

Wuhan to SF infection model

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1 Introduction

This is an exploration of a simple model to see much time there is after the start of an epidemic to stop transportation between two cities in order to prevent the spread from one to another. The model looks similar to [1], but I made a simpler model of spreading, quarantine, and transit between two cities, and I made lots of other simplifying assumptions too. This model is not intended to make quantitative predictions. It's just meant to illustrate that there is a limited amount of time to stop transit before it's too late to contain the spread.

2 Model Definition

I adapted a standard SIR model in the limit where there are lots of people in the susceptible (S) stage and no one in the recovered (R) stage. Therefore this model only tries to measure $I(t)$ in two populations. I also assumed that there is a spreading rate that goes as α and a quarantine rate that gets smaller when there are more people who are sick (assuming that we are using up limited healthcare resources) $\beta + \kappa/I$. Finally, there is some small factor μ that defines the rate of people moving from the infected city to the non-infected city. Combining these assumptions into an equation and setting $\gamma = \alpha - \beta$ we get the primary model equation.

$$\frac{dI_{wu}}{dt} = \gamma I_{wu} - \kappa \tag{1}$$

$$\frac{dI_{sf}}{dt} = \gamma I_{sf} - \kappa + \mu I_{wu} \tag{2}$$

I should point out that the equation for I_{sf} is only valid when it doesn't send I_{sf} negative. Whenever $I_{sf} = 0$ and the equation for $dI_{sf}/dt < 0$ we just set $dI_{sf}/dt = 0$.

This equation still has 3 free parameters, which makes it cumbersome to simulate all possible outcomes. In the next section I renormalize variables to transform this into a single parameter differential equation.

3 Nondimensionalization

Following, [2], I perform a normalization of our variables, t , I_{wu} , and I_{sf} by the constants t_0 , I_0 , giving us

$$w = \frac{I_{wu}}{I_0}, x = \frac{I_{sf}}{I_0}, \tau = \frac{t}{t_0} \quad (3)$$

and

$$\frac{dw}{d\tau} = \frac{dI_{wu}}{dt} \frac{t_0}{I_0} \quad (4)$$

$$\frac{dx}{d\tau} = \frac{dI_{sf}}{dt} \frac{t_0}{I_0} \quad (5)$$

I sub these into equations 1 to get

$$\frac{dw}{d\tau} \frac{I_0}{t_0} = \gamma w I_0 - \kappa \quad (6)$$

$$\frac{dx}{d\tau} \frac{I_0}{t_0} = \gamma x I_0 - \kappa + \mu w I_0 \quad (7)$$

and rearranging to get

$$\frac{dw}{d\tau} = \gamma w t_0 - \kappa \frac{t_0}{I_0} \quad (8)$$

$$\frac{dx}{d\tau} = \gamma x t_0 - \kappa \frac{t_0}{I_0} + \mu w t_0 \quad (9)$$

We can set the first term in each equation to 1 by choosing $t_0 = 1/\gamma$. After that substitution, we can use $I_0 = \kappa/\gamma$ to set the second term to 1 as well, leaving

$$\frac{dw}{d\tau} = w - 1 \quad (10)$$

$$\frac{dx}{d\tau} = x - 1 + \frac{\mu}{\gamma} w \quad (11)$$

Finally, we rename the variable μ/γ to ρ for simplicity leaving our final dimensionless dynamical equation

$$\frac{dw}{d\tau} = w - 1 \quad (12)$$

$$\frac{dx}{d\tau} = x - 1 + \rho w \quad (13)$$

which we can use to simulate the dynamics without worrying about so many parameters.

It's important to remember that for this equation to hold, we must assume that ρ is much less than 1. Otherwise, we would need to account for the $-\rho w$ in our equation for w . This is reasonable though because we expect $\mu \ll \gamma$, since the probability of infecting another person is much higher than the probability of flying to San Francisco.

4 Initial infected population

The solution for $w(\tau)$ with $w(0) = n$ (from [wolfram alpha](#)) is

$$w(\tau) = (n - 1)e^\tau + 1 \quad (14)$$

To make our model interesting we assume that the infected population in Wuhan is going to be growing. This amounts to assuming that the initial value of w at $\tau = 0$ (ie n) is such that $dw/dt > 0$ or $n > 1$.

If we make this assumption it becomes cleaner to describe [14](#) in terms of a new variable $\varepsilon = n - 1$, which is how much the starting infected population in Wuhan exceeds the minimum needed for an outbreak in Wuhan, 1. So we can rewrite [14](#) as

$$w(\tau) = \varepsilon e^\tau + 1 \quad (15)$$

5 Outbreak Critical Time

Now we can get to the interesting part, where we can estimate the critical time where we reach an outbreak in San Francisco, τ_x .

