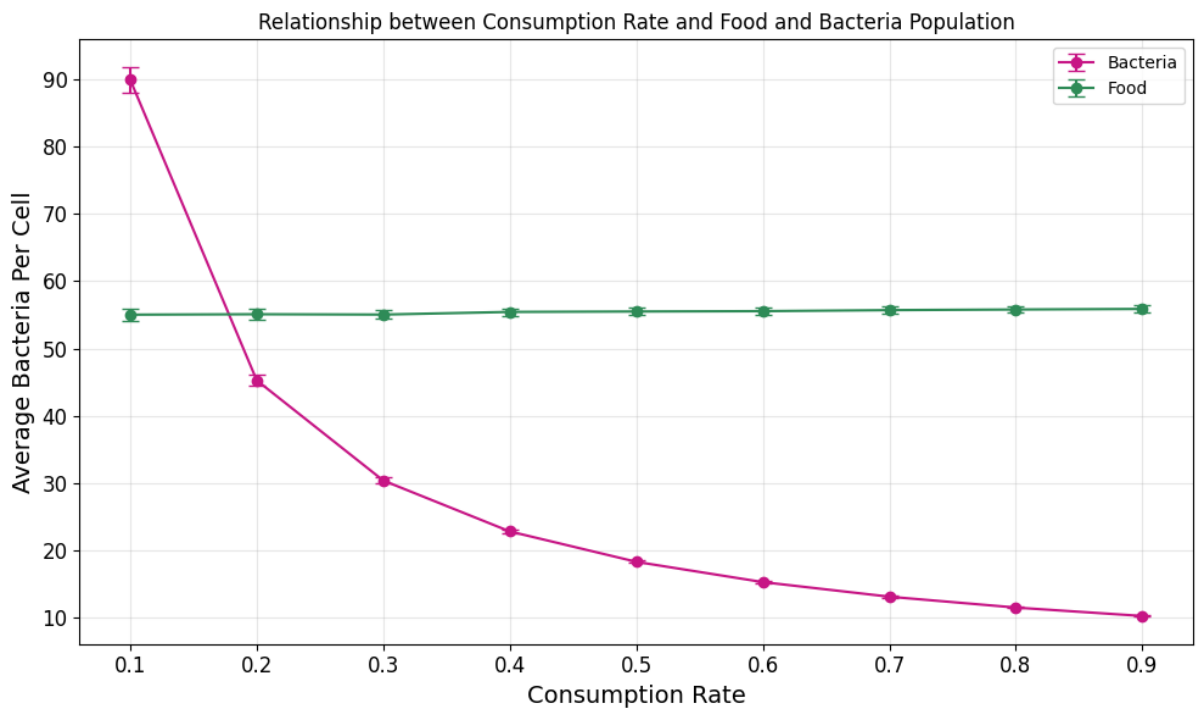


Figure 5: Relationship between consumption rate and food and bacteria population

Figure 5 shows that the bacteria growth rate almost does not influence the food population at all. However, given roughly the same amount of food available all time, the bacteria population decreases exponentially (judge by the shape) as the consumption rate grows.

