

The Spread of COVID-19 in London: Network Effects and Optimal Lockdowns?

Christian Julliard Ran Shi Kathy Yuan

London School of Economics

November 5, 2020

Motivation

- The COVID-19 pandemic: 1.2M world-wide deaths, 47k in UK so far
- Unprecedented policy responses – lockdowns – to control the spread
- Significant social and economic cost
- Urgent need to evaluate effectiveness and inform optimal policy making
- So far, no spatial element in models of COVID-19 spread
- We use London as a salient case study and:
 - embed network elements in the workhorse SIR epidemiology model
 - map heterogeneous interactions among population using commuting networks
 - estimate the model and analyse effectiveness of policy response
- Given the estimated spatial dynamics, we study (counterfactual) optimal policy responses

Motivation

- The COVID-19 pandemic: 1.2M world-wide deaths, 47k in UK so far
- Unprecedented policy responses – lockdowns – to control the spread
- Significant social and economic cost
- Urgent need to evaluate effectiveness and inform optimal policy making
- So far, no spatial element in models of COVID-19 spread
- We use London as a salient case study and:
 - embed network elements in the workhorse SIR epidemiology model
 - map heterogeneous interactions among population using commuting networks
 - estimate the model and analyse effectiveness of policy response
- Given the estimated spatial dynamics, we study (counterfactual) optimal policy responses

Main Findings

- Significant network externalities in COVID-19 epidemics
 - Decomposition of infections: network effects $\approx 42\%$; local/autoregressive $\approx 35\%$; endemic $\approx 23\%$.
 - Network Impulse-Response significantly different across geographic regions

⇒ Most infections \neq most externalities
- Lockdown policy has had major impact, with substantial network effects:
 - Total reduction of infected: 56.5%
 - Reduction due to network externalities: 36.1%
- Counterfactual analyses show that:
 - lockdown was somehow late, but further delay would have had more dramatic effects
 - the targeted lockdown of a small number of highly connected geographic regions is equally effective
 - targeted lockdowns based on a threshold of number of cases are not effective

Main Findings

- Significant network externalities in COVID-19 epidemics
 - Decomposition of infections: network effects $\approx 42\%$; local/autoregressive $\approx 35\%$; endemic $\approx 23\%$.
 - Network Impulse-Response significantly different across geographic regions
 \Rightarrow Most infections \neq most externalities
- Lockdown policy has had major impact, with substantial network effects:
 - Total reduction of infected: 56.5%
 - Reduction due to network externalities: 36.1%
- Counterfactual analyses show that:
 - lockdown was somehow late, but further delay would have had more dramatic effects
 - the targeted lockdown of a small number of highly connected geographic regions is equally effective
 - targeted lockdowns based on a threshold of number of cases are not effective

Main Findings

- Significant network externalities in COVID-19 epidemics
 - Decomposition of infections: network effects $\approx 42\%$; local/autoregressive $\approx 35\%$; endemic $\approx 23\%$.
 - Network Impulse-Response significantly different across geographic regions
 \Rightarrow Most infections \neq most externalities
- Lockdown policy has had major impact, with substantial network effects:
 - Total reduction of infected: 56.5%
 - Reduction due to network externalities: 36.1%
- Counterfactual analyses show that:
 - lockdown was somehow late, but further delay would have had more dramatic effects
 - the targeted lockdown of a small number of highly connected geographic regions is equally effective
 - targeted lockdowns based on a threshold of number of cases are not effective

Literature I

- On-going discussions of trade-off of lockdowns: the planner's problem in balancing economic output and controlling fatality
- Review by Avery, Bossert, Clark, Ellison and Ellison, 2020
- Calibrations of SIR models, cost of life, productivity
 - Formulate and solve a planner's dynamic control problem, (Alvarez, Argente and Lippi 2020)
 - Externalities and history dependent equilibrium (Rowthorn and Toxvaerd, 2020)
 - Value of death is major policy determinant (Garriga, Manuelli, Sanghi 2020)
 - Equilibrium interactions between economic decision and epidemic dynamics: (Eichbaum, Rebelo, and Trabandt 2020)
 - Contagion externalities (Jones, Philippon, and Venkateswaran 2020)
 - Individual optimal reduction in social activities in SIR dynamics: (Farboodi, Jarosch and Shimer, 2020)
 - Heterogeneity in social interaction and productivity among social groups (Acemoglu, Chernozhukov, Werning, Whinston, 2020)
- Estimation of vanilla SIR(D): Fernandez-Villaverde and Jones (2020)
- We depart by estimating the network element and demonstrating its relevance and policy potential.

Literature II

- Network literature on epidemiology
- Jackson 2008, Chapter 7, Sections 7.1, 7.2.
- Easley and Kleinberg, 2010, Chapter 21
- Key players in network (Ballester, Calvo-Armengol, and Zenou 2006; Denbee, Julliard, Li and Yuan 2020)

A Network SIR Model

SIR Models

The triplet: $\{S, I, R\}$ (Susceptible, Infectious, Removed) such that

$$dS = -\beta_I \frac{S}{N} Idt, \quad dI = \underbrace{\left(\beta_I \frac{S}{N} - \beta_R \right)}_{\alpha_t} Idt, \quad dR = \beta_R Idt$$

where: β_I = contact rate, β_R = removal rate; N = population.

Probabilistic counterpart (a continuous-time Markov chain):

$$\mathbb{P} [y \text{ new infections in } (t, t + dt) \mid I_t = x] = \begin{cases} \alpha_t x dt & y = 1 \\ o(dt) & y \geq 2 \end{cases}$$

Let $\mathbb{P} [y \text{ new infections in } (t, t + h) \mid I_t = x] = p_h(y \mid x)$

(holding α_t constant from t to $t + h$ and solving the Kolmogorov forward equation):

$$p_h(y \mid x) = \frac{\Gamma(y+x)}{\Gamma(x)\Gamma(y+1)} \left(e^{-\alpha_t h} \right)^x \left(1 - e^{-\alpha_t h} \right)^y$$

- probability mass function of the negative binomial distribution

SIR Models

The triplet: $\{S, I, R\}$ (Susceptible, Infectious, Removed) such that

$$dS = -\beta_I \frac{S}{N} Idt, \quad dI = \underbrace{\left(\beta_I \frac{S}{N} - \beta_R \right)}_{\alpha_t} Idt, \quad dR = \beta_R Idt$$

where: β_I = contact rate, β_R = removal rate; N = population.

