

# Herd immunity in a network

---

Jens Kinch, Nikolai Plambech Nielsen & Mads Ehrhorn

28th of July 2017

Niels Bohr Institute  
University of Copenhagen

# Herd immunity

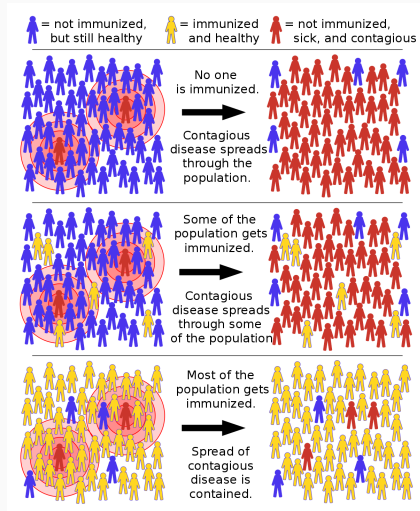


Figure 1: By Tkarcher - Own work, CC BY-SA 4.0,  
<https://commons.wikimedia.org/w/index.php?curid=56760604>

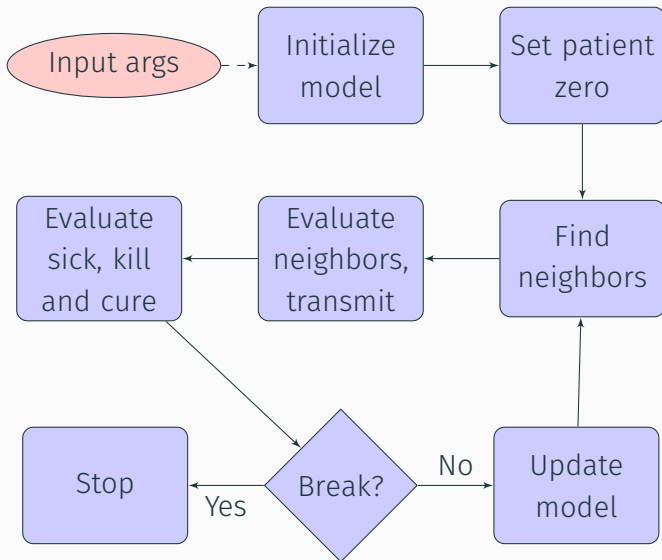
## Let's frontload some variables!

- $R_0$ : 'basic reproduction number'
  - Avg. no. of people infected pr. person
- $HIT = 1 - 1/R_0$
- $N$ : no. of people initially
- $n_{\text{dead}}$ : no. of dead people
- $n_{\text{healthy}}$ : no. of healthy people (not immune)

	$R_0$	Mortality rate	HIT
Ebola	1.5-2.5	0.25-0.90	0.33-0.60
Measles	12-18	0.15	0.92-0.94
Polio	5-7	0.15-0.30	0.80-0.86

Table 1: Data from [https://en.wikipedia.org/wiki/Herd\\_immunity](https://en.wikipedia.org/wiki/Herd_immunity)

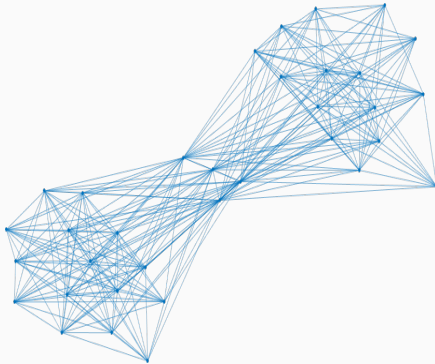
# Code



- Small world
- Scale free
- Random
- Custom network (two cities with commuters. Random networks for each city)

# Custom network

Custom network, simulating two cities with commuters



Movie time: two cities, death



## Movie time: two cities, immunity

Movie time: two cities, immunity

# Success criteria

- Real world
  - Disease no longer endemic
- Our model
  - Discussion of alternate criteria
    - Percolation
    - Effective reproductive number
      - $R_e = \text{fraction of susceptible individuals} \cdot R_0 \leq 1$
    - $n_{\text{sick}} = 0$  and  $n_{\text{healthy}} \neq 0$
    - $n_{\text{healthy}} = 0$
    - **Total sick < arbitrary threshold**

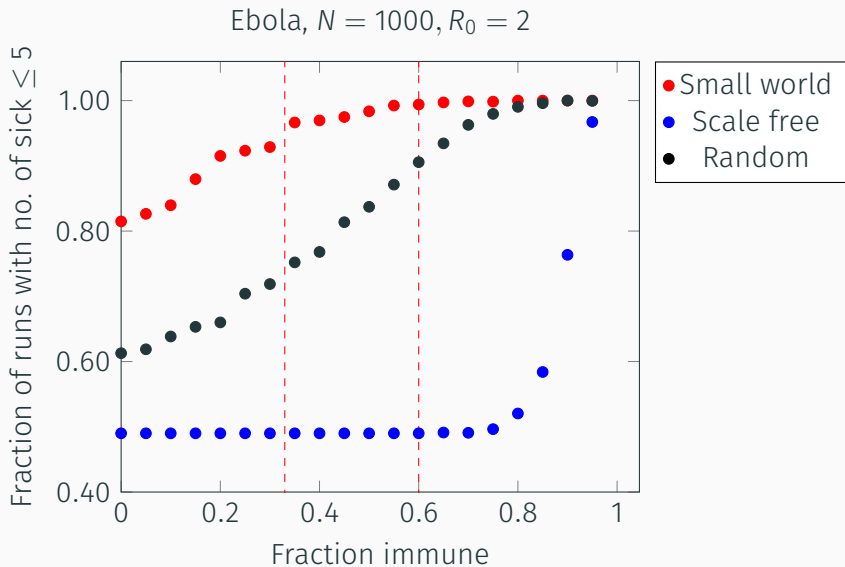
# The simulations

- Test 20 values of  $p_I$  (0 to 0.95, in steps of 0.05)