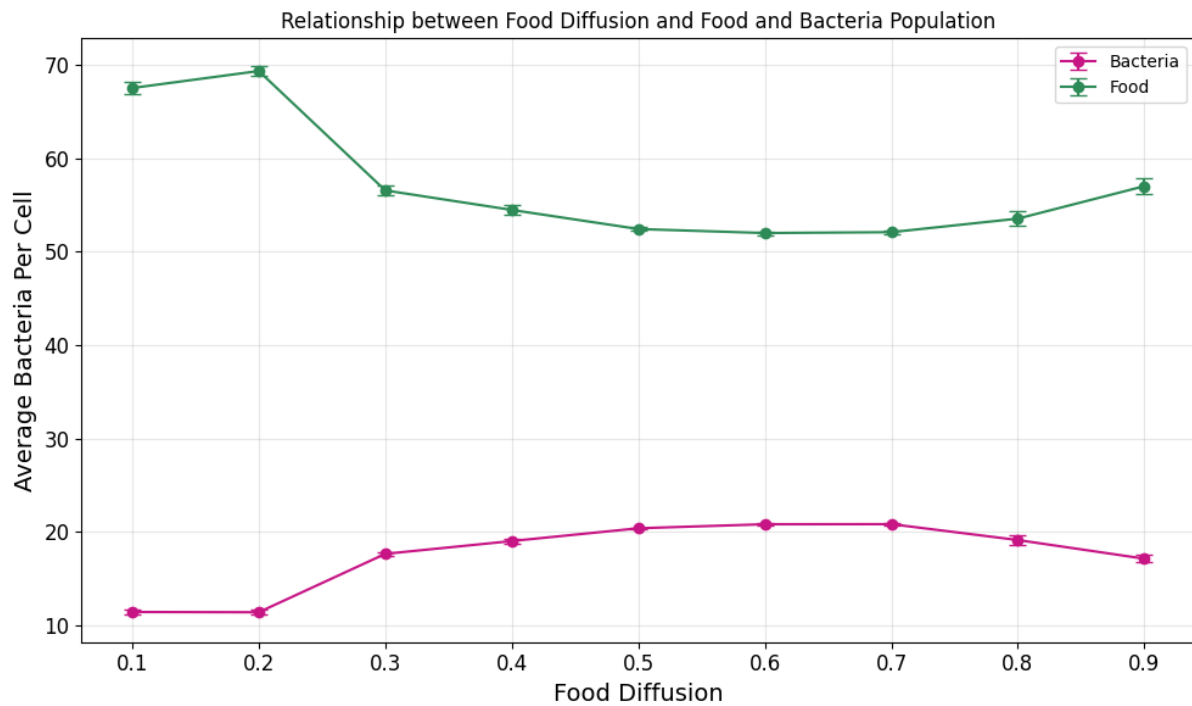


Figure 6: Relationship between food diffusion rate and food and bacteria population

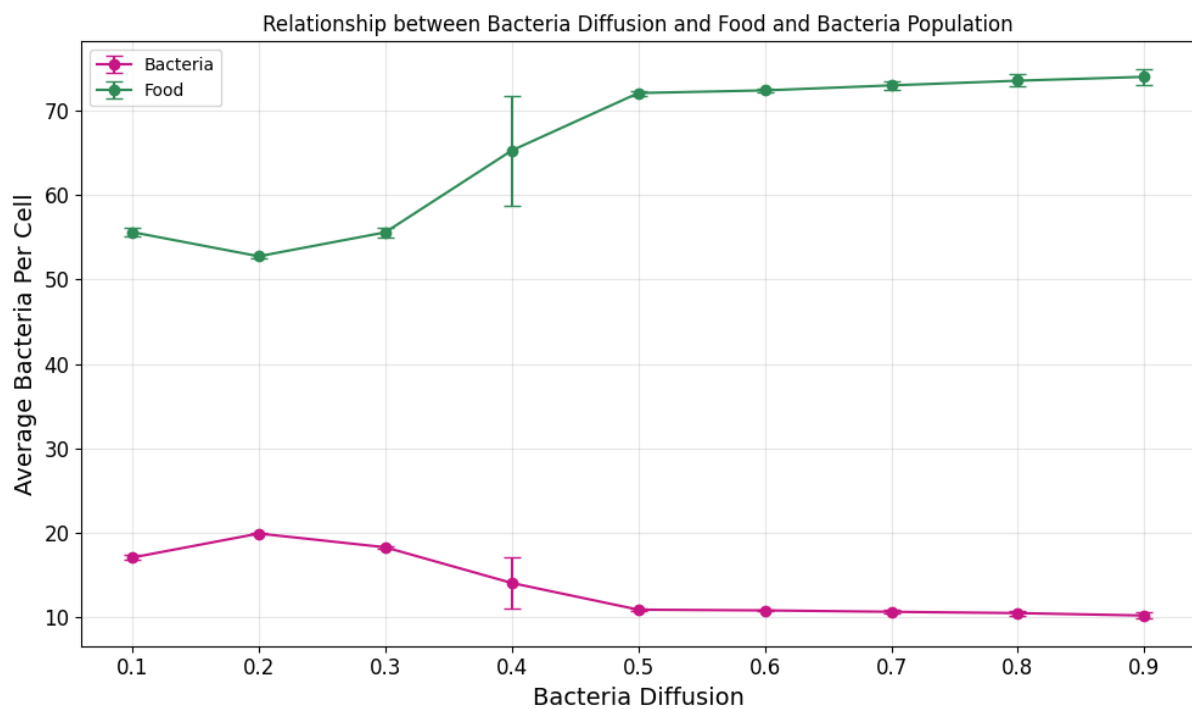


Figure 7: Relationship between bacteria diffusion rate and food and bacteria population

Figure 6 and 7 show that the diffusion rates might influence the two output populations roughly equally, as the bacteria and food population lines on both graphs seem symmetrical to each other.

