```
1 import numpy as np
2 import matplotlib.pyplot as plt
 3
  def printPrettyMatrix(objStr, matrix):
4
5
       objStr+="\t\t["
       for row in matrix:
6
7
           objStr+="["
8
           for col in row:
                objStr+=f" {col:.02f},"
9
           objStr= objStr[:-1]+"],"
10
11
           objStr+="\n \t\t"
       objStr=objStr[:-5]+"]"
12
13
       return objStr
14
15 def showPlot(PlotSS, PlotII, PlotRR, node, ts, main_title):
       plt.title(f"{main_title}\nNode {node} Infctd:{np.max(PlotII)}")
16
       S i = plt.plot(PlotSS[node,:], label=f'S_{node}({ts})') # just plot node 0's
17
   susceptible over time
       I_i = plt.plot(PlotII[node,:], label=f'I_{node}({ts})') # just plot node 0's
18
   infection over time
       R_i = plt.plot(PlotRR[node,:], label=f'R_{node}({ts})') # just plot node 0's
19
   recovery over time
       plt.legend()
20
       plt.show()
21
22
23 def SIR(beta, gamma, A, I0, timesteps, show_plot=False, node=1, main_title="Question
   00"):
       . . .
24
25
       Notes:
26
         - assumes that everything is dimensioned correctly
27
         - all S/I/R are fractions of node's population
         - R0 = 0
28
         - S0 = 1-I0
29
30
31
       n = A.shape[0]
32
33
       # initialize:
       SS = np.zeros([n,timesteps]) # each column in a list of S i's; there is one
34
   column per timestep
35
       II = np.zeros_like(SS) # copy the shape of SS to create II
       RR = np.zeros like(SS) # nobody starts out with immunity
36
37
       II[:,0] = I0 # set the initial infection
       SS[:,0] = np.ones(n)-II[:,0] # if you're not infected, you're susceptible
38
39
40
       for t in range(1,timesteps):
41
           bigS = np.diag(SS[:,t-1])
42
           new_infections = beta * bigS @ A @ II[:,t-1]
43
           heals = gamma*II[:,t-1]
44
45
           SS[:,t] = SS[:,t-1] - new_infections
46
           II[:,t] = II[:,t-1] + new_infections - heals
47
           RR[:,t] = RR[:,t-1] + heals
48
49
       if show plot:
50
           showPlot(SS, II, RR, node, timesteps, main_title)
51
52
       return SS,II,RR
53
```

```
55 # Homework 05: Question 1
57 print("\nQuestion 1:")
58 beta = 0.8
59 \mid gamma = 0.4
60 Q1_og_infection = [0.01, 0.0, 0.0, 0.0, 0.0]
61 Q1_graph_adj = np.array([[0.8, 0.0, 0.0, 0.05, 0.15],
                         [1.0, 0.0, 0.0, 0.0, 0.0],
62
                         [0.3, 0.2, 0.5, 0.0, 0.0],
63
64
                         [0.0, 0.0, 0.05, 0.95, 0.0],
                         [0.0, 0.0, 0.0, 0.2, 0.8]])
65
66 | q1_{timesteps} = 2
67
68 Q1_SS, Q1_II, Q1_RR = SIR(beta, gamma, Q1_graph_adj, Q1_og_infection, q1_timesteps,
   show_plot=True, main_title="Question 01")
69 print("\tS_i(t):\t")
70 print(printPrettyMatrix("", Q1_SS))
71 print("\tI_i(t):\t")
72 print(printPrettyMatrix("", Q1_II))
73 print("\tR i(t):\t")
74 print(printPrettyMatrix("", Q1_RR))
75
77 # Homework 05: Question 2
79 print("\nQuestion 2:")
80 Q2_og_infection = [0.01, 0.0, 0.0, 0.0, 0.0]
81 Q2_graph_adj = np.array([[0.8, 0.0, 0.0, 0.05, 0.15],
82
                         [1.0, 0.0, 0.0, 0.0, 0.0],