Probabilistic counterpart (a continuous-time Markov chain):

$$\mathbb{P} [y \text{ new infections in } (t, t + dt) \mid I_t = x] = \begin{cases} \alpha_t x dt & y = 1 \\ o(dt) & y \geq 2 \end{cases}$$

Let $\mathbb{P} [y \text{ new infections in } (t, t + h) \mid I_t = x] = p_h(y \mid x)$

(holding α_t constant from t to $t + h$ and solving the Kolmogorov forward equation):

$$p_h(y \mid x) = \frac{\Gamma(y + x)}{\Gamma(x)\Gamma(y + 1)} \left(e^{-\alpha_t h} \right)^x \left(1 - e^{-\alpha_t h} \right)^y$$

- probability mass function of the negative binomial distribution

SIR Models

The triplet: $\{S, I, R\}$ (Susceptible, Infectious, Removed) such that

$$dS = -\beta_I \frac{S}{N} Idt, \quad dI = \underbrace{\left(\beta_I \frac{S}{N} - \beta_R \right)}_{\alpha_t} Idt, \quad dR = \beta_R Idt$$

where: β_I = contact rate, β_R = removal rate; N = population.

Probabilistic counterpart (a continuous-time Markov chain):

$$\mathbb{P} [y \text{ new infections in } (t, t + dt) \mid I_t = x] = \begin{cases} \alpha_t x dt & y = 1 \\ o(dt) & y \geq 2 \end{cases}$$

Let $\mathbb{P} [y \text{ new infections in } (t, t + h) \mid I_t = x] = p_h(y \mid x)$

(holding α_t constant from t to $t + h$ and solving the Kolmogorov forward equation):

$$p_h(y \mid x) = \frac{\Gamma(y + x)}{\Gamma(x)\Gamma(y + 1)} \left(e^{-\alpha_t h} \right)^x \left(1 - e^{-\alpha_t h} \right)^y$$

- probability mass function of the negative binomial distribution

SIR Models

- Normalizing h to $h = 1$, we have

$$y_t \mid \mathcal{F}_{t-1} \sim \text{NegBinom}(p_t, x_{t-1})$$

where

$$p_t = 1 - e^{-\alpha_t}$$

- If $S \approx N$, then $\alpha_t \equiv \alpha = \beta_I - \beta_R > 0$
- Infected remain infectious for L periods (weighted by decaying function $v(l)$)

$$x_t = \sum_{l=0}^{L-1} v(l)y_{t-l}$$

- Model moments

$$\mathbb{E}_{t-1}[y_t] = \mu_t = ax_{t-1}, \quad a = e^\alpha - 1 = \frac{p_t}{1-p_t} \in (0, \infty)$$

$$\text{var}_{t-1}[y_t] = (a+1)ax_{t-1}$$

SIR Models

- Normalizing h to $h = 1$, we have

$$y_t \mid \mathcal{F}_{t-1} \sim \text{NegBinom}(p_t, x_{t-1})$$

where

$$p_t = 1 - e^{-\alpha_t}$$

- If $S \approx N$, then $\alpha_t \equiv \alpha = \beta_I - \beta_R > 0$
- Infected remain infectious for L periods (weighted by decaying function $\nu(l)$)

$$x_t = \sum_{l=0}^{L-1} \nu(l) y_{t-l}$$

- Model moments

$$\mathbb{E}_{t-1}[y_t] = \mu_t = ax_{t-1}, \quad a = e^\alpha - 1 = \frac{p_t}{1-p_t} \in (0, \infty)$$

$$\text{var}_{t-1}[y_t] = (a+1)ax_{t-1}$$

SIR Models

- Normalizing h to $h = 1$, we have

$$y_t \mid \mathcal{F}_{t-1} \sim \text{NegBinom}(p_t, x_{t-1})$$

where

$$p_t = 1 - e^{-\alpha_t}$$

- If $S \approx N$, then $\alpha_t \equiv \alpha = \beta_I - \beta_R > 0$
- Infected remain infectious for L periods (weighted by decaying function $\nu(l)$)

$$x_t = \sum_{l=0}^{L-1} \nu(l) y_{t-l}$$

- Model moments

$$\mathbb{E}_{t-1}[y_t] = \mu_t = ax_{t-1}, \quad a = e^\alpha - 1 = \frac{p_t}{1-p_t} \in (0, \infty)$$

$$\text{var}_{t-1}[y_t] = (a+1)ax_{t-1}$$

A Network SIR Model: **Framework**

- Multiple subpopulations (local authorities) $i = 1, \dots, n$
 - Within subpopulation: autoregressive epidemic dynamics
 - Between subpopulations: network epidemic effects
- Distribution:

$$y_{i,t} \mid \mathcal{F}_{t-1} \sim \text{NegBinom}(p_{i,t}, x_{i,t-1}), \text{ and } \mathbb{E}_{t-1} [y_{i,t}] = \mu_{i,t}$$

- Multivariate extensions of $\mu_t = ax_{t-1}$:

$$\underbrace{\mu_t^{AR} + \mu_t^{NE}}_{\text{epidemic effects}} = Ax_{t-1} = \underbrace{\text{diag}(A)x_{t-1}}_{\mu_t^{AR}: \text{autoregressive}} + \underbrace{(A - \text{diag}(A))x_{t-1}}_{\mu_t^{NE}: \text{network}}$$

$$x_t = \sum_{l=0}^{L-1} v(l)y_{t-l}$$

$$\mu_t = \mu_t^{AR} + \mu_t^{NE} + \underbrace{\mu_t^{EN}}_{\text{endemic effects}}$$