3 Theoretical analysis

After exploring the relationship between input parameters and output populations, I first fit linear regression between every pair of them. The result is shown below:

	bacteria_correlation	bacteria_slope	bacteria_intercept	bacteria_r_squared	food_correlation	food_slope	food_intercept	food_r_squared
bacteria_growth_rate	0.445085	4.080122	16.108139	0.198100	-0.971953	-53.765603	72.728525	0.944693
food_growth_rate	0.966541	21.384267	-2.118268	0.934202	0.803780	72.345791	7.227771	0.646062
consumption_rate	-0.824998	-76.945612	67.043009	0.680622	0.971766	1.155175	54.876798	0.944329
food_diffusion	0.667625	9.023062	13.023957	0.445724	-0.694638	-16.801923	65.610579	0.482522
bacteria_diffusion	-0.878447	-12.356714	19.811900	0.771669	0.907580	29.598801	51.184762	0.823701

Figure 8: Linear regression between every pair of input and output

The R^2 are generally not very high, as we can see from figures above that the average population's reaction to different input parameters are very dynamic. However, the food growth rate to bacteria population has an R^2 score of 0.93 and a 96% correlation. The bacteria growth rate to food population has an R^2 of 0.94 and a -97% correlation. We can write the relationship mathematically as:

$$\text{AverageBacteriaPopulation} = \text{FoodGrowthRate} \times 21.38 - 2.11$$

This shows a positively correlated relationship with a slope of 21.38.

$$\text{AverageFoodPopulation} = \text{BacteriaGrowthRate} \times -53.76 + 72.72$$

This shows a negatively correlation relationship with a slope of -53.76, the output values are still all positive as *FoodGrowthRate* will not exceed 1.

Lastly, given the shape of the line in Figure 5, I fit an exponential model to the relationship between bacteria consumption rate and the average food amount per cell (see Figure 8).

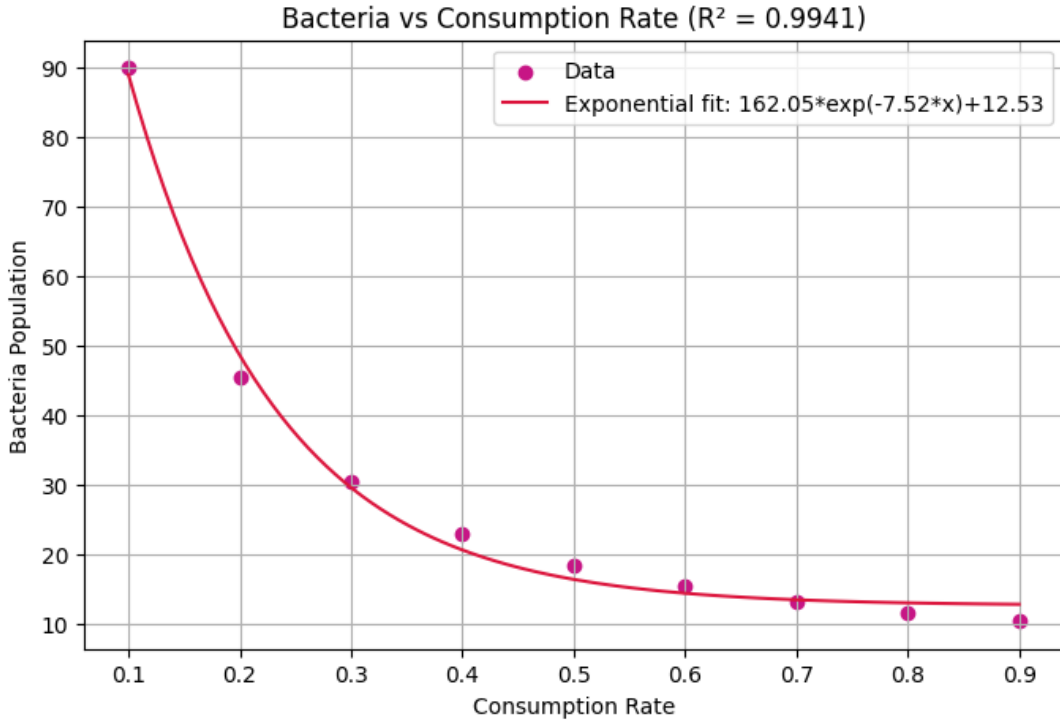


Figure 9: Exponential model between the average bacteria population and the bacteria

The R^2 score is 0.99, meaning that this exponential model can account for 99% of the change in output given the input. The mathematical relation is shown below:

$$\text{AverageBacteriaPopulation} = 162.05e^{-7.52 \cdot \text{BacteriaConsumptionRate}} + 12.53$$

4 Reference

- CS166 Session 6 - [3.2] Simulating cellular automata -for code that implements and animates CA

AI Statement

I used Grammarly to help correct my grammar mistakes. I used Claude to help debug and correct my code in Python. No other AI tools were used.

Collaboration Statement

I did not discuss, cross-check, or collaborate with anyone on this assignment.