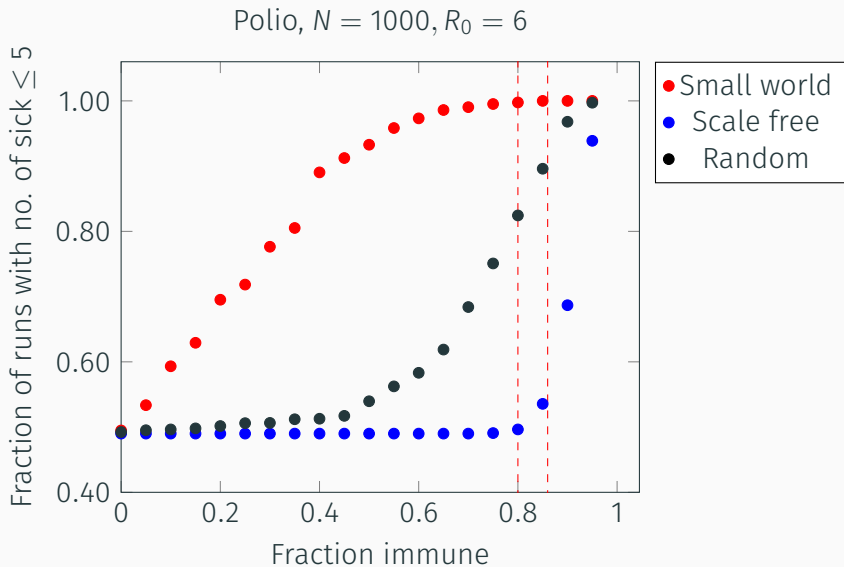
# The simulations

- Test 20 values of  $p_I$  (0 to 0.95, in steps of 0.05)
- For each  $p_I$  test 50 different networks, and test each network 50 times (random distribution of patient zero and immune nodes)

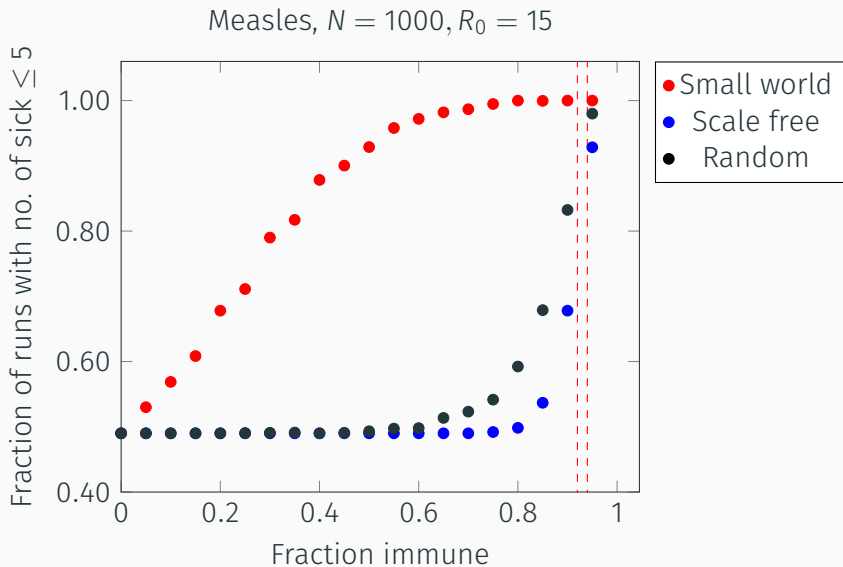
# Results



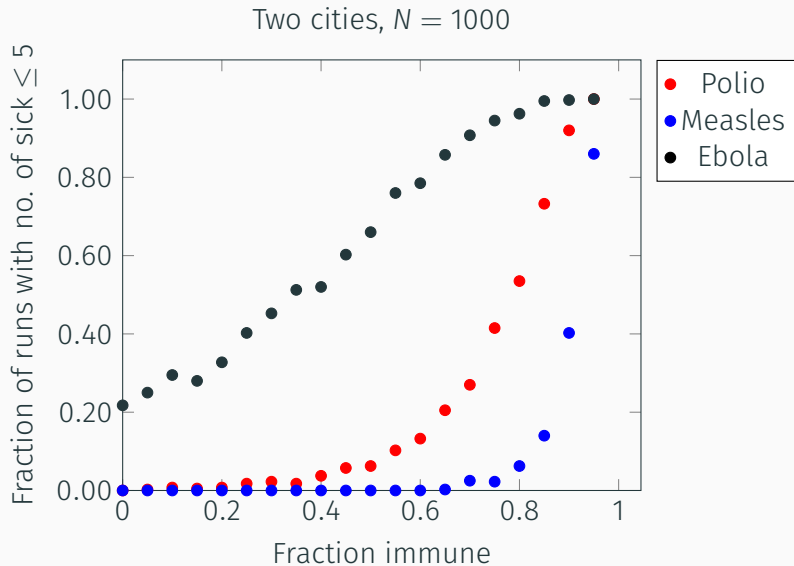
# Results



# Results



# Results





# Conclusion

Simulating the spread of disease is easy. Defining herd immunity is hard.