A Network SIR Model: **Framework**

- Multiple subpopulations (local authorities) $i = 1, \dots, n$
 - Within subpopulation: autoregressive epidemic dynamics
 - Between subpopulations: network epidemic effects
- Distribution:

$$y_{i,t} \mid \mathcal{F}_{t-1} \sim \text{NegBinom}(p_{i,t}, x_{i,t-1}), \text{ and } \mathbb{E}_{t-1} [y_{i,t}] = \mu_{i,t}$$

- **Multivariate extensions** of $\mu_t = ax_{t-1}$:

$$\underbrace{\mu_t^{AR} + \mu_t^{NE}}_{\text{epidemic effects}} = Ax_{t-1} = \underbrace{\text{diag}(A)x_{t-1}}_{\mu_t^{AR}: \text{autoregressive}} + \underbrace{(A - \text{diag}(A))x_{t-1}}_{\mu_t^{NE}: \text{network}}$$

$$x_t = \sum_{l=0}^{L-1} \nu^{(l)} y_{t-l}$$

$$\mu_t = \mu_t^{AR} + \mu_t^{NE} + \underbrace{\mu_t^{EN}}_{\text{endemic effects}}$$

A Network SIR Model: **Spatial Form**

- Autoregressive effects: $\text{diag}(A) = \gamma I$

$$\mu_t^{AR} = \gamma x_{t-1}$$

- Network effects:** driven by an adjacency matrix W ($\text{diag}(W) = 0$)

$$\mu_{i,t}^{NE} = \phi \left(\sum_{j \neq i} w_{ij} x_{j,t-1} \right) \iff \mu_t^{NE} = \phi W x_{t-1}$$

- allowing for multiple graphs $\{W^{(g)}\}_{g=1}^G$:

$$\mu_{i,t}^{NE} = \sum_{g=1}^G \phi^{(g)} \left(\sum_{j \neq i} w_{ij}^{(g)} x_{j,t-1} \right)$$

- Implication: parametrize A as

$$A = \gamma I + \sum_{g=1}^G \phi^{(g)} W^{(g)}$$

A Network SIR Model: **Spatial Form**

- Autoregressive effects: $\text{diag}(A) = \gamma I$

$$\mu_t^{AR} = \gamma x_{t-1}$$

- Network effects:** driven by an adjacency matrix W ($\text{diag}(W) = 0$)

$$\mu_{i,t}^{NE} = \phi \left(\sum_{j \neq i} w_{ij} x_{j,t-1} \right) \iff \mu_t^{NE} = \phi W x_{t-1}$$

- allowing for multiple graphs $\{W^{(g)}\}_{g=1}^G$:

$$\mu_{i,t}^{NE} = \sum_{g=1}^G \phi^{(g)} \left(\sum_{j \neq i} w_{ij}^{(g)} x_{j,t-1} \right)$$

- Implication: parametrize A as

$$A = \gamma I + \sum_{g=1}^G \phi^{(g)} W^{(g)}$$

A Network SIR Model: **Spatial Form**

- Autoregressive effects: $\text{diag}(A) = \gamma I$

$$\mu_t^{AR} = \gamma x_{t-1}$$

- Network effects:** driven by an adjacency matrix W ($\text{diag}(W) = 0$)

$$\mu_{i,t}^{NE} = \phi \left(\sum_{j \neq i} w_{ij} x_{j,t-1} \right) \iff \mu_t^{NE} = \phi W x_{t-1}$$

- allowing for multiple graphs $\{W^{(g)}\}_{g=1}^G$:

$$\mu_{i,t}^{NE} = \sum_{g=1}^G \phi^{(g)} \left(\sum_{j \neq i} w_{ij}^{(g)} x_{j,t-1} \right)$$

- Implication: parametrize A as

$$A = \gamma I + \sum_{g=1}^G \phi^{(g)} W^{(g)}$$

A Network SIR Model: Policy Specification

- Policy regime $D_t \in \{0, 1\}$ may affect fundamental parameters $(\gamma, \phi^{(1)}, \dots, \phi^{(G)})$
- Autoregressive parameters:

$$\gamma(D_t) = \exp(D_t \gamma + \gamma_0)$$

- before policy: $\exp(\gamma_0)$; after policy: $\exp(\gamma_0 + \gamma)$
- policy impact: $\times \exp(\gamma)$
- Network parameters: for $g = 1, \dots, G$

$$\phi^{(g)}(D_t) = \exp(D_t \phi^{(g)} + \phi_0^{(g)})$$

- before policy: $\exp(\phi_0^{(g)})$; after policy: $\exp(\phi_0^{(g)} + \phi^{(g)})$
- policy impact: $\times \exp(\phi^{(g)})$
- Endemic effects:

$$\mu_{i,t}^{EN} = \exp(z_t^\top \beta + \eta_i) N_i$$

- z_t : time trends interacted with D_t / fixed effects / detection ratio / testing
- η_i : subpopulation fixed effects
- N_i : subpopulation size

A Network SIR Model: Policy Specification

- Policy regime $D_t \in \{0, 1\}$ may affect fundamental parameters $(\gamma, \phi^{(1)}, \dots, \phi^{(G)})$
- Autoregressive parameters:

$$\gamma(D_t) = \exp(D_t \gamma + \gamma_0)$$

- before policy: $\exp(\gamma_0)$; after policy: $\exp(\gamma_0 + \gamma)$
 - policy impact: $\times \exp(\gamma)$
- Network parameters: for $g = 1, \dots, G$

$$\phi^{(g)}(D_t) = \exp(D_t \phi^{(g)} + \phi_0^{(g)})$$

- before policy: $\exp(\phi_0^{(g)})$; after policy: $\exp(\phi_0^{(g)} + \phi^{(g)})$
 - policy impact: $\times \exp(\phi^{(g)})$
- Endemic effects:

$$\mu_{i,t}^{EN} = \exp(z_t^\top \beta + \eta_i) N_i$$

- z_t : time trends interacted with D_t / fixed effects / detection ratio / testing
- η_i : subpopulation fixed effects
- N_i : subpopulation size