5 Appendix

5.1 Appendix A - Main code for the bacteria growth model

```
1 class BacteriaGrowthSimulator:
2     """
3     A cellular automaton simulation of bacteria growth and food dynamics.
4     """
5
6     def __init__(self, food_growth_rate, food_diffusion_rate, bacteria_consumption_rate,
7                   bacteria_growth_rate, bacteria_diffusion_rate,
8                   reseed_prob=0.01, grid_size=100, food_max = 100, testing_mode = False):
9         """
10         Initialize the simulation with given parameters.
11
12         Parameters:
13             grid_size (int): Size of the square grid
14             food_growth_rate (float): Growth rate of food
15             food_max (float): Food capacity (maximum food per cell)
16             food_diffusion_rate (float): Food diffusion rate
17             bacteria_consumption (float): Bacteria consumption rate
18             bacteria_diffusion_rate (float): Bacteria diffusion rate
19             reseed_prob (float): Probability of food reseeding
20             food_prob (float): Probability of food in each cell
21             bacteria_prob (float): Probability of bacteria in each cell
22             bacteria_amount (float): Initial amount of bacteria if present
23         """
24         self.grid_size = grid_size
25         self.food = np.zeros((grid_size, grid_size))
26         self.bacteria = np.zeros((grid_size, grid_size))
27
28         # Parameters
```

```

28     self.food_growth_rate = food_growth_rate
29     self.food_max = food_max
30     self.food_diffusion_rate = food_diffusion_rate
31     self.bacteria_consumption_rate = bacteria_consumption_rate
32     self.bacteria_growth_rate = bacteria_growth_rate
33     self.bacteria_diffusion_rate = bacteria_diffusion_rate
34     self.reseed_prob = reseed_prob
35
36     # For tracking system dynamics
37     self.food_history = []
38     self.bacteria_history = []
39
40     # Figure for visualization
41     self.figure = None
42     self.axes = None
43
44     # Whether it's for testing
45     self.testing_mode = testing_mode
46
47 def initialize_ca(self, food_prob=0.3, bacteria_prob=0.1, bacteria_amount=20):
48     """
49     Initialize the grid with random food and bacteria.
50     """
51     # Initialize food randomly by generating a temporary grid for deciding where to put food
52     # Initiate with randomly chosen food amount between 0 and 100
53     food_mask = np.random.random((self.grid_size, self.grid_size)) < food_prob
54     self.food = np.where(food_mask, (self.food_max * np.random.random((self.grid_size, self.grid_size))), 0)
55
56     # Initialize bacteria randomly by generating a temporary grid for deciding where to put bacteria
57     # Initiate with 15 bacteria for the grids with bacteria
58     bacteria_mask = np.random.random((self.grid_size, self.grid_size)) < bacteria_prob
59     self.bacteria = np.where(bacteria_mask, bacteria_amount, 0)
60
61     # Record initial state
62     self.food_history = [np.mean(self.food)]
63     self.bacteria_history = [np.mean(self.bacteria)]
64
65     if self.testing_mode == True:
66         np.set_printoptions(threshold=20) # Ensures all elements are printed
67         print("Initialized CA")
68         print("----- Food distribution -----")
69         print(self.food)
70         print("----- Bacteria population -----")

```

```

71     print(self.bacteria)
72
73
74 def food_growth(self):
75     """
76     Increase food according to logistic growth formula.
77     """
78     self.food = self.food * (1 + self.food_growth_rate * (1 - self.food / self.food_max))
79     if self.testing_mode == True:
80         print("Food grew!")
81         print("----- Food distribution -----")
82         print(self.food)
83
84 def bacteria_reproduction(self):
85     """
86     Increase bacteria according to a fixed growth rate.
87     """
88     self.bacteria = self.bacteria * (1 + self.bacteria_growth_rate)
89     if self.testing_mode == True:
90         print("Bacteria reproduced!")
91         print("----- Bacteria population -----")
92         print(self.bacteria)
93
94 def food_reseeding(self):
95     """
96     Add 1 unit of food to each cell with probability reseed_prob.
97     """
98     # Decide where to reseed food by generating a temporary grid
99     reseed_mask = np.random.random((self.grid_size, self.grid_size)) <
100     self.reseed_prob
101     self.food = np.where(reseed_mask, self.food + 1, self.food)
102
103     # Ensure food doesn't exceed capacity - over food_max values will be turned to
104     food_max
105     self.food = np.clip(self.food, 0, self.food_max)
106     if self.testing_mode == True:
107         print("Food reseed!")
108         print("----- Food distribution -----")
109         print(self.food)
110
111 def diffusion(self, current_config, diffusion_rate):
112     """
113     Apply diffusion to food or bacteria.
114
115     Parameters:
116         current_config (numpy.ndarray): Grids (100 x 100) to apply diffusion to

