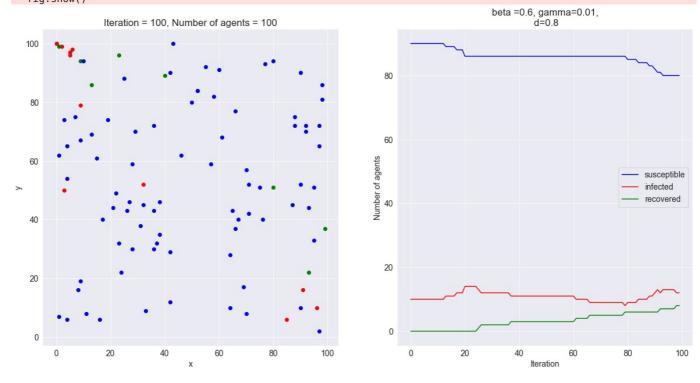
Exercise 11.1

```
In [1]:
         import numpy as np
         import matplotlib.pyplot as plt
         import pylab
         import seaborn as sns
         import random
         import math
In [2]:
         no of individuals = 100
         no_of_iterations = 100
         diffusion rate = 0.8 # diffusion probability
         initial infection rate = 0.1
         beta = \overline{0.6} #Infection Probability
gamma = 0.01 # Recovery Probability
         grid x = 100 # max coordinate in x
         grid_y = 100 #max coordinate in y
         # status 1 = susceptible
         # status 2 = infected
         # status 3 = immune (recovered)
         individual position = []
         individual status = []
         final_status = np.zeros((3, no_of_iterations))
         for i in range(no of individuals):
             individual_position.append([random.randint(0,grid_x), random.randint(0,grid_y)])
             individual_status.append(1)
         initially_infected_individuals = random.sample(range(no_of_individuals), round(no_of_individuals*initial_infection)
         for j in initially_infected_individuals:
             individual_status[j] = 2
         for m in range(no_of_iterations):
             for n in range(no_of_individuals):
                  rand1 = np.random.rand()
                  if rand1 < diffusion_rate:</pre>
                      motion = np.random.choice([1, 2, 3, 4])
                      if motion == 1:
                          if individual_position[n][0] < grid_x:</pre>
                              individual_position[n][0] = individual_position[n][0] + 1
                              individual_position[n][0] = 0
                      elif motion == 2:
                          if individual position[n][0] > 0:
                              individual_position[n][0] = individual_position[n][0] - 1
                          else:
                              individual position[n][0] = grid x
                      elif motion == 3:
                          if individual position[n][1] < grid y:</pre>
                              individual position[n][1] = individual position[n][1] + 1
                          else:
                              individual_position[n][0] = 0
                      elif motion == 4:
                          if individual_position[n][1] > 0:
                              individual position[n][1] = individual position[n][1] - 1
                          else:
                              individual_position[n][0] = grid_y
             for i in range(no_of_individuals):
                  if individual status[i] == 2:
                      infected individual position = individual position[i]
                      rand2 = np.random.rand()
                      if rand2 < beta:</pre>
                          for j in range(no_of_individuals):
                              if individual_position[j] == infected_individual_position and individual_status[j] == 1:
                                  individual_status[j] = 2
                      rand3 = np.random.rand()
                      if rand3 < gamma:</pre>
                          individual_status[i] = 3
             for i in range(no_of_individuals):
                  if individual status[i] == 1:
                      final status[0][m] = final status[0][m] + 1
                  if individual_status[i] == 2:
                      final_status[1][m] = final_status[1][m] + 1
                  if individual status[i] == 3:
```

```
final status[2][m] = final status[2][m] + 1
sns.set style("darkgrid")
fig = plt.figure(figsize=(20,10))
fig.tight_layout()
plt.rcParams.update({'font.size': 14})
ax1 = fig.add_subplot(121)
ax1.clear()
# ax1.set_ylim(0, 10)
# ax1.set_xlim(0, 10)
for i in range(no of individuals):
    if individual status[i] == 1:
         \verb|plt.scatter(individual_position[i][0], individual_position[i][1], color = "b"|)|
    elif individual status[i] == 2:
        plt.scatter(individual_position[i][0],individual position[i][1], color = "r")
    elif individual_status[i] == 3:
         plt.scatter(individual_position[i][0],individual_position[i][1], color = "g")
ax1.set_xlabel('x')
ax1.set ylabel('y')
ax1.set_title(f'Iteration = {no_of_iterations}, Number of agents = {no_of_individuals}')
ax1 = fig.add_subplot(122)
ax1.plot(range(no_of_iterations), final_status[0][:], "b", label = "susceptible")
ax1.plot(range(no_of_iterations), final_status[1][:], "r", label = "infected")
ax1.plot(range(no of iterations), final status[2][:], "g", label = "recovered")
ax1.set_xlabel('Iteration')
ax1.set_ylabel('Number of agents')
plt.legend(ncol = 1, loc = "center right")
ax1.set title(f" beta ={beta}, gamma={gamma}, \n d={diffusion rate}")
fig.show()
```

<ipython-input-2-e54fcf30fc37>:108: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_i
nline, which is a non-GUI backend, so cannot show the figure.
fig.show()