There are actually two distinct phases in this critical time. First, there is the time until the infected population in SF becomes greater than 0, meaning the number of infected people arriving exceeds the capacity to quarantine them fast enough. I'll label this time as τ_+ .

Second, there is the time from when SF's infected population starts to grow up until the point where the infection becomes self-sustaining even if the influx from Wuhan goes to 0. Let's say that we renormalize time after τ_+ as $\tau' = \tau - \tau_+$. So I'll call the time when SF's τ'_x .

So in the end we $\tau_x = \tau_+ + \tau'_x$, and we need to calculate both of these terms

5.1 Calculating time to SF spread

We can calculate τ_+ directly from the solution to our differential equation. The infection only starts spreading in SF once we have $dx/d\tau > 0$. We can sub the equation for $w(\tau)$ [15](#) into the equation for $dx/d\tau$ to get

$$\frac{dx}{d\tau} = x - 1 + \rho(\varepsilon e^\tau + 1) > 0 \quad (16)$$

Since x is by definition 0 up until this point we can set it so in the equation and rearrange to get

$$\rho(\varepsilon e^\tau + 1) > 1 \quad (17)$$

and rearranging further we get to

$$\tau_+ = \ln \left(\frac{1 - \rho}{\rho \varepsilon} \right) \quad (18)$$

5.2 Estimating time to self-sustaining spread in SF

If at some point we stop further input from Wuhan, (ie set $\rho = 0$), the infection in SF will continue to spread anyway if $x > 1$ at that point. We can evaluate $x(\tau)$ in the time after τ_+ by subbing in equation 15, shifting our time variable to $\tau' = \tau - \tau_+$, and solving equation 13 with $x(\tau') = 0$ at $\tau' = 0$ (using [wolfram alpha](#)).

$$x(\tau') = \rho \varepsilon e^{\tau'} + (1 - \rho)(1 - e^{\tau'}) \quad (19)$$

Unfortunately, finding the time τ'_x when this is 1 isn't possible because of the mixture of exponential and linear τ' . However we can estimate it in two different limiting cases, one where τ'_x is really short and another where τ'_x is really long.

5.2.1 Limit where $\tau'_x \ll 1$

If we assume that we're in the limit where the time to outbreak (τ'_x) will be small then we can expand $e^{\tau'}$ as $1 + \tau'$, approximate 19, and find when $x(\tau'_x) = 1$.

$$x(\tau'_x) \approx \tau'_x(\rho \varepsilon - (1 - \rho)) = 1 \quad (20)$$

and rearranging for τ'_x

$$\tau_x = \ln \left(\frac{1 - \rho}{\rho \varepsilon} \right) = \frac{1}{\rho(\varepsilon + 1) - 1} \quad (21)$$

In order for τ'_x to be $\ll 1$, we need the denominator to be large. But since, $\rho \ll 1$ we know that what we really need is $\varepsilon \gg 1/\rho$. In this limit we can say that $\tau'_x \approx 0$ or in other words, $\tau_x \approx \tau_+$

5.2.2 Limit where $\tau'_x \gg 1$

In the opposite limit, I just assume that the first term in 19 dominates to we have

$$x(\tau'_x) \approx \rho \varepsilon e^{\tau'_x} = 1 \quad (22)$$

According to [wolfram alpha](#) this is solved by $\tau'_x = W(1/\rho \varepsilon)$, where W is the product log function. This is some crazy function that I don't know how

to think about, but it seems to be mostly just something like $0.8 \cdot \ln(1/\rho\varepsilon)$ for values of $1/(\rho\varepsilon)$ from 10 to 100000 and asymptotically approaching $\ln(1/\rho\varepsilon)$ as $\rho\varepsilon \rightarrow 0$.

So I'll just approximate it as $0.8 \cdot \ln(1/\rho\varepsilon)$ and give our final result for τ_x in this limit as

$$\tau_x = \ln\left(\frac{1-\rho}{\rho\varepsilon}\right) + 0.8 \cdot \ln(1/\rho\varepsilon) \quad (23)$$

Finally, since we assume that $\rho \ll 1$, we can drop the $\ln(1-\rho)$ and simplify the logarithms to

$$\tau_x = 1.8 * \ln \frac{1}{\rho\varepsilon} \quad (24)$$

6 Conclusion

To see how this compares to simulated results, check out [this jupyter notebook](#).

References

- [1] Aleksa Zlojutro, David Rey, and Lauren Gardner. A decision-support framework to optimize border control for global outbreak mitigation. *Scientific Reports*, 9(1):2216, 2019.
- [2] Steven H. Strogatz. *Nonlinear Dynamics and Chaos: With Applications to Physics, Biology, Chemistry and Engineering*. Westview Press, 2000.