```

```

115         diffusion_rate (float): Rate of diffusion
116
117     Returns:
118         numpy.ndarray: Updated grid after diffusion
119     """
120     # Amount that diffuses out of each cell
121     diffusion_amount = current_config * diffusion_rate
122
123     # Amount that stays in each cell
124     next_config = current_config * (1 - diffusion_rate)
125
126     # Amount that diffuses to each of the 4 neighbors (von Neumann neighborhood)
127     diff_per_neighbor = diffusion_amount / 4
128
129     # Use roll to shift the grid and add diffusion to neighbors
130     next_config += np.roll(diff_per_neighbor, 1, axis=0) # from cell above
131     next_config += np.roll(diff_per_neighbor, -1, axis=0) # from cell below
132     next_config += np.roll(diff_per_neighbor, 1, axis=1) # from cell left
133     next_config += np.roll(diff_per_neighbor, -1, axis=1) # from cell right
134
135     return next_config
136
137 def food_diffusion(self):
138     """
139     Apply diffusion to the food grid.
140     """
141     self.food = self.diffusion(self.food, self.food_diffusion_rate)
142     if self.testing_mode == True:
143         print("Food diffusion!")
144         print("----- Food distribution -----")
145         print(self.food)
146
147 def bacteria_diffusion(self):
148     """
149     Apply diffusion to the bacteria grid.
150     """
151     self.bacteria = self.diffusion(self.bacteria, self.bacteria_diffusion_rate)
152     if self.testing_mode == True:
153         print("Bacteria diffusion!")
154         print("----- Bacteria population -----")
155         print(self.bacteria)
156
157 def consumption_and_starvation(self):
158     """
159     Handle food consumption, bacteria starvation, and reproduction.
160     """

```

```

161 # Calculate how much food bacteria need
162 food_needed = self.bacteria * self.bacteria_consumption_rate
163
164 # Calculate how much food is actually available (minimum of needed and available)
165 food_consumed = np.minimum(food_needed, self.food)
166
167 # Update food after consumption
168 self.food -= food_consumed
169
170 if self.testing_mode == True:
171     print("Bacteria consume food! yum!")
172     print("----- Food distribution -----")
173     print(self.food)
174
175 # Calculate how many bacteria survive (those that got food)
176 die_out_population = np.ceil((food_needed - food_consumed) /
177     self.bacteria_consumption_rate).astype(int)
178 self.food = np.clip(self.food, 0, self.food_max)
179
180 # Make sure there is no negative bacteria
181 assert np.all(self.bacteria >= 0), "Error: Negative bacteria count detected!"
182
183 # Update bacteria population after starvation
184 self.bacteria -= die_out_population
185 self.bacteria = np.clip(self.bacteria, 0, None)
186
187 if self.testing_mode == True:
188     print("Some bacteria died!")
189     print("----- Bacteria population -----")
190     print(self.bacteria)
191
192 def update(self):
193     """
194     Perform one step of the simulation.
195     """
196
197     # Bacteria consume food, and die out if they don't have enough food
198     self.consumption_and_starvation()
199
200     # The remaining bacteria reproduce
201     self.bacteria_reproduction()
202
203     # The remaining food grows
204     self.food_growth()
205
206     # Food reseeds with a probability

```

```

206     self.food_reseeding()
207
208     # Food diffuses
209     self.food_diffusion()
210
211     # Bacteria diffuses
212     self.bacteria_diffusion()
213
214     # Record history
215     self.food_history.append(np.mean(self.food))
216     self.bacteria_history.append(np.mean(self.bacteria))
217
218     if self.testing_mode == True:
219         print("One step done!")
220
221 def run_simulation(self, steps):
222     """
223     Run the simulation for a specified number of steps.
224
225     Parameters:
226         steps (int): Number of steps to run
227
228     Returns:
229         tuple: (food_history, bacteria_history) - Lists of average food and bacteria over
230         time
231     """
232     for _ in range(steps):
233         self.update()
234
235     return self.food_history, self.bacteria_history
236
237 def setup_visualization(self):
238     """Setup the figure and axes for visualization."""
239     self.figure, self.axes = plt.subplots(1, 3, figsize=(15, 5))
240
241     # Title for the whole figure
242     self.figure.suptitle('Bacteria Growth Simulation', fontsize=16)
243
244     # Setup individual plots
245     self.axes[0].set_title(f'Food Distribution \n diffusion rate =
246     {self.food_diffusion_rate}, growth rate = {self.food_growth_rate}')
247     self.food_plot = self.axes[0].imshow(self.food, cmap='YlGn', vmin=0,
248     vmax=self.food_max)
249     self.figure.colorbar(self.food_plot, ax=self.axes[0], label='Food Amount')

