```
In [6]:
         import numpy as np
         import random
         import matplotlib.pyplot as plt
         from matplotlib.colors import ListedColormap
         N = 10
         M = np.zeros((N,N))
         k = 16
         def simplify_sequence(sequence):
             result = []
             i = 0
             while i < len(sequence):</pre>
                                                             # if the current node is repeated
                 if sequence[i] in sequence[i+1:]:
                     i = sequence.index(sequence[i], i+1) # find the next index of the repeated node
                 else:
                     result.append(sequence[i])
                                                            # if the current node is not repeated, append it to the simplif
                     i += 1
             return result
         for i in range(N):
             for j in range(i+1, N):
                 if random.uniform(0,1) < 0.5:
                     M[i,j] = 1
                     M[j,i] = 1
                 else:
                     M[i,j] = 0
                     M[j,i] = 0
         plt.subplot(1,3,1)
         colors = ['White', 'Black']
         colormap = ListedColormap(colors)
         plt.title("Connection Matrix")
         plt.imshow(M,colormap)
         D = np.zeros((N,N))
         W = np.zeros((N,N))
         for i in range(N):
             for j in range(i+1, N):
                 if random.uniform(0,1) < 0.5:
                     if M[i,j] == 1:
                         r = random.uniform(1,100)
                         D[i,j] = r
                         D[j,i] = r
                     else:
                         D[i,j] = 100000000
D[j,i] = 100000000
                     W[i,j] = 1/D[i,j]
                     W[j,i] = 1/D[j,i]
         plt.subplot(1,3,2)
         plt.title("Distance Matrix")
         plt.imshow(D,vmin = 0,vmax = 100, cmap = 'Blues')
         plt.subplot(1,3,3)
         plt.title("Weight Matrix")
         plt.imshow(D, cmap = 'Reds')
         path = []
         start_vertex = random.randint(0,N)
         path.append(start vertex)
         for i in range (k-1):
             consecutive_vertex = path[i]
             consecutive vertex list = M[:, consecutive vertex]
             index = np.where(consecutive vertex list == 1)
             path.append(round(random.choice(random.choice(index))))
         distance = 0
         for i in range(len(path)-1):
             distance += D[path[i], path[i+1]]
         print('Path is ' + str(path))
         print('Distance travelled = ' + str(distance))
         paths = simplify_sequence(path)
         simplified distance = 0
         for i in range(len(paths)-1):
             simplified_distance += D[paths[i], paths[i+1]]
```

```
In [2]:
        import numpy as np
        from scipy.spatial import Delaunay
        import matplotlib.pyplot as plt
        def total distance(sequence,D):
            return sum(D[sequence[i], sequence[i+1]] for i in range(len(sequence)-1))
        #remove all the elements between repeated nodes
        def simplify_sequence(sequence):
           result = []
           i = 0
           while i < len(sequence):</pre>
               if sequence[i] in sequence[i+1:]:
                                                      # if the current node is repeated
                   i = sequence.index(sequence[i], i+1) # find the next index of the repeated node
                   result.append(sequence[i])
                                                     # if the current node is not repeated, append it to the simplif
                   i += 1
            return result
        def branch decision(i,T,D,M,alpha,beta):
            possible_next = np.where(M[i] == 1)[0]
                                                            # can only move to the nodes that are connected to the o
           probability next = np.zeros(len(possible next))
                                                            # probability of moving to each of the possible next noc
           for j in range(len(possible_next)):
               probability_next = probability_next / sum(probability_next)
                                                                                               # normalize the pr
            return np.random.choice(possible_next, p=probability_next)
```