A Network SIR Model: Policy Specification

- Policy regime $D_t \in \{0, 1\}$ may affect fundamental parameters $(\gamma, \phi^{(1)}, \dots, \phi^{(G)})$
- Autoregressive parameters:

$$\gamma(D_t) = \exp(D_t \gamma + \gamma_0)$$

- before policy: $\exp(\gamma_0)$; after policy: $\exp(\gamma_0 + \gamma)$
 - policy impact: $\times \exp(\gamma)$
- Network parameters: for $g = 1, \dots, G$

$$\phi^{(g)}(D_t) = \exp(D_t \phi^{(g)} + \phi_0^{(g)})$$

- before policy: $\exp(\phi_0^{(g)})$; after policy: $\exp(\phi_0^{(g)} + \phi^{(g)})$
 - policy impact: $\times \exp(\phi^{(g)})$
- Endemic effects:

$$\mu_{i,t}^{EN} = \exp(z_t^\top \beta + \eta_i) N_i$$

- z_t : time trends interacted with D_t / fixed effects / detection ratio / testing
- η_i : subpopulation fixed effects
- N_i : subpopulation size

Data and Estimation

Data and Estimation

- A case study of COVID-19 spread in London
 - Disease surveillance data of 32 local authorities
<https://coronavirus.data.gov.uk/>
 - Sample: March 1st to June 4th
 - **March 23rd** (nationwide lockdown announcement) = policy change date
($D_t = 0$ before and $D_t = 1$ afterwards)
- Networks: residence/work links from 2011 UK census
 - A matrix $K = \{k_{ij}\}$: number of people who lives at i & goes to work at j
 - $W^1 = K$: **work-to-home** transmission
 - $W^2 = K^\top$: **home-to-work** transmission
 - $W^3 = KK^\top$: **"home-to-home"** transmission via common work places
 - Rescaled by the largest singular values
- Local authority population sizes
<https://data.london.gov.uk>
- Maximum likelihood estimation

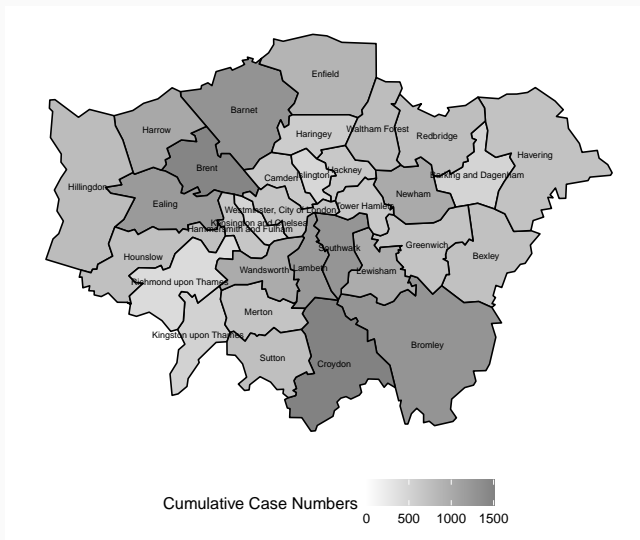
Data and Estimation

- A case study of COVID-19 spread in London
 - Disease surveillance data of 32 local authorities
<https://coronavirus.data.gov.uk/>
 - Sample: March 1st to June 4th
 - **March 23rd** (nationwide lockdown announcement) = policy change date
($D_t = 0$ before and $D_t = 1$ afterwards)
- Networks: residence/work links from 2011 UK census
 - A matrix $K = \{k_{ij}\}$: number of people who lives at i & goes to work at j
 - $W^1 = K$: **work-to-home** transmission
 - $W^2 = K^\top$: **home-to-work** transmission
 - $W^3 = KK^\top$: **“home-to-home”** transmission via common work places
 - Rescaled by the largest singular values
- Local authority population sizes
<https://data.london.gov.uk>
- Maximum likelihood estimation

Data and Estimation

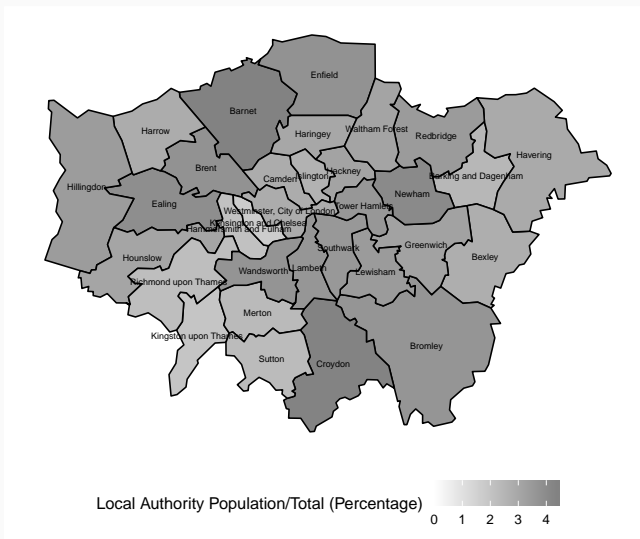
- A case study of COVID-19 spread in London
 - Disease surveillance data of 32 local authorities
<https://coronavirus.data.gov.uk/>
 - Sample: March 1st to June 4th
 - **March 23rd** (nationwide lockdown announcement) = policy change date
($D_t = 0$ before and $D_t = 1$ afterwards)
- Networks: residence/work links from 2011 UK census
 - A matrix $K = \{k_{ij}\}$: number of people who lives at i & goes to work at j
 - $W^1 = K$: **work-to-home** transmission
 - $W^2 = K^\top$: **home-to-work** transmission
 - $W^3 = KK^\top$: **“home-to-home”** transmission via common work places
 - Rescaled by the largest singular values
- Local authority population sizes
<https://data.london.gov.uk>
- Maximum likelihood estimation