```

```

248 self.axes[1].set_title(f'Bacteria Population \n diffusion rate =
    {self.bacteria_diffusion_rate}, growth rate = {self.bacteria_growth_rate}')
249 self.bacteria_plot = self.axes[1].imshow(self.bacteria, cmap='RdPu', vmin=0)
250 self.figure.colorbar(self.bacteria_plot, ax=self.axes[1], label='Bacteria
    Amount')
251
252 self.axes[2].set_title(f'Population Dynamic \n consumption rate =
    {self.bacteria_consumption_rate}')
253 self.axes[2].set_xlabel('Time Step')
254 self.axes[2].set_ylabel('Average Population per Cell')
255 self.axes[2].grid(True)
256
257 # Line plots for average food and bacteria over time
258 self.food_line, = self.axes[2].plot([], [], 'g-', label='Food')
259 self.bacteria_line, = self.axes[2].plot([], [], 'r-', label='Bacteria')
260 self.axes[2].legend()
261
262 plt.tight_layout()
263
264 def observe(self):
265     """Update the visualization with current state."""
266     # Update the grid plots
267     self.food_plot.set_data(self.food)
268     self.bacteria_plot.set_data(self.bacteria)
269
270     # Set appropriate range for bacteria plot
271     self.bacteria_plot.set_clim(0, np.max(self.bacteria))
272
273     # Update the line plots
274     x = range(len(self.food_history))
275     self.food_line.set_data(x, self.food_history)
276     self.bacteria_line.set_data(x, self.bacteria_history)
277
278     # Adjust y-axis limits if needed
279     if len(self.food_history) > 1:
280         max_val = max(max(self.food_history), max(self.bacteria_history)) * 1.1
281         min_val = min(min(self.food_history), min(self.bacteria_history)) * 0.9
282         self.axes[2].set_ylim(min_val, max_val)
283         self.axes[2].set_xlim(0, len(self.food_history))
284
285     # Add step count to title
286     self.figure.suptitle(f'Bacteria Growth Simulation (Step
        {len(self.food_history)-1})', fontsize=16)
287

```


5.2 Appendix B - Plotting and Animating the model

```
1
2 def make_animation(simulation, total_frames, steps_per_frame=1, interval=50):
3     """
4     Create an animation of the simulation.
5
6     Parameters:
7         simulation (BacteriaGrowthSimulator): The simulator to animate
8         total_frames (int): Total number of frames in the animation
9         steps_per_frame (int): Simulation steps per animation frame
10        interval (int): Time between frames in milliseconds
11
12    Returns:
13        HTML: Animation that can be displayed in Jupyter notebook
14    """
15    # Setup visualization
16    simulation.setup_visualization()
17
18    # Animation update function
19    def update(frame):
20        for _ in range(steps_per_frame):
21            simulation.update()
22            progress_bar.update(1)
23        return simulation.observe()
24
25    # Initialize progress bar
26    progress_bar = tqdm(total=total_frames)
27
28    # Create animation
29    animation = FuncAnimation(
30        simulation.figure, update, frames=total_frames,
31        interval=interval, blit=True
32    )
33
34    # Convert to HTML for display
35    html_animation = HTML(animation.to_jshtml())
36
37    return html_animation
38
39 def plot_population_dynamics(simulator):
40     """
41     Plot the food and bacteria population dynamics over time.
42
43     Parameters:
44         simulator (BacteriaGrowthSimulator): The simulator after running simulation
```

```

45     """
46     plt.figure(figsize=(10, 6))
47     plt.plot(simulator.food_history, 'g-', label='Food')
48     plt.plot(simulator.bacteria_history, 'r-', label='Bacteria')
49     plt.xlabel('Time Step')
50     plt.ylabel('Average Population per Cell')
51     plt.title('Population Dynamics')
52     plt.grid(True)
53     plt.legend()
54     plt.tight_layout()
55     plt.show()
56

```

5.3 Appendix C - Simulation for empirical analysis

```

1  import numpy as np
2  import matplotlib.pyplot as plt
3  from scipy import stats
4  import pandas as pd
5  from matplotlib.ticker import MaxNLocator
6  from tqdm import tqdm
7
8  # Function to run the simulation with varying parameters
9  def run_simulation(parameter_name, parameter_values, trials = 100):
10     """
11     Run the bacteria growth simulation with varying parameter values
12     and collect data on the converged population sizes.
13
14     Parameters:
15     - parameter_name: Name of the parameter being varied
16     - parameter_values: List of values for the parameter
17     - trials: Number of simulation runs per parameter value for statistical analysis
18     - grid_size: Size of the simulation grid
19     - steps: Number of simulation steps
20
21     Returns:
22     - DataFrame with parameter values and corresponding population statistics
23     """
24
25     results = []
26
27     for value in tqdm(parameter_values, desc=f"Testing {parameter_name}"):
28         bacteria_final = []
29         food_final = []
30
31         for trial in range(trials):