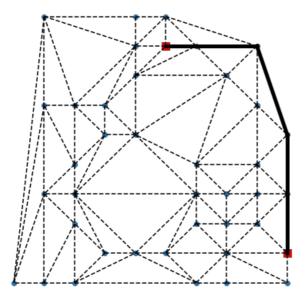
```
In [3]:
         n = 20
                           # number of ants
         alpha = 0.8
                           # alpha parameter
         beta = 1
                          # beta parameter
         rho = 0.5
                         # rho parameter
         max steps = 80  # maximum number of steps
         number of iterations = 100 # number of iterations
         N = 40 # number of nodes
         # generate N points at unique random positions (x,y) with x,y in [0,N]
                                     # Generate an array of all the possible points in grid
         all points = []
         for i in range(10):
             for j in range(10):
                all points.append([i,j])
         # generate N random unique points from all points
         indices = np.random.choice(len(all_points), N, replace=False)
         r = np.array([all_points[i] for i in indices])
         # connect the generated points by a Delaunay triangulation
         tri = Delaunay(r)
         # create the corresponding adjacency matrix
         M = np.zeros((N, N))
         for simplex in tri.simplices:
             for i in range(len(simplex)):
                 for j in range(i+1, len(simplex)):
                     M[simplex[i], simplex[j]] = 1
                     M[simplex[j], simplex[i]] = 1
```

```
# define the Euclidean distance all points in r
D = np.zeros((N, N))
for i in range(N):
    for j in range(i+1,N):
         D[i,j] = np.linalg.norm(r[i]-r[j])
         D[j,i] = D[i,j]
# make the distance inf for points that are not connected
for i in range(N):
    for j in range(i+1,N):
         if M[i,j] == 0:
             D[i,j] = np.inf
              D[j,i] = np.inf
T = M # initialize the Pheromone matrix, initially equal to the adjacency matrix
distance bird = 0
while distance_bird < 7:</pre>
                                           # we don't want the starting and the target node to be too close
     # decide a starting node s
    s = np.random.randint(0, N)
    # decide a target node t
    t = np.random.randint(0, N)
    distance bird = np.sqrt((r[t,0] - r[s,0])**2 + (r[t,1] - r[s,1])**2)
shortestDistance = []
shortestPath = []
for iteration in range(number_of_iterations):
     # initialize the paths of the ants
    T = T*(1-rho)
    paths = []
    for i in range(n):
         paths.append([s])
         for step in range(max_steps):
              next node = branch decision(paths[i][-1], T, D, M, alpha, beta)
              paths[i].append(next node)
              if next node == t:
                  break
    # check if the target node is in the path of any of the ants
    shortestDist = np.inf
    for i in range(n)
         if t in paths[i]:
              paths[i] = simplify_sequence(paths[i])
              path_distance = total_distance(paths[i],D)
              if path distance<shortestDist:</pre>
                  shortestDist = path_distance
                  shortPath = paths[i]
              for j in range(len(paths[i])-1):
                  T[paths[i][j], paths[i][j+1]] += 1/path_distance
                  T[paths[i][j+1],paths[i][j]] += 1/path_distance
     shortestDistance.append(shortestDist)
    shortestPath.append(shortPath)
    if iteration % 40 == 0:
         plt.figure(figsize=(7,7))
         ptt.ligdic(!ig3ize=(/,//)
plt.plot(r[:,0], r[:,1], 'o')
plt.plot(r[s,0], r[s,1], 's', color='red', markersize=10)
plt.plot(r[t,0], r[t,1], 's', color='red', markersize=10)
         for i in range(N):
              for j in range(i+1,N):
                  if M[i,j] == 1:
                       plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'k--')
         p = shortestPath[iteration]
         for q in range(len(p)-1):
             i = p[q]
              j = p[q+1]
              plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'black', linewidth=5)
         plt.title('Iteration ' + str(iteration+1)+' - Shortest Path')
         plt.axis('off')
         plt.show()
    if iteration % 40 == 0:
         plt.figure(figsize=(7,7))
         plt.plot(r[:,0], r[:,1], 'o')
plt.plot(r[s,0], r[s,1], 's', color='red', markersize=10)
plt.plot(r[t,0], r[t,1], 's', color='red', markersize=10)
         for i in range(N):
              for j in range(i+1,N):
                  if M[i,j] == 1:
                       plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'k--')
         for i in range(N):
              for j in range(i+1,N):
         \label{eq:plt.plot} $$ plt.plot([r[i,0],\ r[j,0]],\ [r[i,1],\ r[j,1]],'r',linewidth=T[i,j]*2) $$ plt.title('Iteration' + str(iteration+1) + ' - Pheromone Matrix') $$
```