Data: Cumulative Cases Till June 4th



Total: 27,078 (9.3% of UK)

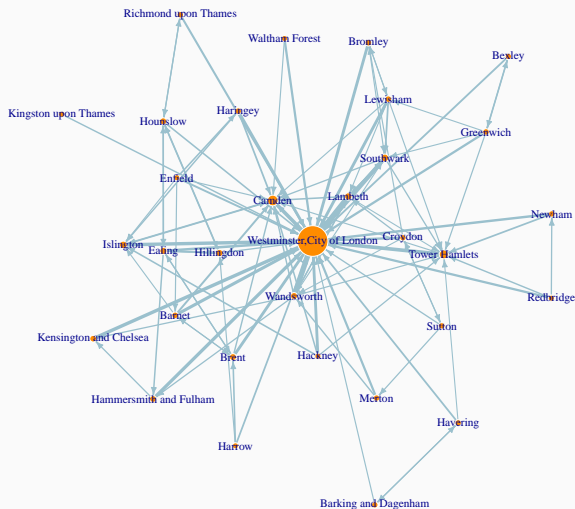
Data: Subpopulation Sizes



Corr(population; cases) = 0.63/0.78 pre/post lockdown

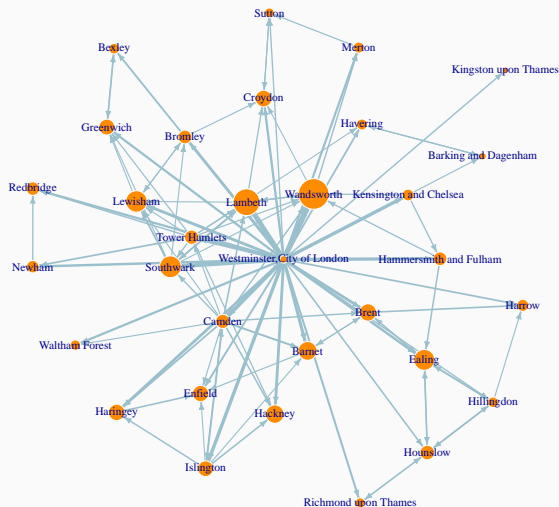
Data: Work-to-Home Transmission Network

Commuting from residential areas to work places: $W^{(1)} = K$



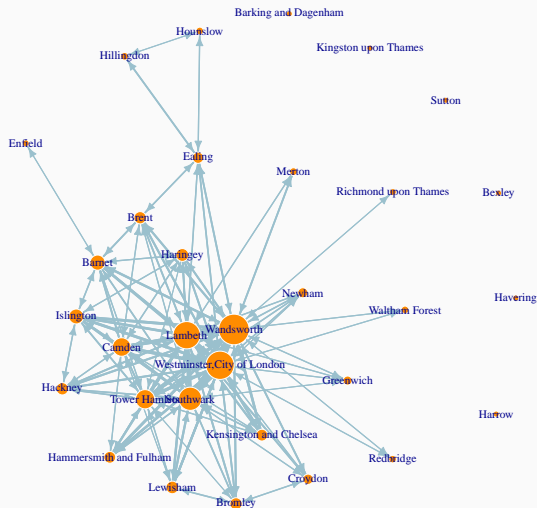
Data: Home-to-Work Transmission Network

Commuting from work places to residential areas: $W^{(2)} = K^\top$



Data: “Home-to-home” Transmission Network

Residential areas links via common work places: $W^{(3)} = KK^T$



Model Estimation

Value	(K)		(K^\top)		(KK^\top)		all		(K, K^\top)	
	est.	se.	est.	se.	est.	se.	est.	se.	est.	se.
Autoregressive effect										
$\exp(\gamma_0)$	0.311	0.035	0.286	0.040	0.286	0.037	0.252	0.041	0.253	0.041
$\exp(\gamma)$	0.617	0.077	0.958	0.136	0.750	0.102	0.760	0.129	0.754	0.127
Network effect										
K:										
$\exp(\phi_0)$	0.477	0.070					0.503	0.144	0.510	0.073
$\exp(\phi)$	0.885	0.136					0.821	0.169	0.827	0.127
K^\top :										
$\exp(\phi_0)$			0.115	0.022			0.062	0.036	0.064	0.023
$\exp(\phi)$			0.374	0.080			0.148	0.218	0.166	0.181
KK^\top :										
$\exp(\phi_0)$					0.261	0.035	0.000	0.085		
$\exp(\phi)$					0.570	0.082	0.284	0.000		
Testing-related endemic effect										
pos-to-test	-1.37	0.47	-1.09	0.28	-1.46	0.41	-1.46	0.48	-1.46	0.49
lag test	1.85	0.31	1.77	0.23	1.88	0.28	1.91	0.32	1.91	0.32
pseudo- R^2	82.09%		82.00%		82.02%		82.12%		82.12%	

est. w.o. control

Model Estimation

Value	(K)		(K [⊤])		(KK [⊤])		all		(K,K [⊤])	
	est.	se.	est.	se.	est.	se.	est.	se.	est.	se.
Autoregressive effect										
$\exp(\gamma_0)$	0.311	0.035	0.286	0.040	0.286	0.037	0.252	0.041	0.253	0.041
$\exp(\gamma)$	0.617	0.077	0.958	0.136	0.750	0.102	0.760	0.129	0.754	0.127
Network effect										
K:										
$\exp(\phi_0)$	0.477	0.070					0.503	0.144	0.510	0.073
$\exp(\phi)$	0.885	0.136					0.821	0.169	0.827	0.127
K [⊤] :										
$\exp(\phi_0)$			0.115	0.022			0.062	0.036	0.064	0.023
$\exp(\phi)$			0.374	0.080			0.148	0.218	0.166	0.181
KK [⊤] :										
$\exp(\phi_0)$					0.261	0.035	0.000	0.085		
$\exp(\phi)$					0.570	0.082	0.284	0.000		
Testing-related endemic effect										
pos-to-test	-1.37	0.47	-1.09	0.28	-1.46	0.41	-1.46	0.48	-1.46	0.49
lag test	1.85	0.31	1.77	0.23	1.88	0.28	1.91	0.32	1.91	0.32
pseudo-R ²	82.09%		82.00%		82.02%		82.12%		82.12%	

