



```

In [42]: from matplotlib import pyplot as plt
import numpy as np

# experiment 2 rendition 2

import numpy as np
import matplotlib.pyplot as plt

# Create some mock data
t = np.arange(0.01, 10.0, 0.01)
data1 = [74.4044311142, 74.2625051138, 73.7763650102, 72.617034761, 71.12093181819999, 69.1502188716, 66.3084190
data2 = [137.52, 124.26, 110.74, 96.14, 82.34, 69.06, 56.36, 45.68, 37.52, 30.14]

vac = data1
sus = data2

vac_std = [0.01202053, 1.04794591, 1.24330532, 1.70601292, 2.19675224, 2.834807, 3.71498164, 4.25674477, 4.57118
sus_std = [7.7051671, 7.21612084, 6.99945712, 7.93727913, 8.2331282, 8.28832914, 8.65507943, 7.93080072, 7.032

fig, ax1 = plt.subplots()

ax1.set_xlabel('Duration', fontsize=14)
ax1.set_ylabel('Sum of Vaccines', color='blue', fontsize=14)
ax1.tick_params(axis='y', labelcolor='blue')
ax1.plot([i*5 for i in range(len(vac))], vac, color='blue')
ax1.plot([i*5 for i in range(len(vac))], [vac[i] + vac_std[i] for i in range(len(vac))], color='blue', linestyle=
ax1.plot([i*5 for i in range(len(vac))], [vac[i] - vac_std[i] for i in range(len(vac))], color='blue', linestyle=
ax1.fill_between([i*5 for i in range(len(vac))], vac, [vac[i] + vac_std[i] for i in range(len(vac))], color='#87
ax1.fill_between([i*5 for i in range(len(vac))], vac, [vac[i] - vac_std[i] for i in range(len(vac))], color='#87

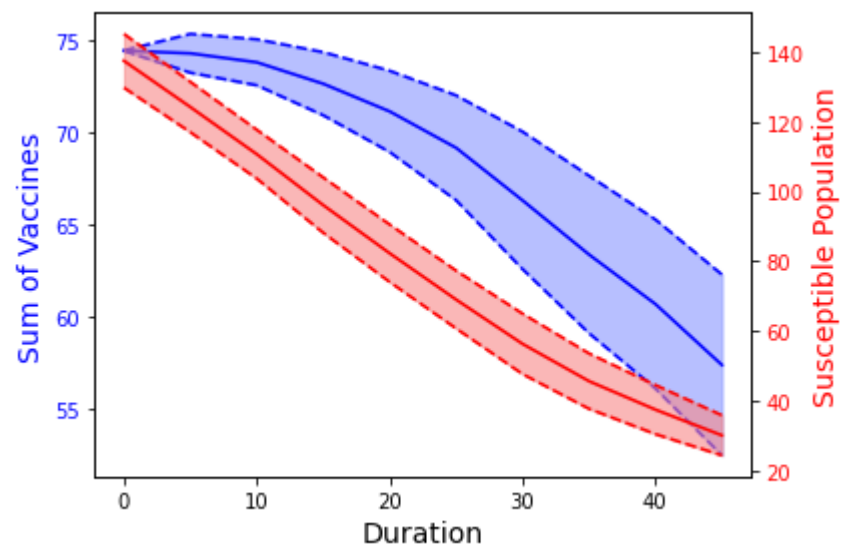
ax2 = ax1.twinx()

ax2.set_ylabel('Susceptible Population', color='red', fontsize=14)
ax2.tick_params(axis='y', labelcolor='red')
ax2.plot([i*5 for i in range(len(sus))], sus, color='red')
ax2.plot([i*5 for i in range(len(sus))], [sus[i] + sus_std[i] for i in range(len(sus))], color='red', linestyle=
ax2.plot([i*5 for i in range(len(sus))], [sus[i] - sus_std[i] for i in range(len(sus))], color='red', linestyle=
ax2.fill_between([i*5 for i in range(len(sus))], sus, [sus[i] + sus_std[i] for i in range(len(sus))], color='#f7
ax2.fill_between([i*5 for i in range(len(sus))], sus, [sus[i] - sus_std[i] for i in range(len(sus))], color='#f7

fig.tight_layout() # otherwise the right y-label is slightly clipped
plt.savefig('experiment_condition_2_3_final.png', dpi=300)