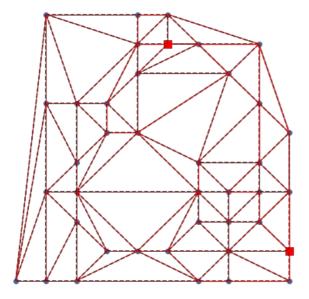
```
plt.axis('off')
    plt.show()

plt.plot(np.linspace(0,number_of_iterations-1,number_of_iterations),shortestDistance)
plt.xlabel("No. of Iterations")
plt.ylabel("Shortest Path Distance")
```

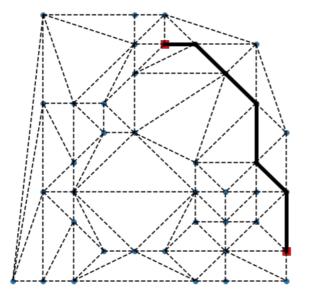
Iteration 1 - Shortest Path



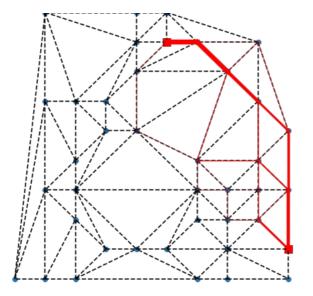
Iteration 1 - Pheromone Matrix



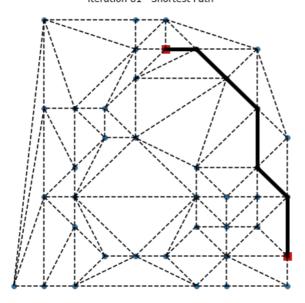
Iteration 41 - Shortest Path



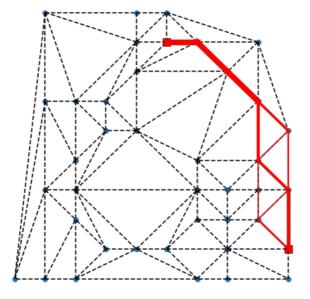
Iteration 41 - Pheromone Matrix



Iteration 81 - Shortest Path



Iteration 81 - Pheromone Matrix



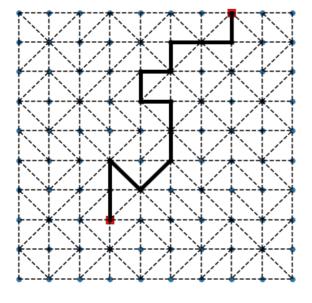
Out[3]: Text(0, 0.5, 'Shortest Path Distance')



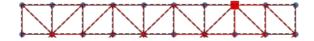
```
In [4]:
         n = 20
                            # number of ants
         alpha = 0.8
                            # alpha parameter
         beta = 1
                          # beta parameter
         rho = 0.5
                          # rho parameter
         max_steps = 80
                         # maximum number of steps
         number of iterations = 100 # number of iterations
         N = 100 # number of nodes
         # generate N points at unique random positions (x,y) with x,y in [0,N]
                                     # Generate an array of all the possible points in grid
         all_points = []
         for i in range(10):
             for j in range(10):
                 all_points.append([i,j])
         # generate N random unique points from all_points
         indices = np.random.choice(len(all_points), N, replace=False)
         r = np.array([all_points[i] for i in indices])
         # connect the generated points by a Delaunay triangulation
         tri = Delaunay(r)
         # create the corresponding adjacency matrix
         M = np.zeros((N, N))
         for simplex in tri.simplices:
             for i in range(len(simplex)):
                 for j in range(i+1, len(simplex)):
                     M[simplex[i], simplex[j]] = 1
                     M[simplex[j], simplex[i]] = 1
         # define the Euclidean distance all points in r
         D = np.zeros((N, N))
         for i in range(N):
             for j in range(i+1,N):
                 D[i,j] = np.linalg.norm(r[i]-r[j])
                 D[j,i] = D[i,j]
         # make the distance inf for points that are not connected
         for i in range(N):
             for j in range(i+1,N):
                 if M[i,j] == 0:
                     D[i,j] = np.inf
                     D[j,i] = np.inf
         T = M # initialize the Pheromone matrix, initially equal to the adjacency matrix
         distance bird = 0
         while distance_bird < 7:</pre>
                                               # we don't want the starting and the target node to be too close
             # decide a starting node s
             s = np.random.randint(0, N)
             # decide a target node t
             t = np.random.randint(0, N)
             distance_bird = np.sqrt((r[t,0] - r[s,0])**2 + (r[t,1] - r[s,1])**2)
         shortestDistance = []
         shortestPath = []
         for iteration in range(number of iterations):
             # initialize the paths of the ants
             T = T*(1-rho)
             paths = []
             for i in range(n):
                 paths.append([s])
                 for step in range(max_steps):
                     next_node = branch_decision(paths[i][-1], T, D, M, alpha, beta)
                     paths[i].append(next_node)
                     if next_node == t:
                         break
```