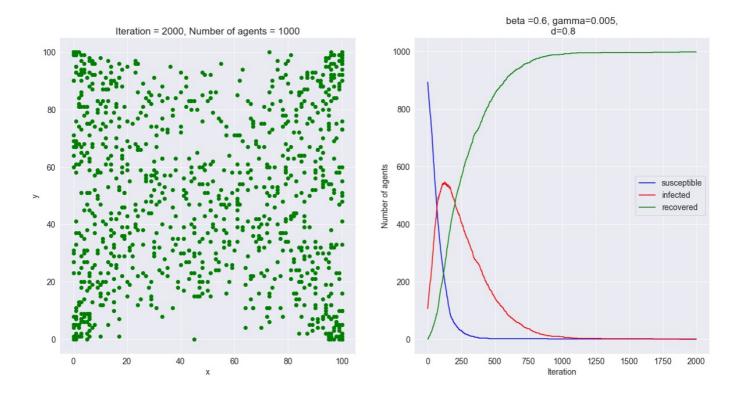
```
no_of_individuals = 1000
no_of_iterations = 2000
diffusion_rate = 0.8  # diffusion probability
initial_infection_rate = 0.1
beta = 0.6  #Infection Probability
gamma = 0.005  # Recovery Probability
grid_x = 100  # max coordinate in x
grid_y = 100  #max coordinate in y

# status 1 = susceptible
# status 2 = infected
# status 3 = immune (recovered)
individual_position = []
```

```
individual status = []
 final_status = np.zeros((3, no_of_iterations))
for i in range(no of individuals):
         individual_position.append([random.randint(0,grid_x), random.randint(0,grid_y)])
         individual_status.append(1)
initially_infected_individuals = random.sample(range(no_of_individuals), round(no_of_individuals*initial_infection)
for j in initially_infected_individuals:
         individual status[j] = 2
for m in range(no_of_iterations):
         for n in range(no_of_individuals):
                  rand1 = np.random.rand()
                  if rand1 < diffusion_rate:</pre>
                           motion = np.random.choice([1, 2, 3, 4])
                           if motion == 1:
                                   if individual_position[n][0] < grid_x:</pre>
                                            individual_position[n][0] = individual_position[n][0] + 1
                                             individual position[n][0] = 0
                           elif motion == 2:
                                   if individual_position[n][0] > 0:
                                           individual_position[n][0] = individual_position[n][0] - 1
                                    else:
                                            individual position[n][0] = grid x
                           elif motion == 3:
                                   if individual position[n][1] < grid y:</pre>
                                            individual position[n][1] = individual position[n][1] + 1
                                   else:
                                            individual_position[n][0] = 0
                           elif motion == 4:
                                   if individual position[n][1] > 0:
                                            individual position[n][1] = individual position[n][1] - 1
                                   else:
                                            individual_position[n][0] = grid_y
         for i in range(no of individuals):
                  if individual status[i] == 2:
                           infected individual position = individual position[i]
                           rand2 = np.random.rand()
                           if rand2 < beta:</pre>
                                   for j in range(no_of_individuals):
                                            \textbf{if} \  \, \text{individual\_position} \, [\texttt{j}] \, = \, \text{infected\_individual\_position} \, \, \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_status} [\texttt{j}] \, = \, 1 : \, \\ \textbf{and} \, \, \text{individual\_s
                                                    individual status[j] = 2
                           rand3 = np.random.rand()
                           if rand3 < gamma:</pre>
                                   individual status[i] = 3
         for i in range(no of individuals):
                  if individual status[i] == 1:
                           final status[0][m] = final status[0][m] + 1
                  if individual_status[i] == 2:
                           final_status[1][m] = final_status[1][m] + 1
                  if individual status[i] == 3:
                           final_status[2][m] = final_status[2][m] + 1
sns.set_style("darkgrid")
fig = plt.figure(figsize=(20,10))
fig.tight layout()
plt.rcParams.update({'font.size': 14})
ax1 = fig.add_subplot(121)
ax1.clear()
# ax1.set_ylim(0, 10)
# ax1.set xlim(0, 10)
for i in range(no of individuals):
         if individual status[i] == 1:
                  plt.scatter(individual_position[i][0],individual_position[i][1], color = "b")
         elif individual_status[i] == 2:
                 \verb|plt.scatter(individual_position[i][0], individual_position[i][1], | color = "r"|)|
         elif individual_status[i] == 3:
                 plt.scatter(individual_position[i][0],individual_position[i][1], color = "g")
ax1.set xlabel('x'
ax1.set_ylabel('y')
ax1.set_title(f'Iteration = {no_of_iterations}, Number of agents = {no_of_individuals}')
ax1 = fig.add subplot(122)
ax1.plot(range(no_of_iterations), final_status[0][:], "b", label = "susceptible")
ax1.plot(range(no_of_iterations), final_status[1][:], "r", label = "infected")
ax1.plot(range(no_of_iterations), final_status[2][:], "g", label = "recovered")
ax1.set_xlabel('Iteration')
```

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ax1.set_ylabel('Number of agents')
plt.legend(ncol = 1, loc = "center right")
ax1.set_title(f" beta ={beta}, gamma={gamma}, \n d={diffusion_rate}")
fig.show()
```

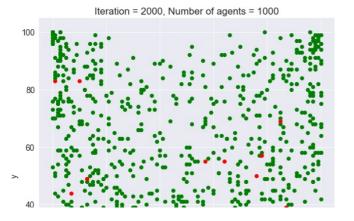
<ipython-input-3-9476ed2905c1>:108: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_i
nline, which is a non-GUI backend, so cannot show the figure.
fig.show()

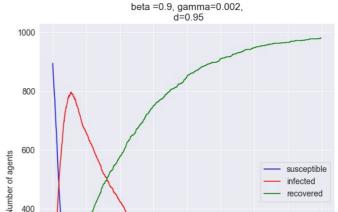