est. w.o. control

Model Estimation

Value	(K)		(K^\top)		(KK^\top)		all		(K, K^\top)	
	est.	se.	est.	se.	est.	se.	est.	se.	est.	se.
Autoregressive effect										
$\exp(\gamma_0)$	0.311	0.035	0.286	0.040	0.286	0.037	0.252	0.041	0.253	0.041
$\exp(\gamma)$	0.617	0.077	0.958	0.136	0.750	0.102	0.760	0.129	0.754	0.127
Network effect										
K:										
$\exp(\phi_0)$	0.477	0.070					0.503	0.144	0.510	0.073
$\exp(\phi)$	0.885	0.136					0.821	0.169	0.827	0.127
K^\top :										
$\exp(\phi_0)$			0.115	0.022			0.062	0.036	0.064	0.023
$\exp(\phi)$			0.374	0.080			0.148	0.218	0.166	0.181
KK^\top :										
$\exp(\phi_0)$					0.261	0.035	0.000	0.085		
$\exp(\phi)$					0.570	0.082	0.284	0.000		
Testing-related endemic effect										
pos-to-test	-1.37	0.47	-1.09	0.28	-1.46	0.41	-1.46	0.48	-1.46	0.49
lag test	1.85	0.31	1.77	0.23	1.88	0.28	1.91	0.32	1.91	0.32
pseudo- R^2	82.09%		82.00%		82.02%		82.12%		82.12%	

est. w.o. control

Model Estimation

Value	(K)		(K^\top)		(KK^\top)		all		(K, K^\top)	
	est.	se.	est.	se.	est.	se.	est.	se.	est.	se.
Autoregressive effect										
$\exp(\gamma_0)$	0.311	0.035	0.286	0.040	0.286	0.037	0.252	0.041	0.253	0.041
$\exp(\gamma)$	0.617	0.077	0.958	0.136	0.750	0.102	0.760	0.129	0.754	0.127
Network effect										
K :										
$\exp(\phi_0)$	0.477	0.070					0.503	0.144	0.510	0.073
$\exp(\phi)$	0.885	0.136					0.821	0.169	0.827	0.127
K^\top :										
$\exp(\phi_0)$			0.115	0.022			0.062	0.036	0.064	0.023
$\exp(\phi)$			0.374	0.080			0.148	0.218	0.166	0.181
KK^\top :										
$\exp(\phi_0)$					0.261	0.035	0.000	0.085		
$\exp(\phi)$					0.570	0.082	0.284	0.000		
Testing-related endemic effect										
pos-to-test	-1.37	0.47	-1.09	0.28	-1.46	0.41	-1.46	0.48	-1.46	0.49
lag test	1.85	0.31	1.77	0.23	1.88	0.28	1.91	0.32	1.91	0.32
pseudo- R^2	82.09%		82.00%		82.02%		82.12%		82.12%	

est. w.o. control

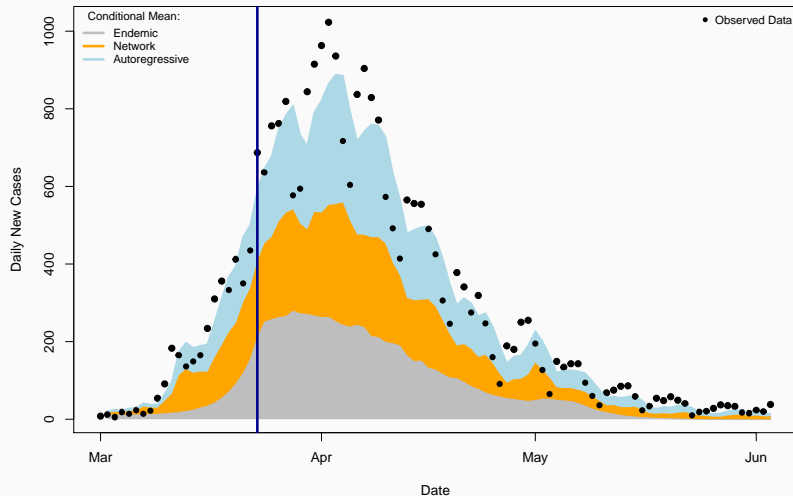
Model Estimation

Value	(K)		(K^\top)		(KK^\top)		all		(K, K^\top)	
	est.	se.	est.	se.	est.	se.	est.	se.	est.	se.
Autoregressive effect										
$\exp(\gamma_0)$	0.311	0.035	0.286	0.040	0.286	0.037	0.252	0.041	0.253	0.041
$\exp(\gamma)$	0.617	0.077	0.958	0.136	0.750	0.102	0.760	0.129	0.754	0.127
Network effect										
K:										
$\exp(\phi_0)$	0.477	0.070					0.503	0.144	0.510	0.073
$\exp(\phi)$	0.885	0.136					0.821	0.169	0.827	0.127
K^\top :										
$\exp(\phi_0)$			0.115	0.022			0.062	0.036	0.064	0.023
$\exp(\phi)$			0.374	0.080			0.148	0.218	0.166	0.181
KK^\top :										
$\exp(\phi_0)$					0.261	0.035	0.000	0.085		
$\exp(\phi)$					0.570	0.082	0.284	0.000		
Testing-related endemic effect										
pos-to-test	-1.37	0.47	-1.09	0.28	-1.46	0.41	-1.46	0.48	-1.46	0.49
lag test	1.85	0.31	1.77	0.23	1.88	0.28	1.91	0.32	1.91	0.32
pseudo- R^2	82.09%		82.00%		82.02%		82.12%		82.12%	