```

```
plt.show()
```



In [41]: # Experiment 1

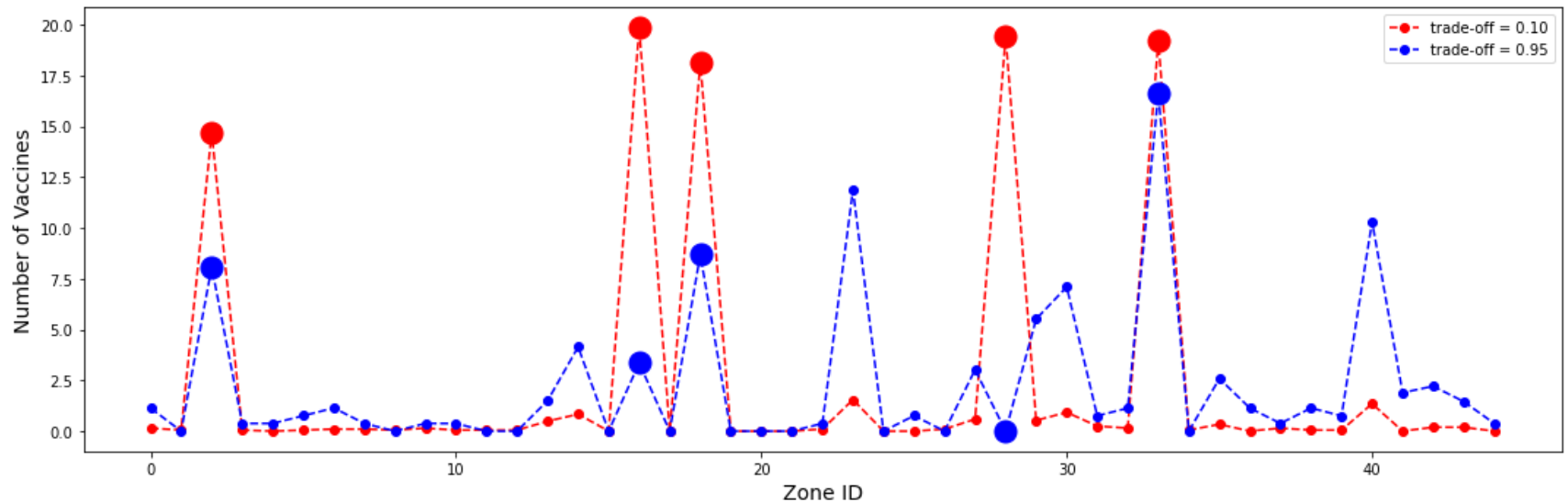
```
p0_1 = [0.14747104, 0.049438067, 14.685921, 0.049310038, 0.0, 0.049428155, 0.09890848, 0.098838255, 0.049349479,
p0_95 = [1.1307041, 0.0, 8.0685203, 0.37807464, 0.37894179, 0.75796056, 1.1375408, 0.37891106, 0.0, 0.37909878,

LW = [18, 28, 16, 2, 33]

plt.figure(figsize=(15,5))
plt.plot([i for i in range(45)], p0_1, color='red', label='trade-off = 0.10', linestyle='--', marker='o')
plt.plot([i for i in range(45)], p0_95, color='blue', label='trade-off = 0.95', linestyle='--', marker='o')

plt.plot([i for i in LW], [p0_1[i] for i in LW], color='red', marker='o', markersize=15, linewidth=0)
plt.plot([i for i in LW], [p0_95[i] for i in LW], color='blue', marker='o', markersize=15, linewidth=0)

plt.ylabel('Number of Vaccines', fontsize=14)
plt.xlabel('Zone ID', fontsize=14)
plt.legend()
plt.tight_layout()
plt.savefig('experiment_condition_1_3_final.png', dpi=300)
plt.show()
```