```
In [4]:
         no of individuals = 1000
         no of iterations = 2000
         diffusion_rate = 0.95 # diffusion probability
         initial_infection_rate = 0.1
         beta = 0.9
                     #Infection Probability
         gamma = 0.002 # Recovery Probability
         grid_x = 100 \# max coordinate in x
         grid\ y = 100 \ \#max\ coordinate\ in\ y
         # status 1 = susceptible
         # status 2 = infected
         # status 3 = immune (recovered)
         individual_position = []
         individual status = []
         final_status = np.zeros((3, no_of_iterations))
         for i in range(no of individuals):
             individual_position.append([random.randint(0,grid_x), random.randint(0,grid_y)])
             individual status.append(1)
         initially infected individuals = random.sample(range(no of individuals), round(no of individuals*initial infection
         for j in initially_infected_individuals:
             individual status[j] = 2
         for m in range(no_of_iterations):
             for n in range(no_of_individuals):
                 rand1 = np.random.rand()
                 if rand1 < diffusion_rate:</pre>
                     motion = np.random.choice([1, 2, 3, 4])
                     if motion == 1:
                         if individual position[n][0] < grid x:</pre>
                             individual_position[n][0] = individual_position[n][0] + 1
                         else:
                             individual position[n][0] = 0
                     elif motion == 2:
                         if individual position[n][0] > 0:
                             individual_position[n][0] = individual_position[n][0] - 1
                              individual position[n][0] = grid x
```

```
elif motion == 3:
                  if individual_position[n][1] < grid_y:</pre>
                       individual_position[n][1] = individual_position[n][1] + 1
                  else:
                       individual_position[n][0] = 0
              elif motion == 4:
                  if individual_position[n][1] > 0:
                       individual_position[n][1] = individual_position[n][1] - 1
                  else:
                       individual position[n][0] = grid y
     for i in range(no_of_individuals):
         if individual status[i] == 2:
              infected_individual_position = individual_position[i]
              rand2 = np.random.rand()
              if rand2 < beta:</pre>
                  for j in range(no_of_individuals):
                       if individual_position(j) == infected_individual_position and individual_status(j) == 1:
                           individual_status[j] = 2
              rand3 = np.random.rand()
              if rand3 < gamma:</pre>
                  individual status[i] = 3
     for i in range(no_of_individuals):
         if individual_status[i] == 1:
              final status[0][m] = final status[0][m] + 1
         if individual_status[i] == 2:
              final status[1][m] = final status[1][m] + 1
         if individual status[i] == 3:
              final_status[2][m] = final_status[2][m] + 1
sns.set_style("darkgrid")
fig = plt.figure(figsize=(20,10))
fig.tight layout()
plt.rcParams.update({'font.size': 14})
ax1 = fig.add_subplot(121)
ax1.clear()
# ax1.set_ylim(0, 10)
# ax1.set xlim(0, 10)
for i in range(no_of_individuals):
     if individual_status[i] == 1:
         \verb|plt.scatter| (individual_position[i][0], individual_position[i][1], color = "b")|
     elif individual_status[i] == 2:
         plt.scatter(individual_position[i][0],individual_position[i][1], color = "r")
     elif individual status[i] == 3:
         plt.scatter(individual_position[i][0],individual_position[i][1], color = "g")
ax1.set xlabel('x')
ax1.set_ylabel('y')
ax1.set title(f'Iteration = {no of iterations}, Number of agents = {no of individuals}')
ax1 = fig.add subplot(122)
ax1.plot(range(no_of_iterations), final_status[0][:], "b", label = "susceptible")
ax1.plot(range(no_of_iterations), final_status[1][:], "r", label = "infected")
ax1.plot(range(no_of_iterations), final_status[2][:], "g", label = "recovered")
ax1.set_xlabel('Iteration')
ax1.set_ylabel('Number of agents')
plt.legend(ncol = 1, loc = "center right")
ax1.set_title(f" beta ={beta}, gamma={gamma}, \n d={diffusion_rate}")
fig.show()
<ipython-input-4-5e10c0a5e74d>:108: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend i
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<ipython-input-4-bel0c0abe/4d>:108: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_i
nline, which is a non-GUI backend, so cannot show the figure.
 fig.show()