83
                         [0.3, 0.2, 0.5, 0.0, 0.0],
84
                         [0.0, 0.0, 0.05, 0.95, 0.0],
85
                         [0.0, 0.0, 0.0, 0.2, 0.8]]
86 q2_timesteps = 0
87
88 while True:
89
      Q2_SS, Q2_II, Q2_RR = SIR(beta, gamma, Q2_graph_adj, Q2_og_infection,
   q2_timesteps+1, show_plot=False)
90
      if np.all(Q2_II[:, q2_timesteps] != 0.0):
91
          print(f"all nodes are infected at time step {q2_timesteps} \t{Q2_II[:,
   q2 timesteps]}")
92
          break
93
      if q2_timesteps % 2 == 0:
94
          showPlot(Q2_SS, Q2_II, Q2_RR, 2, q2_timesteps, main_title="Question 02")
95
96
97
      q2_{timesteps} += 1
98
99 print("\tS_i(t):\t")
100 print(printPrettyMatrix("", Q2_SS))
101 print("\tI_i(t):\t")
102 print(printPrettyMatrix("", Q2_II))
103 print("\tR_i(t):\t")
104 print(printPrettyMatrix("", Q2_RR))
105
106
108 # Homework 05: Question 3
110 print("\nQuestion 3:")
111 print("\tSuppose that the connection between node 2 and node 1 is removed. If\n",
```

```
"\tnothing else changes in the network, would this change the spread of\n",
112
113
         "\tthe epidemic? Can you predict how many people in node 2 would eventually\n",
         "\tget sick?\n")
114
115 Q3_og_infection = [0.01, 0.0, 0.0, 0.0, 0.0]
116 Q3_graph_adj = np.array([[0.8, 0.0, 0.0, 0.05, 0.15],
                            [0.0, 0.0, 0.0, 0.0, 0.0],
117
118
                            [0.3, 0.2, 0.5, 0.0, 0.0],
119
                            [0.0, 0.0, 0.05, 0.95, 0.0],
                            [0.0, 0.0, 0.0, 0.2, 0.8]])
120
121 q3_{timesteps} = 0
122
123 while True:
124
       Q3_SS, Q3_II, Q3_RR = SIR(beta, gamma, Q3_graph_adj, Q3_og_infection,
   q3 timesteps+1, node=2, show plot=False)
       if (Q3_II[:, q3_timesteps][1] != 0.0):
125
126
           print(f"Node 2 is infected at time step {q3_timesteps} \t{Q3_II[:,
   q3_timesteps]}")
127
           break
128
       if q3_timesteps % 5 == 0:
129
           print(f"Step:\t{q3_timesteps}")
           print(f"Node 2 infection %:{Q3_II[:, q3_timesteps][1]}")
130
           print(F"The rest of the network {Q3_II[:, q3_timesteps]}")
131
132
           showPlot(Q3_SS, Q3_II, Q3_RR, 2, q3_timesteps, "Question 03")
133
       if q3_timesteps > 50:
134
           break
135
       q3 timesteps += 1
136
137 print(f"Last ten time steps of {q3_timesteps-1}")
138 print("\tS i(t):\t")
139 print(printPrettyMatrix("",Q3_SS[:, q3_timesteps-10:q3_timesteps]))
140 print("\tI_i(t):\t")
141 print(printPrettyMatrix("", Q3_II[:, q3_timesteps-10:q3_timesteps]))
142 print("\tR_i(t):\t")
143 print(printPrettyMatrix("", Q3_RR[:, q3_timesteps-10:q3_timesteps]))
144
145
147 # Homework 05: Question 4
149 print("\nQuestion 4:")
150 print("\tRepeat that same thought experiment, but this time let the initial\n",
         "\tinfection start on node 2 (so S_i(0)=0.1 and S_2(0)=0.99, but all other\n",
151
         "\tnodes have S_i(0)=1, I_i(0)=0 and R_i(0)=0. Now can you predict how
152
   many\n"
         "\tpeople in node 2 would eventually get sick?\n")