In [ ]: *# Learning Rate Results*

```
lr0_1 = [0.4, 0.4099999999375003, 0.3832857142780485, 0.35924285713391707, 0.36617571427154577, 0.3438438571280
lr0_2 = [0.4, 0.32, 0.330999999990625, 0.31479999998625, 0.30183999998275, 0.316471999976825, 0.30317759997521,
lr0_002 = [0.4, 0.4001999999987504, 0.399399599998753, 0.3996008007997505, 0.3990515991981198, 0.3985034959996

plt.plot([i for i in range(len(lr0_1))], lr0_1, color='red', label='lr = 0.1')
plt.plot([i for i in range(len(lr0_2))], lr0_2, color='blue', label='lr = 0.2')
plt.plot([i for i in range(len(lr0_002))], lr0_002, color='green', label='lr = 0.002')
plt.plot([i for i in range(len(lr0_002))], [0.2 for i in range(len(lr0_002))], linestyle='--', color='black', la
plt.ylabel('Learning Rate', fontsize=14)
plt.xlabel('Duration', fontsize=14)
plt.gca().invert_yaxis()
plt.legend()
plt.tight_layout()
plt.savefig('learning_rate_630.png', dpi=300)
plt.show()
```

```
In [ ]: import pulp
import random
import networkx as nx
import simpy
import numpy as np
import pickle
import random
import math
import matplotlib.pyplot as plt
import operator
import itertools
import pandas as pd
import decimal
import matplotlib.pyplot as plt
from geopy import distance
from geopy.distance import geodesic
from scipy.spatial.distance import *
from scipy.optimize import minimize, differential_evolution
from scipy import optimize
from sklearn.cluster import KMeans
from sklearn.metrics import silhouette_score, calinski_harabasz_score

final_arr = [[] for i in range(50)]
learning_rate_change = []
double_arr = []
sus_arr = [[] for i in range(50)]

for alphabetagammaepsilon in range(50):

    class Node(object):

        def __init__(self, env, ID, coor, state, alpha, beta, gamma, delta, sigma, zone):

            global T, d

            self.ID = ID
            self.env = env
            self.alpha = alpha
            self.beta = beta
            self.gamma = gamma
            self.delta = delta
            self.sigma = sigma
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self.zone = zone

# Neighbor List
self.nlist = []

self.old_coor = None
self.new_coor = coor
self.start = True

self.ti = 3

self.state = state

if self.ID == 1:
    self.env.process(self.time_increment())
    self.env.process(self.optimizer())
    d = []

self.env.process(self.move())
self.env.process(self.influence())

def move(self):

    global Xlim, Ylim, W, a, zone_coordinates, R

    while True:

        # if T % mho == 0 and self.state != 'D':
        if T % mho == 0:

            c = zone_coordinates[self.zone]
            r = R[self.zone]

            # Define a set of k random points (potential next positions) within the circle of my current
            k = 10
            P = []

            # Calculate the distance between current location (p) and each potential next hop
            D = []

            for i in range(k):
                x = random.uniform(c[0] - r, c[0] + r)
                y = random.uniform(c[1] - r, c[1] + r)

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        P.append((x, y))
        D.append(euclidean(self.new_coor, (x, y)))

        # Select the next destination from P preferring short distances over long distances
        likelihood_of_selecting = [1.0/D[i] for i in range(k)]
        likelihood_of_selecting = [likelihood_of_selecting[i]/np.sum(likelihood_of_selecting) for i
        ind = np.random.choice([i for i in range(k)], p = likelihood_of_selecting, size = 1)[0]

        # New position of current agent
        self.new_coor = P[ind]

    yield self.env.timeout(minimumWaitingTime)

def scan_neighbors(self):

    global eG, sensing_range, entities, Coor

    while True:

        if T % PT == 2:

            self.nlist = []

            if self.start:
                for u in range(eG):
                    if euclidean(self.new_coor, Coor[u]) <= sensing_range:
                        self.nlist.append(u)
                self.start = False

            else:
                for u in range(eG):
                    if euclidean(self.new_coor, entities[u].new_coor) <= sensing_range:
                        self.nlist.append(u)

            self.nlist = [u for u in self.nlist if u != self.ID]

        yield self.env.timeout(minimumWaitingTime)

def influence(self):

    global minimumWaitingTime

    while True:

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if T % PT == (self.ti + 1) % PT:

    state_change = False

    if self.state == 'S':
        for u in self.nlist:
            if entities[u].state == 'E' and random.uniform(0, 1) <= self.alpha:
                self.state = 'E'
                state_change = True
                break
        ...
    if self.state == 'S' and state_change == False:
        for u in self.nlist:
            if entities[u].state == 'I' and random.uniform(0, 1) <= self.beta:
                self.state = 'I'
                state_change = True
                break
        ...

    if self.state == 'E' and state_change == False:
        if random.uniform(0, 1) <= self.gamma:
            self.state = 'I'
            state_change = True

        ...
    if self.state == 'E' and state_change == False:
        if random.uniform(0, 1) <= pi:
            self.state = 'R'
            state_change = True
        ...

    if self.state == 'I' and state_change == False:
        if random.uniform(0, 1) <= self.delta:
            self.state = 'R'
            state_change = True

    if self.state == 'I' and state_change == False:
        if random.uniform(0, 1) <= self.sigma:
            self.state = 'D'
            state_change = True

    yield self.env.timeout(minimumWaitingTime)
```