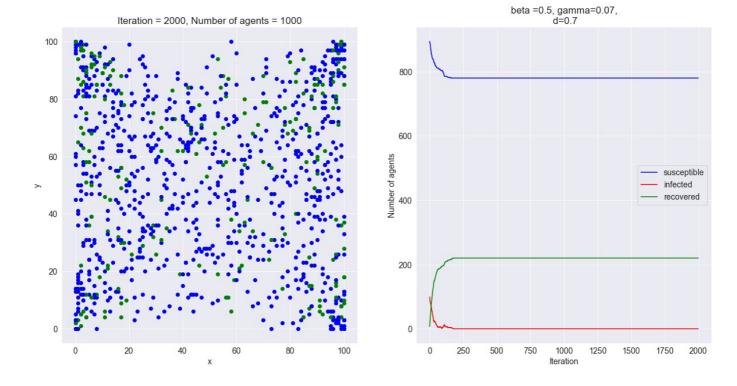




```
In [5]:
         no of individuals = 1000
         no_of_iterations = 2000
         diffusion rate = 0.7 # diffusion probability
         initial infection rate = 0.1
                      #Infection Probability
         beta = 0.5
         gamma = 0.07 # Recovery Probability
         grid x = 100 # max coordinate in x
         grid_y = 100 #max coordinate in y
         # status 1 = susceptible
         # status 2 = infected
         # status 3 = immune (recovered)
         individual position = []
         individual status = []
         final status = np.zeros((3, no of iterations))
         for i in range(no_of_individuals):
              individual_position.append([random.randint(0,grid_x), random.randint(0,grid_y)])
             individual status.append(1)
         initially infected individuals = random.sample(range(no of individuals), round(no of individuals*initial infection
         for j in initially infected individuals:
              individual_status[j] = \overline{2}
         for m in range(no_of_iterations):
              for n in range(no of individuals):
                  rand1 = np.random.rand()
                  if rand1 < diffusion_rate:</pre>
                      motion = np.random.choice([1, 2, 3, 4])
                      if motion == 1:
                          if individual_position[n][0] < grid_x:</pre>
                              individual_position[n][0] = individual_position[n][0] + 1
                              individual position[n][0] = 0
                      elif motion == 2:
                          if individual_position[n][0] > 0:
   individual_position[n][0] = individual_position[n][0] - 1
                              individual position[n][0] = grid x
                      elif motion == 3:
                          if individual_position[n][1] < grid_y:</pre>
                              individual position[n][1] = individual position[n][1] + 1
                          else:
                              individual_position[n][0] = 0
                      elif motion == 4:
                          if individual_position[n][1] > 0:
                              individual_position[n][1] = individual_position[n][1] - 1
                          else:
                              individual_position[n][0] = grid_y
             for i in range(no of individuals):
                  if individual status[i] == 2:
                      infected_individual_position = individual_position[i]
                      rand2 = np.random.rand()
                      if rand2 < beta:</pre>
                          for j in range(no_of_individuals):
                              if individual_position[j] == infected_individual_position and individual_status[j] == 1:
                                  individual status[j] = 2
                      rand3 = np.random.rand()
                      if rand3 < gamma:</pre>
                          individual status[i] = 3
             for i in range(no_of_individuals):
                  if individual status[i] == 1:
                      final_status[0][m] = final_status[0][m] + 1
```

```
if individual status[i] == 2:
              final_status[1][m] = final_status[1][m] + 1
         if individual_status[i] == 3:
              final status[2][m] = final status[2][m] + 1
sns.set style("darkgrid")
fig = plt.figure(figsize=(20,10))
fig.tight_layout()
plt.rcParams.update({'font.size': 14})
ax1 = fig.add_subplot(121)
ax1.clear()
# ax1.set ylim(0, 10)
# ax1.set_xlim(0, 10)
for i in range(no of individuals):
    if individual status[i] == 1:
         \verb|plt.scatter(individual_position[i][0], individual_position[i][1], color = "b"|)|
     elif individual_status[i] == 2:
         plt.scatter(individual_position[i][0],individual_position[i][1], color = "r")
     elif individual_status[i] == 3:
         plt.scatter(individual_position[i][0],individual_position[i][1], color = "g")
ax1.set_xlabel('x')
ax1.set ylabel('y')
ax1.set_title(f'Iteration = {no_of_iterations}, Number of agents = {no_of_individuals}')
ax1 = fig.add subplot(122)
ax1.plot(range(no_of_iterations), final_status[0][:], "b", label = "susceptible")
ax1.plot(range(no_of_iterations), final_status[1][:], "r", label = "infected")
ax1.plot(range(no_of_iterations), final_status[2][:], "g", label = "recovered")
ax1.set xlabel('Iteration')
ax1.set_ylabel('Number of agents')
plt.legend(ncol = 1, loc = "center right")
ax1.set title(f" beta ={beta}, gamma={gamma}, \n d={diffusion rate}")
fig.show()
```

<ipython-input-5-af2bfc2bd229>:108: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_i
nline, which is a non-GUI backend, so cannot show the figure.
fig.show()