est. w.o. control

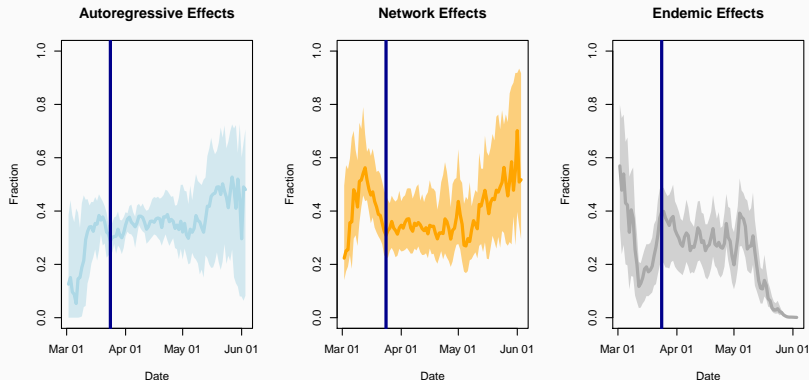
Decomposing the Conditional Mean

Conditional mean of disease cases: $\mu_t = \mu_t^{AR} + \mu_t^{NE} + \mu_t^{EN}$



Decomposing the Conditional Mean cont'd

Fraction of conditional mean explained: $\frac{\mu_{i,t}^{AR}}{\mu_{i,t}}, \frac{\mu_{i,t}^{NE}}{\mu_{i,t}}, \frac{\mu_{i,t}^{EN}}{\mu_{i,t}}$



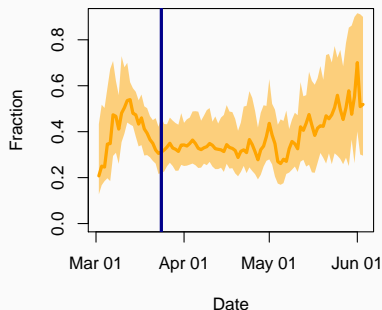
Bands represents 10- and 90-percent quantiles across 32 locations

Note: pre (post) lockdown average NE share: 45% (40%)

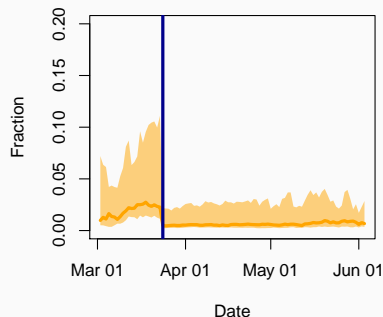
Decomposing the Conditional Mean

Fraction of conditional mean explained by each of the three networks: $\frac{\mu_{i,t}^{NE}}{\mu_{i,t}}(W^{(g)})$

Network Effects: K



Network Effects: K^T



Bands represents 10- and 90-percent quantiles across 32 locations

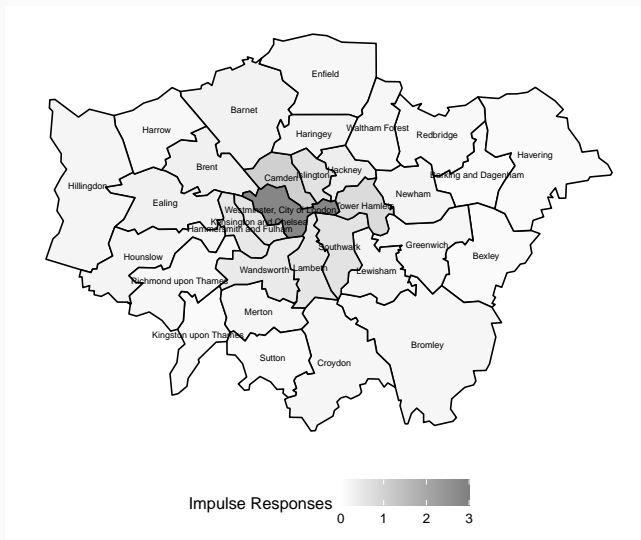
Network Impulse Responses

- “Shocks” to $y_{i,t}$ propagate through the network
- Let $\epsilon_{i,t} = y_{i,t} - \mu_{i,t}$
- We have the s -periods Network Impulse-Response Function:

$$NIRF_i(s) = \sum_{j=1}^N \frac{\partial \mathbb{E}_t[y_{j,t+s}]}{\partial \epsilon_{i,t}}$$

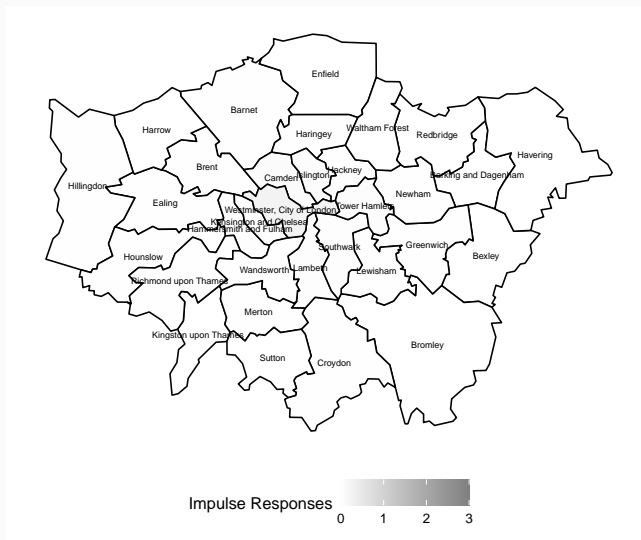
- Can be computed in closed form

Network Impulse Responses: Pre-lockdown, ($s = 7$ days)



Max N-IRF: 2.8

Network Impulse Responses: Post-lockdown, ($s = 7$ days)