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def time_increment(self):

    global Tracker, T, D, sus, exp, inf, rec, dth

    while True:

        T = T + 1
        sus = len([i for i in range(eG) if entities[i].state == 'S'])
        exp = len([i for i in range(eG) if entities[i].state == 'E'])
        inf = len([i for i in range(eG) if entities[i].state == 'I'])
        rec = len([i for i in range(eG) if entities[i].state == 'R'])
        dth = len([i for i in range(eG) if entities[i].state == 'D'])

        # print('sus: ' + str(sus) + ', exp: ' + str(exp) + ', inf: ' + str(inf) + ', rec: ' + str(rec)

        d.append((inf, rec, dth))

        # print (self.new_coor)
        if T % mho == 0 and self.old_coor != None:
            plt.scatter(self.new_coor[0], self.new_coor[1], s = 10, c = 'green')
            plt.plot([self.old_coor[0], self.new_coor[0]], [self.old_coor[1], self.new_coor[1]], line

        yield self.env.timeout(minimumWaitingTime)

def optimizer(self):

    global I, E, S, z, r, T, vaccines, f_interval, learning_rate_change

    while True:
        if T % vaccine_interval == 0:

            vaccines_per_zone = resource_allocation(0.3, vaccines, T)
            arr_infected, arr_suspected, arr_exposed = np.zeros(z), np.zeros(z), np.zeros(z)

            for delta in range(z):
                arr_infected[delta] = len([i for i in range(eG) if entities[i].state == 'I' and enti
                arr_suspected[delta] = len([i for i in range(eG) if entities[i].state == 'S' and ent
                arr_exposed[delta] = len([i for i in range(eG) if entities[i].state == 'E' and entit

                available_vaccine = vaccines_per_zone[delta]
                arr = [iota for iota in range(len(agent_zones)) if ((agent_zones[iota]==delta and er

                immune, vaccinated = [], []