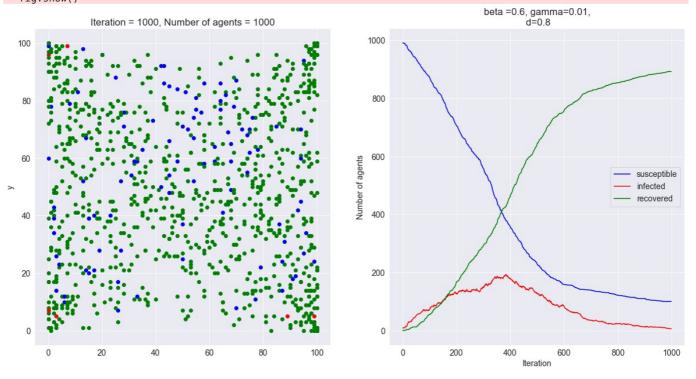
```
In [6]:
    no_of_individuals = 1000
    no_of_iterations = 1000
    diffusion_rate = 0.8  # diffusion probability
    initial_infection_rate = 0.01
    beta = 0.6  #Infection Probability
    gamma = 0.01  # Recovery Probability
    grid_x = 100  # max coordinate in x
    grid_y = 100  #max coordinate in y

# status 1 = susceptible
```

```
# status 2 = infected
# status 3 = immune (recovered)
individual_position = []
individual status = []
final_status = np.zeros((3, no_of_iterations))
for i in range(no_of_individuals):
    individual_position.append([random.randint(0,grid_x), random.randint(0,grid_y)])
    individual status.append(1)
initially_infected_individuals = random.sample(range(no_of_individuals), round(no_of_individuals*initial_infection)
for j in initially_infected_individuals:
    individual_status[j] = 2
for m in range(no_of_iterations):
    for n in range(no of individuals):
        rand1 = np.random.rand()
        if rand1 < diffusion_rate:</pre>
            motion = np.random.choice([1, 2, 3, 4])
            if motion == 1:
                if individual_position[n][0] < grid_x:</pre>
                    individual position[n][0] = individual position[n][0] + 1
                    individual_position[n][0] = 0
            elif motion == 2:
                if individual position[n][0] > 0:
                    individual_position[n][0] = individual_position[n][0] - 1
                     individual position[n][0] = grid x
            elif motion == 3:
                if individual position[n][1] < grid y:</pre>
                    individual_position[n][1] = individual_position[n][1] + 1
                 else:
                     individual position[n][0] = 0
            elif motion == 4:
                if individual_position[n][1] > 0:
                    individual position[n][1] = individual position[n][1] - 1
                else:
                     individual position[n][0] = grid y
    for i in range(no_of_individuals):
        if individual_status[i] == 2:
            infected individual position = individual position[i]
            rand2 = np.random.rand()
            if rand2 < beta:</pre>
                for j in range(no_of_individuals):
                    if individual_position[j] == infected_individual_position and individual_status[j] == 1:
                        individual_status[j] = 2
            rand3 = np.random.rand()
            if rand3 < gamma:</pre>
                individual status[i] = 3
    for i in range(no_of_individuals):
        if individual status[i] == 1:
            final status[0][m] = final status[0][m] + 1
        if individual_status[i] == 2:
            final status[1][m] = final status[1][m] + 1
        if individual_status[i] == 3:
            final_status[2][m] = final_status[2][m] + 1
sns.set_style("darkgrid")
fig = plt.figure(figsize=(20,10))
fig.tight layout()
plt.rcParams.update({'font.size': 14})
ax1 = fig.add subplot(121)
ax1.clear()
# ax1.set ylim(0, 10)
# ax1.set_xlim(0, 10)
for i in range(no_of_individuals):
    if individual_status[i] == 1:
        plt.scatter(individual_position[i][0],individual_position[i][1], color = "b")
    elif individual status[i] == 2:
        \verb|plt.scatter(individual_position[i][0], individual_position[i][1], | color = "r"|)|
    elif individual_status[i] == 3:
        plt.scatter(individual_position[i][0],individual_position[i][1], color = "g")
# ax1.set xlabel('x')
ax1.set vlabel('v')
ax1.set_title(f'Iteration = {no_of_iterations}, Number of agents = {no_of_individuals}')
ax1 = fig.add_subplot(122)
```

```
ax1.plot(range(no_of_iterations), final_status[0][:], "b", label = "susceptible")
ax1.plot(range(no_of_iterations), final_status[1][:], "r", label = "infected")
ax1.plot(range(no_of_iterations), final_status[2][:], "g", label = "recovered")
ax1.set_xlabel('Iteration')
ax1.set_ylabel('Number of agents')
plt.legend(ncol = 1, loc = "center right")
ax1.set_title(f" beta ={beta}, gamma={gamma}, \n d={diffusion_rate}")
fig.show()
```

<ipython-input-6-8f74b89cf45f>:108: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_i
nline, which is a non-GUI backend, so cannot show the figure.
fig.show()



Exercise 11.2 (part -a)

