

Covid-19 Simulation In A Society.

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#COVID-19 Simulation
library(ggplot2)

#initializing the parameters for the simulation
N_total_population=10000
recovery_time=10
simulation_time=150#days
initially_infect=2
mortality_rate=0.002
infection_rate=0.3
immune_coeff=0.1

#initialise citizen
#0--->healthy citizen
#1--->initial infected citizen
#2--->dead citizen

citizens=rep(0, N_total_population)
citizens[sample(1:N_total_population, initially_infect)]=1

#data frame for simulation
df_simulate=data.frame(day=0, sick=sum(citizens==1), deaths=0, new_infections=0)

#loop for simulation
for(day in 1:simulation_time)
{

  #we start by assuming all citizens are alive and healthy
  new_infections=0
  deaths=0
  sick_citizens=which(citizens==1)

  #no. of deaths based on a mortality rate.
  deaths=sum(rnorm(length(sick_citizens))<mortality_rate)

  #every citizen is meeting a randomly selected person from the population b/w 0 to 20
  who is sick.
  contacts=sample(N_total_population,
size=sum(sample(0:20,length(sick_citizens),replace=TRUE)))

  #total no. of new infections based on contacts and infection rate
  new_infections=sum(citizens[contacts]==0 & runif(length(contacts))<infection_rate)

  #After 10 days a sick citizen becomes healthy and stops spreading virus.
  #Hence checking if the person is recovered and marking them healthy(3)
  recovery_citizens=sick_citizens[runif(length(sick_citizens))<inverse_recovery_time]
  citizens[recovery_citizens]=3

  #citizens who are recovered(partial immunity) after surviving the infection and
  getting healthy.
  #which reduces the chances of getting sick in future if they meet a sick person.
  immune_citizens=which(citizens == 3)
  contacts=sample(N_total_population, size=sum(sample(0:20,length(immune_citizens),
replace=TRUE)))
  new_infections_immunity=sum(citizens[contacts] == 0 & runif(length(contacts))
<infection_rate*immune_coeff)

  #updating the status of the citizens to know if the citizens are immune or still
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sick.
  citizens[sick_citizens]=ifelse(runif(length(sick_citizens)) < inverse_recovery_time,
3, 1)
  citizens[immune_citizens]=ifelse(runif(length(immune_citizens)) <
infection_rate*immune_coeff, 1, 3)

#updating the simulation data by adding a row which contains current day,
#the no. of sick individuals, the number of deaths, and the number of new infections.
df_simulate=rbind(df_simulate, data.frame(day=day, sick=sum(citizens == 1),
                                           deaths=deaths,
new_infections=new_infections+new_infections_immunity))
}

#printing the data
print("Simulation Data")
print(df_simulate)

#plotting the simulation data as line plot graph.
ggplot(df_simulate, aes(x = day)) +
  geom_line(aes(y = sick, color = "Sick"), size = 1) +
  geom_line(aes(y = deaths, color = "Deaths"), size = 1) +
  geom_line(aes(y = new_infections, color = "New Infections"), size = 1) +
  labs(x = "Day", y = "Number of Citizens", color = "Status") +
  scale_color_manual(values = c("Sick" = "orange", "Deaths" = "red", "New Infections" =
"cornflowerblue")) +
  ggtitle("COVID-19 Simulation") +
  theme_dark()

#plotting the simulation data as pie chart.
status_sum <- data.frame(
  Status = c("Sick", "Death", "New Infection"),
  Count = c(
    sum(df_simulate$sick),
    sum(df_simulate$deaths),
    sum(df_simulate$new_infections)
  )
)
status_sum$Percentage=status_sum$Count/sum(status_sum$Count)*100
ggplot(status_sum, aes(x = "", y = Count, fill = Status)) +
  geom_bar(stat = "identity", width = 1) +
  coord_polar(theta = "y") +
  labs(x = NULL, y = NULL, fill = "Status") +
  ggtitle("COVID-19 Simulation") +
  geom_text(aes(label = paste0(round(Percentage, 1), "%")),
            position = position_stack(vjust = 0.5)) +
  theme_dark()

library(csv)
# Assuming you have a data frame called "df_simulate"
# Specify the file path and name
file_path <- "C:/Users/spand/OneDrive/Desktop/New folder/COVID STIMULATION.csv"
# Export the data frame to Excel
write.csv(df_simulate, file_path, row.names = FALSE)
# Confirm the export
print("Data exported to Excel successfully.")

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