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        for phi in range(len(arr)):
            while(available_vaccine > 0):
                initial_state = entities[arr[phi]].state
                entities[arr[phi]].state = np.random.choice([initial_state, 'R'], size=1, p=
                vaccinated.append(arr[phi])
                if(entities[arr[phi]].state == "R"):
                    immune.append(arr[phi])
                    if initial_state == 'S':
                        arr_suspected[delta] -= 1
                    elif initial_state == 'E':
                        arr_exposed[delta] -= 1
                available_vaccine -= 1
                break

            r[delta] = r[delta] + (((len(immune))/(len(vaccinated)+0.00000001)) - r[delta]) * le

        learning_rate_change.append(r[2])

        I = np.array(arr_infected)
        S = np.array(arr_suspected)
        E = np.array(arr_exposed)

        yield self.env.timeout(minimumWaitingTime)

def prospect(x, beta, lambdas):

    if x == 0:
        return 1

    return lambdas * math.pow(x, beta)

def count_extra(A):

    how_many_used = 0
    extra = 0

    for i in range(A.shape[0]):
        print (A[i])

        if np.sum(A[i]) > 0:
            how_many_used += 1

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        if np.sum(A[i]) < 1.0:
            extra += 1.0 - np.sum(A[i])
    return how_many_used, extra

def latp(pt, W, a):

    # Available Locations
    AL = [k for k in W.keys() if euclidean(W[k], pt) > 0]
    AL = cutoff(AL, W, pt)

    if len(AL) == 0:
        return pt

    den = np.sum([1.0 / math.pow(float(euclidean(W[k], pt))), a] for k in sorted(AL))

    plist = [(1.0 / math.pow(float(euclidean(W[k], pt))), a) / den] for k in sorted(AL)]

    next_stop = np.random.choice([k for k in sorted(AL)], p = plist, size = 1)

    return W[next_stop[0]]

def least_distance_per_cluster(C, arr):

    array = np.zeros((len(arr), len(arr)))
    for row in range(len(arr)):
        for column in range(len(arr)):
            array[row][column] = euclidean(C[arr[row]], C[arr[column]])
    avg_distance = (np.mean(array, axis=1)).reshape(len(arr), )
    least = np.amin(avg_distance)
    result = 0
    for et in range(len(avg_distance)):
        if avg_distance[et] == least:
            result = arr[et]
            break
    return result

def def_zone_and_coor():

    global population_val, C

    population_val = np.array(file['Population'].values)
    pop = np.true_divide(population_val, np.sum(population_val))

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C = []
coordinates = np.array(file['Location'].values)
for a in range(0, len(coordinates)):
    arr = coordinates[a].split(',')
    for b in range(0, len(arr)):
        arr[b] = float(arr[b])
    C.append((arr[1], arr[0]))
agent_zones = np.random.choice(len(C), size=eG, p=pop)
agent_initial_coordinates = [C[d] for d in agent_zones]

return C, agent_zones, agent_initial_coordinates

def radius():

    global population_val, file

    area = np.true_divide(np.array(file['Population'].values), np.array(file['Population Density'].values))
    rad = [math.sqrt(area[i]/math.pi) for i in range(len(area))] / ((np.max(population_val))/eG)
    return rad

def initial_state():

    global infected_ratio, susceptible_ratio, exposed_ratio, population_val

    infected_ratio = np.true_divide(np.array(file['Total Infected'].values), np.array(file['Population'].values))
    exposed_ratio = pe * (1 - infected_ratio)
    susceptible_ratio = 1 - (infected_ratio + exposed_ratio)
    initial_state = [(np.random.choice(['S', 'E', 'I'], size=1,
                                         p=[susceptible_ratio[agent_zones[c]],
                                              exposed_ratio[agent_zones[c]],
                                              infected_ratio[agent_zones[c]]])[0]) for c in range(len(agent_zones))]

    return initial_state

def initial_state():