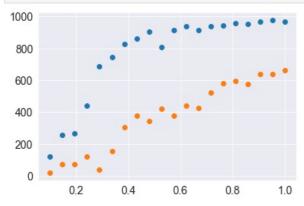
```
In [8]:
         \# for gamma = 0.01
         import numpy as np
         import random
         import matplotlib.pyplot as plt
         no of individuals = 1000
         no\_of\_iterations = 1000
         diffusion_rate = 0.8
                                 # diffusion probability
         initial infection rate = 0.01
         beta = \overline{np.linspace}(0.1, 1, 20)
                                            #Infection Probability
         gamma = 0.01 # Recovery Probability
grid_x = 100 # max coordinate in x
         grid_y = 100 #max coordinate in y
         # status 1 = susceptible
         # status 2 = infected
         # status 3 = immune (recovered)
         repetition = 3
         recovered_individuals_1 = []
         for b in beta:
              total_recovered = 0
              for rep in range(repetition):
                    print(rep)
                  individual_position = []
                  individual_status = []
                  for i in range(no of individuals):
                      individual\_position.append([random.randint(0,grid\_x), random.randint(0,grid\_y)])
                      individual status.append(1)
                  initially_infected_individuals = random.sample(range(no_of_individuals), round(no_of_individuals*initial)
                  for j in initially_infected_individuals:
                      individual_status[j] = 2
```

```
for m in range(no of iterations):
             for n in range(no_of_individuals):
                 rand1 = np.random.rand()
                 if rand1 < diffusion rate:</pre>
                     motion = np.random.choice([1, 2, 3, 4])
                     if motion == 1:
                         if individual position[n][0] < grid x:</pre>
                             individual position[n][0] = individual position[n][0] + 1
                         else:
                             individual position[n][0] = 0
                     elif motion == 2:
                         if individual_position[n][0] > 0:
                             individual position[n][0] = individual position[n][0] - 1
                             individual_position[n][0] = grid_x
                     elif motion == 3:
                         if individual_position[n][1] < grid_y:</pre>
                             individual_position[n][1] = individual_position[n][1] + 1
                             individual position[n][0] = 0
                     elif motion == 4:
                         if individual_position[n][1] > 0:
                             individual_position[n][1] = individual_position[n][1] - 1
                         else:
                             individual position[n][0] = grid y
             for i in range(no_of_individuals):
                 if individual status[i] == 2:
                     infected individual position = individual position[i]
                     rand2 = np.random.rand()
                     if rand2 < b:</pre>
                         for j in range(no of individuals):
                             if individual_position[j] == infected_individual_position and individual_status[j] ==
  individual_status[j] = 2
                     rand3 = np.random.rand()
                     if rand3 < gamma:</pre>
                         individual_status[i] = 3
        recovered individuals = 0
        for i in range(no of individuals):
             if individual_status[i] == 3:
                 recovered_individuals = recovered_individuals + 1
          print('recovered', recovered individuals)
#
        total_recovered = total_recovered + recovered_individuals
#
          print('total recovered', total recovered)
    recovered individuals 1.append(total recovered/repetition)
      print(recovered individuals 1)
# for gamma = 0.02
```

```
In [9]:
         no_of_individuals = 1000
         no of iterations = 1000
         diffusion rate = 0.8 # diffusion probability
         initial_infection_rate = 0.01
         beta = \overline{np.linspace}(0.1, 1, 20)
                                            #Infection Probability
         gamma = 0.02 # Recovery Probability
grid_x = 100 # max coordinate in x
         grid y = 100 #max coordinate in y
         final_status = np.zeros((3, no of iterations))
         # status 1 = susceptible
         # status 2 = infected
         # status 3 = immune (recovered)
         repetition = 3
         recovered individuals 2 = []
         for b in beta:
              total_recovered = 0
              for rep in range(repetition):
                    print(rep)
                  individual_position = []
                  individual_status = []
                  for i in range(no_of_individuals):
                      individual\_position.append([random.randint(0,grid\_x), random.randint(0,grid\_y)])
                      individual_status.append(1)
                  initially_infected_individuals = random.sample(range(no_of_individuals), round(no_of_individuals*initial_
                  for j in initially_infected_individuals:
                      individual status[j] = 2
                  for m in range(no_of_iterations):
                      for n in range(no of individuals):
                          rand1 = np.random.rand()
```

```
if rand1 < diffusion rate:</pre>
                    motion = np.random.choice([1, 2, 3, 4])
                     if motion == 1:
                         if individual position[n][0] < grid x:</pre>
                             individual position[n][0] = individual position[n][0] + 1
                             individual position[n][0] = 0
                    elif motion == 2:
                         if individual_position[n][0] > 0:
                             individual position[n][0] = individual position[n][0] - 1
                         else:
                             individual_position[n][0] = grid_x
                    elif motion == 3:
                         if individual_position[n][1] < grid_y:</pre>
                             individual position[n][1] = individual position[n][1] + 1
                             individual_position[n][0] = 0
                     elif motion == 4:
                         if individual_position[n][1] > 0:
                             individual_position[n][1] = individual_position[n][1] - 1
                             individual position[n][0] = grid y
            for i in range(no_of_individuals):
                if individual_status[i] == 2:
                    infected_individual_position = individual_position[i]
                     rand2 = np.random.rand()
                     if rand2 < b:</pre>
                         for j in range(no_of_individuals):
                             if individual_position[j] == infected_individual_position and individual_status[j] ==
                                 individual_status[j] = 2
                     rand3 = np.random.rand()
                    if rand3 < gamma:</pre>
                        individual_status[i] = 3
        recovered_individuals = 0
        for i in range(no_of_individuals):
            if individual_status[i] == 3:
                recovered individuals = recovered individuals + 1
          print('recovered', recovered individuals)
#
        total recovered = total recovered + recovered individuals
#
          print('total recovered', total recovered)
    recovered_individuals_2.append(total_recovered/repetition)
      print(recovered_individuals_1)
plt.scatter(beta, recovered_individuals_1)
plt.scatter(beta, recovered_individuals_2)
plt.show()
```

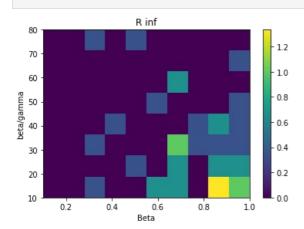
In [10]:



Exercise 11.2 (part -b)

```
In [3]:
         import numpy as np
         import random
         import matplotlib.pyplot as plt
         no_of_individuals = 1000
         no of iterations = 1000
         diffusion rate = 0.8 # diffusion probability
         initial_infection_rate = 0.01
beta = np.linspace(0.1, 1, 10)
                                              #Infection Probability
         # gamma = 0.01 # Recovery Probability
         grid_x = 100  # max coordinate in x
```