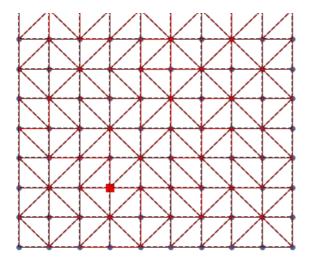
```
# check if the target node is in the path of any of the ants
     shortestDist = np.inf
     for i in range(n):
         if t in paths[i]:
              paths[i] = simplify_sequence(paths[i])
              path_distance = total_distance(paths[i],D)
              if path_distance<shortestDist:</pre>
                   shortestDist = path distance
                   shortPath = paths[i]
              for j in range(len(paths[i])-1):
                   T[paths[i][j], paths[i][j+1]] += 1/path distance
                   T[\mathsf{paths[i][j+1]}, \mathsf{paths[i][j]]} \; +\!\!= \; 1/\mathsf{path\_distance}
     shortestDistance.append(shortestDist)
     shortestPath.append(shortPath)
    if iteration % 40 == 0:
         plt.figure(figsize=(7,7))
         plt.plot(r[:,0], r[:,1], 'o')
plt.plot(r[s,0], r[s,1], 's', color='red', markersize=10)
plt.plot(r[t,0], r[t,1], 's', color='red', markersize=10)
         for i in range(N):
              for j in range(i+1,N):
                   if M[i,j] == 1:
                        plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'k--')
         p = shortestPath[iteration]
         for q in range(len(p)-1):
              i = p[q]
              j = p[q+1]
              plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'black', linewidth=5)
         plt.title('Iteration ' + str(iteration+1)+' - Shortest Path')
         plt.axis('off')
         plt.show()
    if iteration % 40 == 0:
         plt.figure(figsize=(7,7))
         plt.plot(r[:,0], r[:,1], 'o')
plt.plot(r[s,0], r[s,1], 's', color='red', markersize=10)
plt.plot(r[t,0], r[t,1], 's', color='red', markersize=10)
         for i in range(N):
              for j in range(i+1,N):
                   if M[i,j] == 1:
                        plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'k--')
         for i in range(N):
              for j in range(i+1,N):
         plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]],'r',linewidth=T[i,j]*2)
plt.title('Iteration ' + str(iteration+1) + ' - Pheromone Matrix')
         plt.axis('off')
         plt.show()
\verb|plt.plot(np.linspace(0,number_of_iterations-1,number_of_iterations)|, shortestDistance)|
plt.xlabel("No. of Iterations")
plt.ylabel("Shortest Path Distance")
```