    global infected_ratio, susceptible_ratio, exposed_ratio, population_val

    infected_ratio = np.true_divide(np.array(file['Total Infected'].values), np.array(file['Population'].values))
    exposed_ratio = pe * (1 - infected_ratio)
    susceptible_ratio = 1 - (infected_ratio + exposed_ratio)
    initial_state = [(np.random.choice(['S', 'E', 'I'], size=1,
                                         p=[susceptible_ratio[agent_zones[c]],
                                              exposed_ratio[agent_zones[c]],
                                              infected_ratio[agent_zones[c]]])[0]) for c in range(len(agent_zones))]

```

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infected_ratio[(agent_zones[c])][0]) for c in range(len(agent_zones))]

return initial_state

def resource_allocation(p, T, time):

    global trade_off, B, N, I, E, S, C, r

    # Low trade-off favor economic and high trade-off favors vaccine formulation
    trade_off = 0.95

    how_many = warehouse

    kmeans = KMeans(n_clusters=warehouse, random_state=0).fit(C)
    cluster_center = kmeans.cluster_centers_
    cluster_labels = kmeans.labels_

    label_arr = [[] for alpha in range(warehouse)]
    for beta in range(len(cluster_labels)):
        label_arr[cluster_labels[beta]].append(beta)

    # List of warehouses
    LW = [least_distance_per_cluster(C, label_arr[gamma]) for gamma in range(warehouse)]
    print('LW: ', LW)

    ...

    #Add the function to generate warehouses
    array = np.zeros((z, len(LW)))
    for row in range(z):
        for column in range(len(LW)):
            array[row][column] = euclidean(C[row], C[LW[column]])
    ...

    # Equally distributing vaccines across warehouse zones
    VW = {}
    current_warehouse = 0
    for f in range(1, T + 1):
        VW[f - 1] = LW[current_warehouse]
        if (f % (T / warehouse) == 0):
            current_warehouse += 1
    #print(VW.values())

    # Defining the parameters for the optimization
    B = np.array(file['Population Density'].values) * p # rate of disease spread

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N = np.array(num_agent_per_zone) # total population for each zone
if time <= vaccine_interval:
    r = np.array([0.4 for app in range(z)])
    I = np.array([infected_ratio[t]*num_agent_per_zone[t] for t in range(len(N))])
    E = (N-I) * pe
    S = N - (I + E)

    ...

B = np.array([i + 1 for i in range(z)])
N = np.array([z - i for i in range(z)])
I = np.array([random.uniform(1, N[i]) for i in range(z)])
E = (N-I) * pe
S = N - (I + E)

B = np.array([2 for i in range(z)])
N = np.array([100 for i in range(z)])
I = np.array([25 for i in range(z)])
E = (N-I) * pe
S = N - (I + E)
...

ir = I/(N+0.00001)

model = pulp.LpProblem("Vaccine problem", pulp.LpMinimize)
X = pulp.LpVariable.dicts("X", ((i, j) for i in range(warehouse) for j in range(len(B))), lowBound = 0.0)

dist_array = [geodesic(C[VW[j]], C[b]).miles for j in range(T) for b in range(z)]
max_dist = np.max(dist_array)
den_economic = float(T * max_dist)
model += np.sum([X[LW.index(VW[j]), b] * geodesic(C[VW[j]], C[b]).miles for j in range(T) for b in range(z)])

# Constraint 1 -----
for i in range(warehouse):
    # Condition 1: If you must assign all the vaccines generated by a warehouse
    #model += pulp.lpSum([X[(i, j)] for j in range(len(B))]) == int(T/warehouse)

    # Condition 2: If you want to minimize the number of vaccines
    model += pulp.lpSum([X[(i, j)] for j in range(len(B))]) <= int(T/warehouse)

# Constraint 2 -----
s = 0.0
for i in range(warehouse):

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s += pulp.lpSum([X[(i, j)] for j in range(len(B))])

# Condition 1: If you must assign all the vaccines generated by a warehouse
#model += s == T

# Condition 2: If you want to minimize the number of vaccines
model += s <= T

# Constraint 3 (fairness lower) -----
for i in range(len(B)):

    # Condition 3: If calculation is based on susceptible population
    c = (S[i] - r[i] * pulp.lpSum([X[(j, i)] for j in range(warehouse)])) / sum(S)

    # Condition 4: To include population density
    #c = c * B[i] / np.median(B)

    # Condition 5: To include infected population
    #c = c * ir[i] / np.median(ir)

    model += pulp.lpSum([X[(j, i)] for j in range(warehouse)]) >= trade_off * c * T

# -----

model.solve()
print(pulp.LpStatus[model.status])

# Transferred the pulp decision to the numpy array (A)
A = np.zeros((T, len(B)))
for i in range(warehouse):
    for j in range(len(B)):
        A[i, j] = X[(i, j)].varValue

global double_arr
double_arr.append(["Value: " + str(pulp.value(model.objective))])

vaccines_per_zone = []
for i in range(len(B)):
    vaccines_per_zone.append(np.sum(A[:, i]))

print('Trade-off value: ' + str(trade_off))

global final_arr, sus

