Combining Behavioral Choice with a Branching Process Model of Disease

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Disease Spread

- ► (Newman 2002) describes a class of networks on which an SIR model can be solved exactly.
- Social network is an infinite random graph described by degree distribution $\{p_k\}$
- ► Contagion can spread along each edge with probability *T*

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$$T_{c} = \frac{\sum_{k} (p_{k}k)}{\sum_{k} (p_{k}k(k-1))}$$

▶ If $T < T_c$, epidemic occurs with zero probability.

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- ightharpoonup Given degree distribution, there is a critical transmissibility threshold T_c
- ▶ When $T > T_c$, the probability an epidemic occurs equals the expected portion of nodes which become infected. Denoted R_{∞}

$$R_{\infty} = 1 - \sum_{k} (p_k \cdot (1 - (1 - v) T)^k)$$

where $v \in (0,1)$ is the solution to

$$v = \frac{\sum_{k} \left(p_{k} k \cdot (1 - (1 - v) T)^{k} \right)}{\sum_{k} \left(p_{k} k \right)}$$

Important Variables so Far

- $ightharpoonup \{p_k\}$ is the degree distribution of the network.
- ► *T* is transmissibility.
- $ightharpoonup T_c$ is the critical transmissibility threshold.
- $ightharpoonup R_{\infty}$ is the probability and size of epidemic when $T > T_c$
- ightharpoonup v can be thought of as the chance a random neighbor remains uninfected.
- lacktriangle Finally, define the risk of disease from a neighbor Ψ as

$$\psi \equiv \begin{cases} 0 & \text{if } T \le T_c \\ (1 - v)T & \text{if } T > T_c \end{cases}$$

Individual Choice and Equilibrium

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► Each person *makes this choice exactly once*, when news of a potential epidemic arrives.

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► The critical transmissibility threshold is given by:

$$T_c(\{N_i\}) = \frac{\sum_i \alpha_i N_i}{\sum_i \alpha_i N_i^2}$$

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▶ The probability and size of the epidemic when $T > T_c$ is given by

$$R_{\infty} = 1 - \sum_{i} \left[\alpha_{i} e^{-(1-v)TN_{i}} \right]$$

where v is the solution to

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- And finally, let $Ψ^*({N_i})$ be the value of Ψ, taken as a function of the set of choices.
 - When $T \le T_c(\{N_i\}), \Psi^*(\{N_i\}) = 0$
 - When $T > T_c(\{N_i\})$, $\Psi^*(\{N_i\})$ is the solution $\Psi \in (0,1)$ to:

$$\Psi = T \frac{\sum_{i} A_{i} N_{i} (1 - e^{-\Psi N_{i}})}{\sum_{i} A_{i} N_{i}}$$

► The payoff for a person of type *i* is

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- ▶ $1 e^{-\Psi N_i}$ is the probability of getting sick during this outbreak.
- $ightharpoonup \delta_i$ is the disutility from getting sick.

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 - The total payoff $U_i(N_i; \Psi)$ is continuous and concave down,
 - and $N_i^*(\Psi)$, the person's optimal policy function, is a continuous and bounded function of Ψ over $\Psi \in [0,1]$

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- ▶ If $\delta_i = 1$ for all i, then the following function has these properties:

$$u_i(N) = \ln\left(\frac{N}{\theta_i}\right) - \frac{N}{\theta_i}$$

where θ_i is the person's optimal choice when $\Psi = 0$

Equilibrium

Given exogenous T, $\{\alpha_i\}$, an equilibrium in this model consists of Ψ , N_i such that

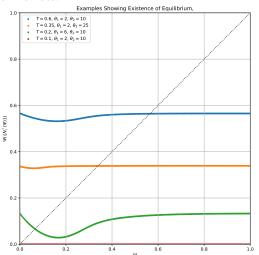
$$\Psi = \Psi^* \left(\{ N_i \} \right)$$
 $N_i = N_i^* (\Psi) \equiv \arg \max U_i(N_i; \Psi)$

Equilibrium Existence

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- ▶ Proposition 2: Iff $T \le T_c(\{N_i^*(0)\})$, then there is an equilibrium without any risk of epidemic exists, where $\Psi = 0$ and $N_i = N_i^*(0)$ for all i.

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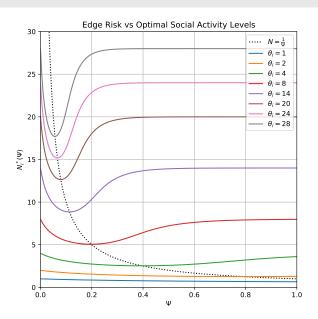
- ightharpoonup An individual's disease risk is an increasing function of both Ψ and N_i
- Nowever, the marginal disease risk from N_i may sometimes decrease as Ψ increases.

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- ► However, the marginal disease risk from N_i may sometimes decrease as Ψ increases.

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• When $\Psi > \frac{1}{N_i}$, an *increase* in disease risk may lead to individuals trying *less* hard to avoid getting sick.



When can *N_i* have *positive* externalities?

▶ **Proposition 3:** Suppose $\{N_i\}$ is such that $T > T_c(\{N_i\})$. In this case,

$$\frac{\partial \Psi^*\left(\{N_i\}\right)}{\partial N_j} < 0$$

$$\updownarrow$$

$$(1 - e^{-\Psi^*\left(\{N_i\}\right)N_j}) < \frac{\Psi\left(\{N_i\}\right)}{T} \left(1 - TN_j e^{-\Psi\left(\{N_i\}\right)N_j}\right)$$

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- ▶ Corollary: If $N_j > \frac{1}{\Psi^*(\{N_i\})}$ and $T > T_c(\{N_i\})$, then $\frac{\partial \Psi^*(\{N_i\})}{\partial N_j} > 0$

Related Literature

- ► (Newman 2002) Describes a model of disease spread based on branching processes and uses it to explicitly solve SIR models for a class of networks.
- ► (Meyers et al. 2005) shows that this works well to approximate the behavior of complex social networks.
- ► (Kremer 1996) Demonstrates similar results regarding fatalism and counter-intuitive externalities in a model of the steady-state of an endemic disease, rather than a disease outbreak.