153
154 Q4_og_infection = [0.0, 0.01, 0.0, 0.0, 0.0]
155 Q4_graph_adj = np.array([[0.8, 0.0, 0.0, 0.05, 0.15],
156
                            [0.0, 0.0, 0.0, 0.0, 0.0],
                            [0.3, 0.2, 0.5, 0.0, 0.0],
157
158
                            [0.0, 0.0, 0.05, 0.95, 0.0],
159
                            [0.0, 0.0, 0.0, 0.2, 0.8]]
160 | q4\_timesteps = 0
161
162 while True:
       Q4_SS, Q4_II, Q4_RR = SIR(beta, gamma, Q4_graph_adj, Q4_og_infection,
163
   q4_timesteps+1, node = 2, show_plot=False, main_title="Question 04")
164
       if q4_timesteps % 10 == 0:
           print(f"Step:\t{q4_timesteps}")
165
166
           print(f"Node 2 infection %:{Q4_II[:, q4_timesteps][1]}")
167
           print(F"The rest of the network {Q4_II[:, q4_timesteps]}")
```

```
showPlot(Q4_SS, Q4_II, Q4_RR, 2, q4_timesteps, main_title="Question 04")
168
169
       if q4_timesteps > 50:
170
           break
171
       q4 timesteps += 1
172
173 print(f"Last ten time steps of {q4_timesteps-1}")
174 print("\tS_i(t):\t")
175 print(printPrettyMatrix("",Q4_SS[:, q4_timesteps-10:q4_timesteps]))
176 print("\tI i(t):\t")
177 print(printPrettyMatrix("", Q4_II[:, q4_timesteps-10:q4_timesteps]))
178 print("\tR_i(t):\t")
179 print(printPrettyMatrix("", Q4_RR[:, q4_timesteps-10:q4_timesteps]))
180
181
183 # Homework 05: Question 5
185 print("\nQuestion 5:")
186 print("\tInstead, suppose you remove the connection from node 2 to node 1\n",
         "\tand replace it with a self-loop on 2 with weight 1. If nothing else\n",
187
188
         "\tchanges in the network (and the infection starts at node 1), would \n"
189
         "\tthis change the spread of the epidemic? Can you predict how many\n",
190
         "\tpeople in node 2 would eventually get sick?\n")
191 Q5_og_infection = [0.0, 0.01, 0.0, 0.0, 0.0]
192 Q5_graph_adj = np.array([[0.8, 0.0, 0.0, 0.05, 0.15],
                           [0.0, 1.0, 0.0, 0.0, 0.0],
193
194
                           [0.3, 0.2, 0.5, 0.0, 0.0],
195
                           [0.0, 0.0, 0.05, 0.95, 0.0],
196
                           [0.0, 0.0, 0.0, 0.2, 0.8]])
197
198 q5 timesteps = 0
199
200 while True:
       Q5_SS, Q5_II, Q5_RR = SIR(beta, gamma, Q5_graph_adj, Q5_og_infection,
201
   q5 timesteps+1, node = 2, show plot=False, main title="Question 05")
       if q5 timesteps % 10 == 0:
202
203
           print(f"Step:\t{q5_timesteps}")
           print(f"Node 2 infection %:{Q5_II[:, q5_timesteps][1]}")
204
           print(F"The rest of the network {Q5 II[:, q5 timesteps]}")
205
           showPlot(Q5_SS, Q5_II, Q5_RR, 2, q5_timesteps, "Question 05")
206
207
208
       if q5_timesteps > 50:
209
           break
210
       q5_timesteps += 1
211
212 print(f"Last ten time steps of {q5_timesteps-1}")
213 print("\tS_i(t):\t")
214 print(printPrettyMatrix("",Q5_SS[:, q5_timesteps-10:q5_timesteps]))
215 print("\tI i(t):\t")
216 print(printPrettyMatrix("", Q5_II[:, q5_timesteps-10:q5_timesteps]))
217 print("\tR i(t):\t")
218 print(printPrettyMatrix("", Q5_RR[:, q5_timesteps-10:q5_timesteps]))
```