```



```
final_arr[alphabetaepsilon].append(np.sum(vaccines_per_zone))
sus_arr[alphabetaepsilon].append(sus)

plt.figure(figsize=(10,4))
plt.bar([i for i in range(z)], vaccines_per_zone)
print(vaccines_per_zone)
#plt.tight_layout()
#plt.savefig('plot.png', dpi=300)
plt.show()
print('var', alphabetaepsilon)
return np.array(vaccines_per_zone)

''' Variables and Parameters for Simulation '''

# Create Simpy environment and assign nodes to it.
env = simpy.Environment()

# Susceptible-> Exposed
alpha = 0.2

# Susceptible-> Infected
beta = 0.2

# Exposed-> Infected
gamma = 0.2

# Infected-> Recovered
delta = 0.2

# Infected-> Death
sigma = 0.2

PT = 10

# Fraction of susceptible/exposed nodes
pe = 0.3

# Simulation area --> not relevant right now
Xlim, Ylim = 100, 100

# Number of agents, zones, warehouses, and vaccines (optimization parameters)
eG = 200
```

```
z = 45
warehouse = 5
vaccines = 100

# Simulation time-variable
T = 0

# Simulation duration
Duration = 50

# Move how often
mho = 3

# Time intervals for administering vaccines
vaccine_interval = 5

# Minimum waiting time
minimumWaitingTime = 1

# Variable used to increase proportion of infected (in case it's too low)
infected_bias = 0.0

sensing_range = 30

# File used for importing data
file = pd.read_csv("covid_confirmed_NY_july.csv")
file = file.iloc[0:z,:]

# Initial position and coordinates of node based on population density Likelihood PW
zone_coordinates, agent_zones, Coor = def_zone_and_coor()

# Scaled radius of each zone
R = radius()

# Number of agents in each zone
num_agent_per_zone = [0 for i in range(z)]
for epsilon in range(len(agent_zones)):
    num_agent_per_zone[agent_zones[epsilon]] += 1

print("Number of agents per zone: ", num_agent_per_zone)

# Learning rate variables
expectedR = [0.2 for var in range(z)]
```

```
learning_rate = 0.15
learning_rate_change.append(0.4)

# List of node initial states
STATE = initial_state()

entities = [Node(env, i, Coord[i], STATE[i], alpha, beta, gamma, delta, sigma, agent_zones[i]) for i in range
env.run(until = Duration)

print('\n\n-----')
print('vac-avg', np.sum(final_arr, axis=0))
print('vac-std', np.std(final_arr, axis=0))
print('sus-avg', np.sum(sus_arr, axis=0))
print('sus-std', np.std(sus_arr, axis=0))
```