```

```

32
33 if parameter_name == "bacteria_growth_rate":
34     # Create a simulator with default parameters
35     simulator = BacteriaGrowthSimulator(
36         food_growth_rate=0.9, # Food growth rate
37         food_diffusion_rate=0.35, # Food diffusion rate
38         bacteria_consumption_rate=0.5, # Bacteria consumption rate
39         bacteria_growth_rate=value, # Bacteria growth rate
40         bacteria_diffusion_rate=0.3, # Bacteria diffusion rate
41         grid_size = 100
42     )
43
44     # Initialize with random food and bacteria
45     simulator.initialize_ca(food_prob=0.5, bacteria_prob=0.5,
46         bacteria_amount=20)
47     food_history, bacteria_history = simulator.run_simulation(steps=5000)
48
49 elif parameter_name == "food_growth_rate":
50     # Create a simulator with default parameters
51     simulator = BacteriaGrowthSimulator(
52         food_growth_rate=value, # Food growth rate
53         food_diffusion_rate=0.35, # Food diffusion rate
54         bacteria_consumption_rate=0.5, # Bacteria consumption rate
55         bacteria_growth_rate=0.2, # Bacteria growth rate
56         bacteria_diffusion_rate=0.3, # Bacteria diffusion rate
57         grid_size = 100
58     )
59
60     # Initialize with random food and bacteria
61     simulator.initialize_ca(food_prob=0.5, bacteria_prob=0.5,
62         bacteria_amount=20)
63     food_history, bacteria_history = simulator.run_simulation(steps=5000)
64
65 elif parameter_name == "consumption_rate":
66     # Create a simulator with default parameters
67     simulator = BacteriaGrowthSimulator(
68         food_growth_rate=0.9, # Food growth rate
69         food_diffusion_rate=0.35, # Food diffusion rate
70         bacteria_consumption_rate=value, # Bacteria consumption rate
71         bacteria_growth_rate=0.2, # Bacteria growth rate
72         bacteria_diffusion_rate=0.3, # Bacteria diffusion rate
73         grid_size = 100
74     )
75
76     # Initialize with random food and bacteria

```

```

75     simulator.initialize_ca(food_prob=0.5, bacteria_prob=0.5,
76     bacteria_amount=20)
77
78     food_history, bacteria_history = simulator.run_simulation(steps=5000)
79
80 elif parameter_name == "food_diffusion":
81     # Create a simulator with default parameters
82     simulator = BacteriaGrowthSimulator(
83         food_growth_rate=0.9, # Food growth rate
84         food_diffusion_rate=value, # Food diffusion rate
85         bacteria_consumption_rate=0.5, # Bacteria consumption rate
86         bacteria_growth_rate=0.2, # Bacteria growth rate
87         bacteria_diffusion_rate=0.3, # Bacteria diffusion rate
88         grid_size = 100
89     )
90
91     # Initialize with random food and bacteria
92     simulator.initialize_ca(food_prob=0.5, bacteria_prob=0.5,
93     bacteria_amount=20)
94     food_history, bacteria_history = simulator.run_simulation(steps=5000)
95
96 elif parameter_name == "bacteria_diffusion":
97     # Create a simulator with default parameters
98     simulator = BacteriaGrowthSimulator(
99         food_growth_rate=0.9, # Food growth rate
100         food_diffusion_rate=0.35, # Food diffusion rate
101         bacteria_consumption_rate=0.5, # Bacteria consumption rate
102         bacteria_growth_rate=0.2, # Bacteria growth rate
103         bacteria_diffusion_rate=value, # Bacteria diffusion rate
104         grid_size = 100
105     )
106
107     # Initialize with random food and bacteria
108     simulator.initialize_ca(food_prob=0.5, bacteria_prob=0.5,
109     bacteria_amount=20)
110     food_history, bacteria_history = simulator.run_simulation(steps=5000)
111
112 else:
113     # Default pattern
114     # Create a simulator with default parameters
115     simulator = BacteriaGrowthSimulator(
116         food_growth_rate=0.9, # Food growth rate
117         food_diffusion_rate=0.35, # Food diffusion rate
118         bacteria_consumption_rate=0.5, # Bacteria consumption rate
119         bacteria_growth_rate=0.2, # Bacteria growth rate
120         bacteria_diffusion_rate=0.3, # Bacteria diffusion rate
121         grid_size = 100