Iteration 1 - Shortest Path

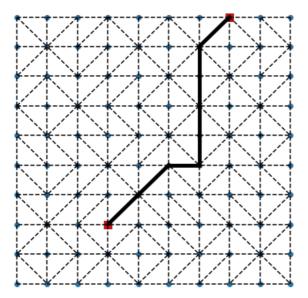


Iteration 1 - Pheromone Matrix

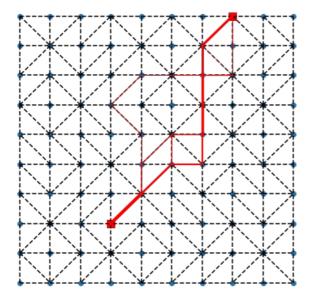




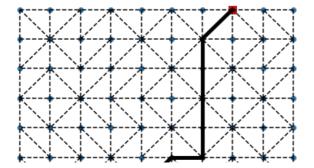
Iteration 41 - Shortest Path

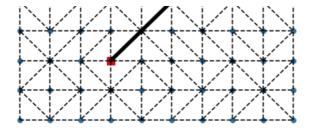


Iteration 41 - Pheromone Matrix

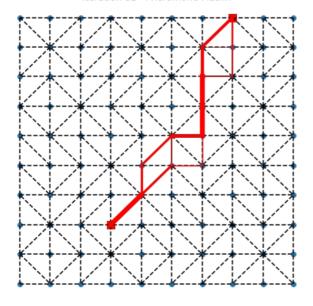


Iteration 81 - Shortest Path

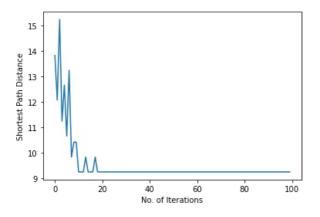




Iteration 81 - Pheromone Matrix



Out[4]: Text(0, 0.5, 'Shortest Path Distance')



```
In [5]: from scipy.spatial import Voronoi
          n = 20
                              # number of ants
          alpha = 0.8
                              # alpha parameter
          beta = 1
                            # beta parameter
          rho = 0.5
                            # rho parameter
          max steps = 80  # maximum number of steps
          number of iterations = 300 # number of iterations
          N = 100 # number of nodes
          \# generate N points at unique random positions (x,y) with x,y in [0,N]
          all_points = []
                                        # Generate an array of all the possible points in grid
          for i in range(10):
              for j in range(10):
                  all_points.append([i,j])
          # generate N random unique points from all_points
indices = np.random.choice(len(all_points), N, replace=False)
          r = np.array([all_points[i] for i in indices])
```

```
# connect the generated points by a Delaunay triangulation
tri = Voronoi(r)
# create the corresponding adjacency matrix
M = np.zeros((N, N))
for simplex in tri.ridge_points:
    for i in range(len(simplex)):
        for j in range(i+1, len(simplex)):
            M[simplex[i], simplex[j]] = 1
M[simplex[j], simplex[i]] = 1
# define the Euclidean distance all points in r
D = np.zeros((N, N))
for i in range(N):
    for j in range(i+1,N):
        D[i,j] = np.linalg.norm(r[i]-r[j])
        D[j,i] = D[i,j]
# make the distance inf for points that are not connected
for i in range(N):
    for j in range(i+1,N):
        if M[i,j] == 0:
            D[i,j] = np.inf
             D[j,i] = np.inf
T = M # initialize the Pheromone matrix, initially equal to the adjacency matrix
distance bird = 0
while distance bird < 7:</pre>
                                         # we don't want the starting and the target node to be too close
    # decide a starting node s
    s = np.random.randint(0, N)
    # decide a target node t
    t = np.random.randint(0, N)
    distance_bird = np.sqrt((r[t,0] - r[s,0])**2 + (r[t,1] - r[s,1])**2)
shortestDistance = []
shortestPath = []
for iteration in range(number of iterations):
    # initialize the paths of the ants
    T = T*(1-rho)
    paths = []
    for i in range(n):
        paths.append([s])
        for step in range(max_steps):
             next_node = branch_decision(paths[i][-1], T, D, M, alpha, beta)
             paths[i].append(next_node)
             if next node == t:
                 break
    # check if the target node is in the path of any of the ants
    shortestDist = np.inf
    for i in range(n)
        if t in paths[i]:
             paths[i] = simplify sequence(paths[i])
             path distance = total distance(paths[i],D)
             if path_distance<shortestDist:</pre>
                 shortestDist = path_distance
                 shortPath = paths[i]
             for j in range(len(paths[i])-1):
    T[paths[i][j],paths[i][j+1]] += 1/path_distance
                 T[paths[i][j+1],paths[i][j]] += 1/path distance
    shortestDistance.append(shortestDist)
    shortestPath.append(shortPath)
    if iteration % 40 == 0:
        plt.figure(figsize=(7,7))
        plt.plot(r[:,0], r[:,1], 'o')
plt.plot(r[s,0], r[s,1], 's', color='red', markersize=10)
plt.plot(r[t,0], r[t,1], 's', color='red', markersize=10)
        for i in range(N):
             for j in range(i+1,N):
                 if M[i,j] == 1:
                     plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'k--')
        p = shortestPath[iteration]
        for q in range(len(p)-1):
            i = p[q]
             j = p[q+1]
             plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], black', linewidth=5)
        plt.title('Iteration ' + str(iteration+1)+' - Shortest Path')
        plt.axis('off')
        plt.show()
    if iteration % 40 == 0:
        plt.figure(figsize=(7,7))
```