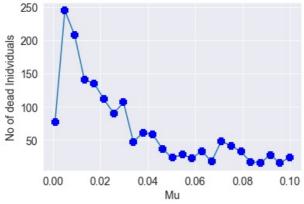
```
grid\ y = 100 \ \#max\ coordinate\ in\ y
repetition = 3
# status 1 = susceptible
# status 2 = infected
# status 3 = immune (recovered)
# final status = np.zeros((3, no of iterations))
BG = [80,70,60,50,40,30,20,10]
beta_list = []
for bg in range(len(BG)):
    bglist = []
    for b in beta:
        total recovered = 0
        for rep in range(repetition):
            gamma = 1/(BG[bg]/b)
            individual_position = []
            individual status = []
            final_status = np.zeros((3, no_of_iterations))
            for i in range(no_of individuals):
                individual_position.append([random.randint(0,grid_x), random.randint(0,grid_y)])
                individual status.append(1)
            initially infected individuals = random.sample(range(no of individuals), round(no of individuals*init
            for j in initially_infected_individuals:
                individual_status[j] = 2
            for m in range(no of iterations):
                for n in range(no_of_individuals):
                     rand1 = np.random.rand()
                     if rand1 < diffusion rate:</pre>
                         motion = np.random.choice([1, 2, 3, 4])
                         if motion == 1:
                             if individual position[n][0] < grid x:</pre>
                                 individual_position[n][0] = individual_position[n][0] + 1
                         else:
                             individual position[n][0] = 0
                     elif motion == 2:
                         if individual_position[n][0] > 0:
                             individual position[n][0] = individual position[n][0] - 1
                         else:
                             individual position[n][0] = grid x
                     elif motion == 3:
                         if individual_position[n][1] < grid_y:</pre>
                             individual_position[n][1] = individual_position[n][1] + 1
                             individual position[n][0] = 0
                     elif motion == 4:
                         if individual position[n][1] > 0:
                             individual position[n][1] = individual position[n][1] - 1
                         else:
                             individual position[n][0] = grid y
            for i in range(no_of_individuals):
                if individual_status[i] == 2:
                    infected individual position = individual position[i]
                     rand2 = np.random.rand()
                     if rand2 < b:</pre>
                         for j in range(no of individuals):
                             if individual_position[j] == infected_individual_position and individual_status[j] ==
   individual_status[j] = 2
                     rand3 = np.random.rand()
                     if rand3 < gamma:</pre>
                         individual_status[i] = 3
            for i in range(no_of_individuals):
                if individual_status[i] == 1:
                     final_status[0][m] = final_status[0][m] + 1
                if individual status[i] == 2:
                     final status[1][m] = final status[1][m] + 1
                if individual_status[i] == 3:
                    final_status[2][m] = final_status[2][m] + 1
            total recovered = final status[2][no of iterations-1]
        bglist.append(total_recovered/repetition)
    beta list.append(bglist)
plt.imshow(beta_list,aspect='auto', extent = [min(beta), max(beta), min(BG), max(BG)])
plt.xlabel("Beta")
plt.ylabel("beta/gamma")
plt.title("R inf")
plt.colorbar()
plt.show()
```



Exercise 11.3

```
In [13]:
                       import numpy as np
                       import random
                       import matplotlib.pyplot as plt
                       no_of_individuals = 1000
                       no_of_iterations = 1000
                       diffusion_rate = 0.8 # diffusion probability
                       initial_infection_rate = 0.01
                       beta = 0.6 #Infection Probability
                       gamma = 0.01 # Recovery Probability
                       grid_x = 100  # max coordinate in x
grid_y = 100  #max coordinate in y
                       Mu = np.linspace(0.001, 0.1, 25)
                       # status 1 = susceptible
                       # status 2 = infected
                       # status 3 = immune (recovered)
                       repetition = 3
                       no_of_dead = []
                       for d in range(len(Mu)):
                                total_dead = 0
                                for rep in range(repetition):
                                         mu = Mu[d]
                                             print('mu', mu)
print('rep', rep)
                       #
                       #
                                         individual_position = []
                                         individual_status = []
                       #
                                             StatusList = np.zeros((3,numberOfLoops))
                                         for i in range(no_of_individuals):
                                                   individual position.append([random.randint(0,grid x), random.randint(0,grid y)])
                                                  individual status.append(1)
                                         initially\_infected\_individuals = random.sample(range(no\_of\_individuals), \ round(no\_of\_individuals*initial\_initial), \ round(no\_of\_individuals*initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial\_initial
                                         for j in initially infected individuals:
                                                   individual_status[j] = 2
                                         for m in range(no of iterations):
                                                  for n in range(no_of_individuals):
                                                           rand1 = np.random.rand()
                                                           if rand1 < diffusion_rate:</pre>
                                                                    motion = np.random.choice([1, 2, 3, 4])
                                                                    if motion == 1:
                                                                              if individual position[n][0] < grid x:</pre>
                                                                                       individual_position[n][0] = individual_position[n][0] + 1
                                                                              else:
                                                                                       individual position[n][0] = 0
                                                                     elif motion == 2:
                                                                              if individual_position[n][0] > 0:
                                                                                       individual_position[n][0] = individual_position[n][0] - 1
                                                                                       individual position[n][0] = grid x
                                                                    elif motion == 3:
                                                                              if individual_position[n][1] < grid_y:</pre>
                                                                                       individual position[n][1] = individual position[n][1] + 1
                                                                                       individual_position[n][0] = 0
                                                                    elif motion == 4:
```

```
if individual position[n][1] > 0:
                             individual_position[n][1] = individual_position[n][1] - 1
                         else:
                             individual position[n][0] = grid y
            for i in range(no_of_individuals):
                if individual_status[i] == 2:
                    infected individual position = individual position[i]
                     rand2 = np.random.rand()
                     if rand2 < beta:</pre>
                         for j in range(no_of_individuals):
                             if individual_position[j] == infected_individual_position and individual_status[j] ==
                                 individual_status[j] = 2
                     rand3 = np.random.rand()
                     if rand3 < gamma:</pre>
                         individual_status[i] = 3
                     rand4 = np.random.rand()
                     if rand4 < mu:</pre>
                        individual_status[i] = 4
        dead individuals = 0
        for i in range(no_of_individuals):
            if individual_status[i] == 4:
                dead individuals = dead individuals + 1
#
          print('dead', dead_individuals)
        total_dead = total_dead + dead_individuals
#
          print('total dead', total dead)
    no_of_dead.append(total_dead/repetition)
#
      print(no_of_dead)
plt.plot(Mu,no_of_dead,marker = "+",markeredgecolor = 'blue',markeredgewidth = 10)
plt.xlabel('Mu')
plt.ylabel('No of dead Inidviduals')
plt.show()
```