```

```

118         )
119
120         # Initialize with random food and bacteria
121         simulator.initialize_ca(food_prob=0.5, bacteria_prob=0.5,
122                                bacteria_amount=20)
123
124         # Ensure values are positive
125         food_history = food_history[4500:]
126         bacteria_history = bacteria_history[4500:]
127
128         food_final.append(np.mean(food_history))
129         bacteria_final.append(np.mean(bacteria_history))
130
131         # Calculate statistics for this parameter value
132         bacteria_mean = np.mean(bacteria_final)
133         bacteria_std = np.std(bacteria_final)
134         food_mean = np.mean(food_final)
135         food_std = np.std(food_final)
136
137         results.append({
138             parameter_name: value,
139             'bacteria_mean': bacteria_mean,
140             'bacteria_std': bacteria_std,
141             'food_mean': food_mean,
142             'food_std': food_std,
143             'total_result': (bacteria_final, food_final)
144         })
145
146         return pd.DataFrame(results)
147
148     # Define the parameters to investigate
149     parameters = {
150         'bacteria_growth_rate': np.linspace(0.1, 0.9, 9),
151         'food_growth_rate': np.linspace(0.1, 0.9, 9),
152         'consumption_rate': np.linspace(0.1, 0.9, 9),
153         'food_diffusion': np.linspace(0.1, 0.9, 9),
154         'bacteria_diffusion': np.linspace(0.1, 0.9, 9),
155     }
156
157     # Run simulations and collect results
158     all_results = {}
159     for param_name, param_values in parameters.items():
160         all_results[param_name] = run_simulation(param_name, param_values)

```

5.4 Appendix D - Using the empirical analysis result for theoretical analysis

```
1 # Function to analyze the relationship between parameters and population sizes
2 def analyze_relationships(results):
3     """
4     Analyze and quantify the relationships between parameters and population sizes.
5     """
6     analysis_results = {}
7
8     for param_name, df in results.items():
9         # Calculate correlation
10        bacteria_corr = np.corrcoef(df[param_name], df['bacteria_mean'])[0, 1]
11        food_corr = np.corrcoef(df[param_name], df['food_mean'])[0, 1]
12
13        # Fit linear regression
14        bacteria_slope, bacteria_intercept, bacteria_r, _, _ =
15        stats.linregress(df[param_name], df['bacteria_mean'])
16        food_slope, food_intercept, food_r, _, _ = stats.linregress(df[param_name],
17        df['food_mean'])
18
19        analysis_results[param_name] = {
20            'bacteria_correlation': bacteria_corr,
21            'bacteria_slope': bacteria_slope,
22            'bacteria_intercept': bacteria_intercept,
23            'bacteria_r_squared': bacteria_r**2,
24            'food_correlation': food_corr,
25            'food_slope': food_slope,
26            'food_intercept': food_intercept,
27            'food_r_squared': food_r**2
28        }
29
30    return pd.DataFrame.from_dict(analysis_results, orient='index')
31
32 # Analyze the relationships
33 relationship_analysis = analyze_relationships(all_results)
34 relationship_analysis.head(5)
```

5.5 Appendix E - Fitting an exponential line to the model

```
1 from scipy.optimize import curve_fit
2
3 def fit_exponential_to_consumption_rate(results):
4     # Get consumption rate data
5     df = results['consumption_rate']
6     x = df['consumption_rate'].values
```

```

7 y = df['bacteria_mean'].values
8
9 # Define exponential function
10 def exp_func(x, a, b, c):
11     return a * np.exp(b * x) + c
12
13 # Fit the model
14 params, _ = curve_fit(exp_func, x, y, p0=[50.0, -5.0, 5.0], maxfev=10000)
15 a, b, c = params
16
17 # Calculate R-squared
18 y_pred = exp_func(x, a, b, c)
19 ss_res = np.sum((y - y_pred)**2)
20 ss_tot = np.sum((y - np.mean(y))**2)
21 r_squared = 1 - (ss_res / ss_tot)
22
23 # Create a simple plot
24 plt.figure(figsize=(8, 5))
25 plt.scatter(x, y, color='mediumvioletred', label='Data')
26
27 # Plot the fitted curve
28 x_curve = np.linspace(min(x), max(x), 100)
29 y_curve = exp_func(x_curve, a, b, c)
30 plt.plot(x_curve, y_curve, 'crimson', label=f'Exponential fit:
31 {a:.2f}*exp({b:.2f}*x)+{c:.2f}')
32
33 plt.xlabel('Consumption Rate')
34 plt.ylabel('Bacteria Population')
35 plt.title(f'Bacteria vs Consumption Rate ( $R^2 = {r\_squared:.4f}$ )')
36 plt.legend()
37 plt.grid(True)
38
39 # Return just the key results
40 return {
41     'r_squared': r_squared,
42     'a': a,
43     'b': b,
44     'c': c,
45     'equation': f'{a:.4f} * exp({b:.4f} * x) + {c:.4f}'
46 }
47
48 results = fit_exponential_to_consumption_rate(all_results)
49 print(results)

```