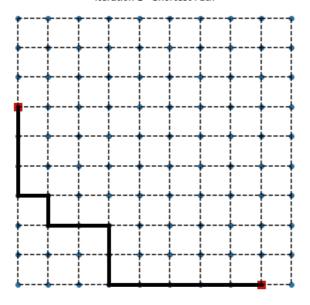
```
plt.plot(r[:,0], r[:,1], 'o')
plt.plot(r[s,0], r[s,1], 's', color='red', markersize=10)
plt.plot(r[t,0], r[t,1], 's', color='red', markersize=10)

for i in range(N):
    for j in range(i+1,N):
        if M[i,j] == 1:
            plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'k--')

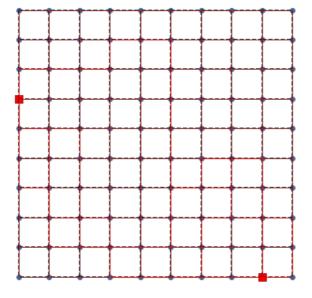
for i in range(N):
    for j in range(i+1,N):
        plt.plot([r[i,0], r[j,0]], [r[i,1], r[j,1]], 'r',linewidth=T[i,j]*2)
plt.title('Iteration ' + str(iteration+1) + ' - Pheromone Matrix')
plt.axis('off')
plt.show()

plt.plot(np.linspace(0,number_of_iterations-1,number_of_iterations),shortestDistance)
plt.xlabel("No. of Iterations")
plt.ylabel("Shortest Path Distance")
```