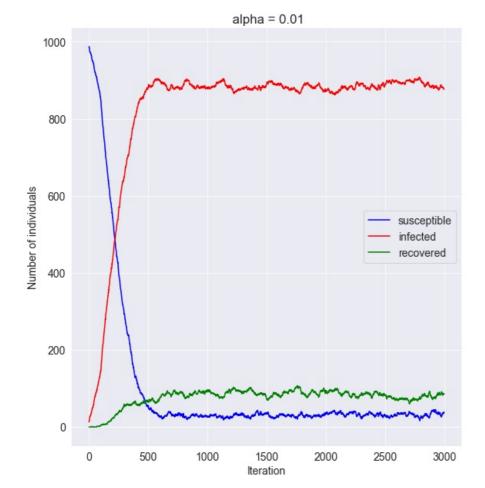


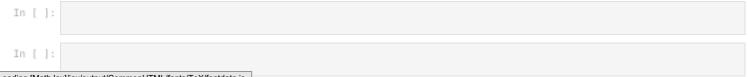
Exercise 11.4

```
In [14]:
          import numpy as np
          import matplotlib.pyplot as plt
          import pylab
          import seaborn as sns
          import random
          import math
          no_of_individuals = 1000
          no_of_iterations = 3000
          diffusion rate = 0.8 # diffusion probability
          initial infection rate = 0.01
          beta = \overline{0.4} #Infection Probability
          gamma = 0.001 # Recovery Probability
          grid x = 100 \# max coordinate in x
          grid_y = 100 #max coordinate in y
          alpha = 0.01
          # status 1 = susceptible
          # status 2 = infected
          # status 3 = immune (recovered)
          individual_position = []
          individual status = []
          final_status = np.zeros((3, no_of_iterations))
          for i in range(no of individuals):
```

```
individual position.append([random.randint(0,grid x), random.randint(0,grid y)])
     individual_status.append(1)
initially infected individuals = random.sample(range(no of individuals), round(no of individuals*initial infection
for j in initially_infected individuals:
     individual_status[j] = 2
for m in range(no of iterations):
     for n in range(no_of_individuals):
         rand1 = np.random.rand()
         if rand1 < diffusion rate:</pre>
              motion = np.random.choice([1, 2, 3, 4])
              if motion == 1:
                  if individual position[n][0] < grid x:</pre>
                       individual position[n][0] = individual position[n][0] + 1
                  else:
                       individual_position[n][0] = 0
              elif motion == 2:
                  if individual_position[n][0] > 0:
                       individual position[n][0] = individual position[n][0] - 1
                  else:
                       individual_position[n][0] = grid_x
              elif motion == 3:
                  if individual_position[n][1] < grid_y:</pre>
                       individual_position[n][1] = individual_position[n][1] + 1
                       individual position[n][0] = 0
              elif motion == 4:
                  if individual_position[n][1] > 0:
                       individual_position[n][1] = individual_position[n][1] - 1
                  else:
                       individual position[n][0] = grid y
     for i in range(no_of_individuals):
         if individual_status[i] == 2:
              infected individual position = individual position[i]
              rand2 = np.random.rand()
              if rand2 < beta:</pre>
                  for j in range(no of individuals):
                       if individual_position[j] == infected_individual_position and individual_status[j] == 1:
                           individual status[j] = 2
              rand3 = np.random.rand()
              if rand3 < gamma:</pre>
                  individual_status[i] = 3
         if individual status[i] == 3:
              rand4 = np.random.rand()
if rand4 < alpha:</pre>
                  individual_status[i] = 1
     for i in range(no of individuals):
          if individual status[i] == 1:
              final status[0][m] = final status[0][m] + 1
         if individual_status[i] == 2:
              final_status[1][m] = final_status[1][m] + 1
         if individual status[i] == 3:
              final_status[2][m] = final_status[2][m] + 1
# sns.set style("darkgrid")
fig = plt.figure(figsize=(20,10))
fig.tight layout()
plt.rcParams.update({'font.size': 14})
ax1 = fig.add_subplot(121)
ax1.clear()
ax1.plot(range(no_of_iterations), final_status[0][:], "b", label = "susceptible")
ax1.plot(range(no_of_iterations), final_status[1][:], "r", label = "infected")
ax1.plot(range(no_of_iterations), final_status[2][:], "g", label = "recovered")
ax1.set_xlabel('Iteration')
ax1.set_ylabel('Number of individuals')
plt.legend(ncol = 1, loc = "center right")
ax1.set\_title(f" beta = \{beta\}, gamma=\{gamma\}, \\ \  \  d=\{diffusion\_rate\}, \\ \  \  \  \  n \ alpha = \{alpha\}")
fig.show()
<ipython-input-14-ed4290cf6112>:106: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_
```

<ipython-input-14-ed4290cf6112>:106: UserWarning: Matplotlib is currently using module://ipykernel.pylab.backend_
inline, which is a non-GUI backend, so cannot show the figure.
 fig.show()





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