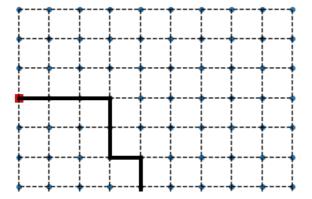
Iteration 1 - Shortest Path

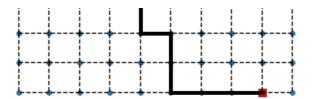


Iteration 1 - Pheromone Matrix

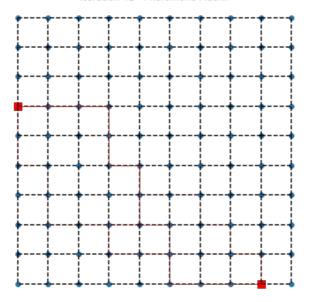


Iteration 41 - Shortest Path

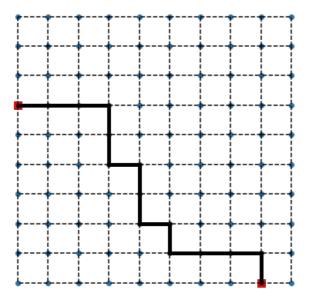




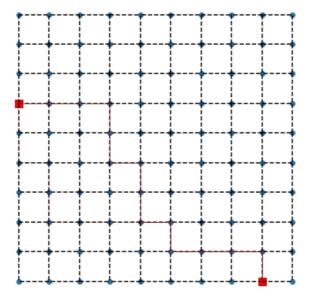
Iteration 41 - Pheromone Matrix



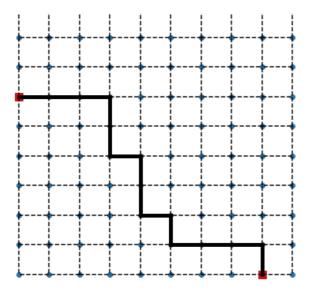
Iteration 81 - Shortest Path



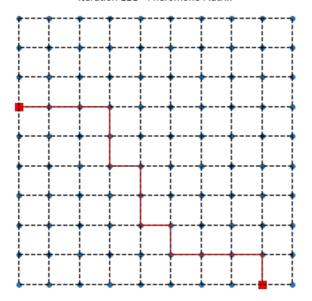
Iteration 81 - Pheromone Matrix



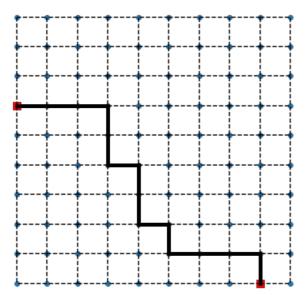
Iteration 121 - Shortest Path



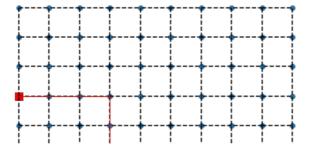
Iteration 121 - Pheromone Matrix

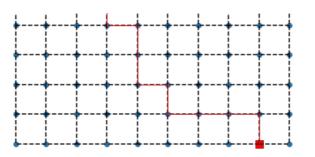


Iteration 161 - Shortest Path

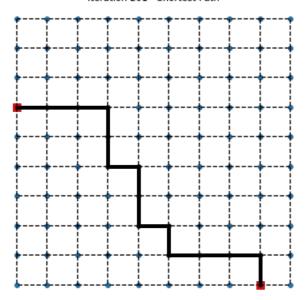


Iteration 161 - Pheromone Matrix

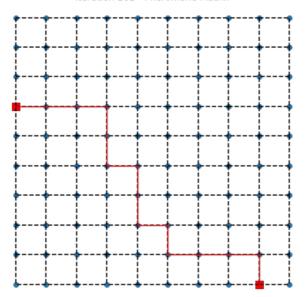




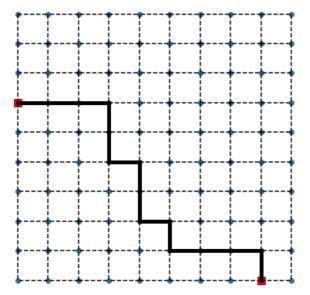
Iteration 201 - Shortest Path

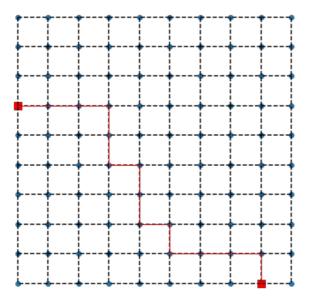


Iteration 201 - Pheromone Matrix

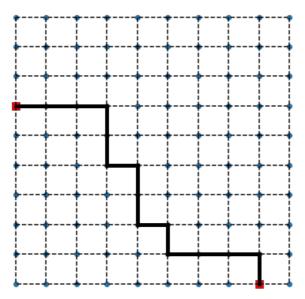


Iteration 241 - Shortest Path

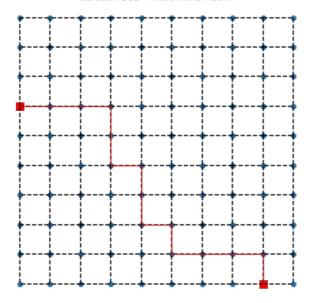




Iteration 281 - Shortest Path

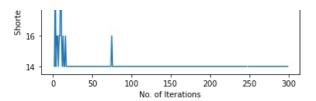


Iteration 281 - Pheromone Matrix



Out[5]: Text(0, 0.5, 'Shortest Path Distance')





In []:

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