Max N-IRF: 0.3

Epidemic Parameters

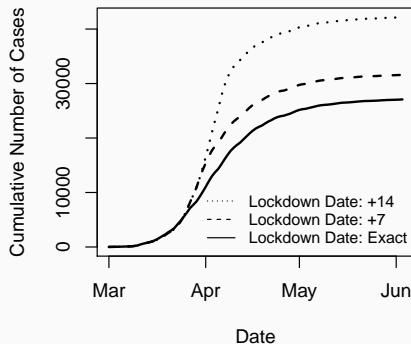
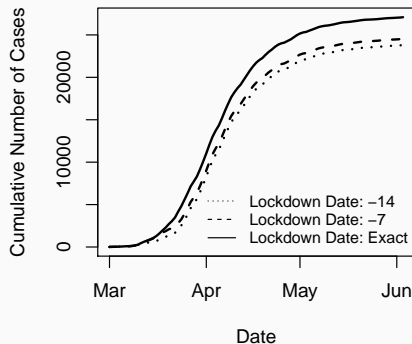
- The dominant (largest) eigenvalue σ_{\max} of $A(D_t) = \gamma(D_t)I + \sum_{g=1}^G \phi^{(g)}(D_t)W^{(g)}$ is key for epidemic dynamics since: $\mu_t \propto A(D_t)x_t$
- The disease epidemic R_0 (basic reproduction number) can be approximated as

$$R_0 \leq \sum_{\ell=0}^L \hat{v}(\ell) \sigma_{\max}(\hat{A}).$$

Model \mathcal{G}	(1)		(2)		(3)		(1,2,3)		(1,2)	
Value	<i>est.</i>	<i>se.</i>	<i>est.</i>	<i>se.</i>	<i>est.</i>	<i>se.</i>	<i>est.</i>	<i>se.</i>	<i>est.</i>	<i>se.</i>
Before Lockdown Policy: $D_t = 0$										
σ_{\max}	0.840	0.590	0.511	0.106	0.639	0.287	0.853	0.533	0.766	0.537
R_0	1.467	1.030	1.577	0.327	1.531	0.688	1.391	0.876	1.369	0.953
After Lockdown Policy: $D_t = 1$										
σ_{\max}	0.510	0.369	0.270	0.038	0.301	0.123	0.501	0.398	0.422	0.392
R_0	0.891	0.644	0.832	0.118	0.722	0.296	0.818	0.649	0.754	0.701

Counterfactuals and Policy Experiments

Counterfactual: Different Lockdown Dates?



Each path is the median of 10,000 simulations

-14 \Rightarrow -3,283 (-12.1%)

-7 \Rightarrow -2,553 (-9.4%)

+7 \Rightarrow +4,514 (+16.7%)

+14 \Rightarrow +15,085 (+55.7%)

Optimal Lockdown Policies

Given the transition dynamics

$$\mu_t \propto Ax_{t-1}$$

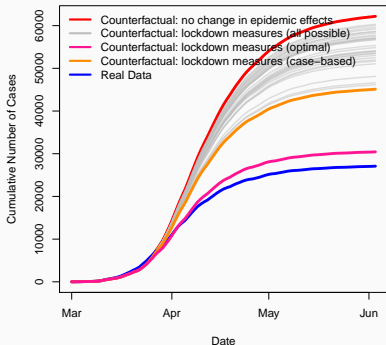
there are two naive candidate optimal lockdown policies:

- Locking down areas with the largest number of cases (“case-based” policy):
 - targeting on x_{t-1}
- Locking down areas showing large externalities (larger NIRF):
 - targeting on A

Partial Lockdowns: One Local Authority

Total lockdown

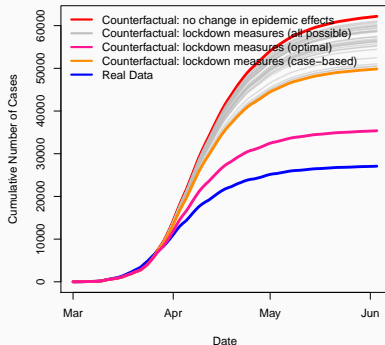
Locking down 1 regions on 2020-03-23



Optimal: Westminster, City of London

Partial lockdown

Locking down 1 regions on 2020-03-23



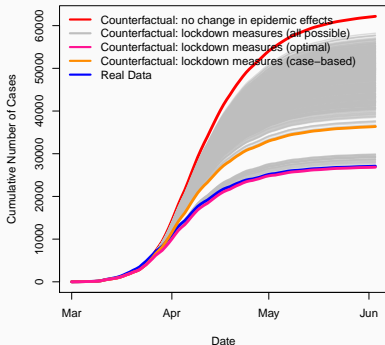
Optimal: Westminster, City of London

Note: Largest number of cases: Lambeth
Largest (integrated) NIRF: Westminster, City of London

Partial Lockdowns: Two Local Authorities

Total lockdown

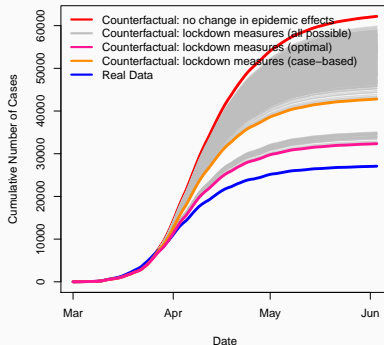
Locking down 2 regions on 2020-03-23



Optimal: Westminster, City of London/Southwark

Partial lockdown

Locking down 2 regions on 2020-03-23



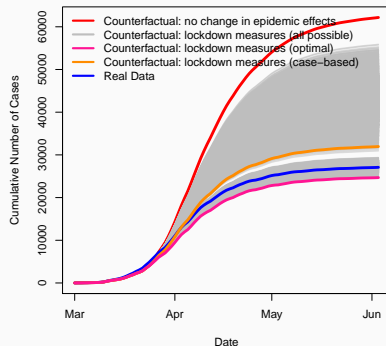
Optimal: Westminster, City of London/Southwark

Note: Largest number of cases: Lambeth/Southwark
Largest (integrated) NIRF: Westminster, City of London/Camden

Partial Lockdowns: Three Local Authorities

Total lockdown

Locking down 3 regions on 2020-03-23

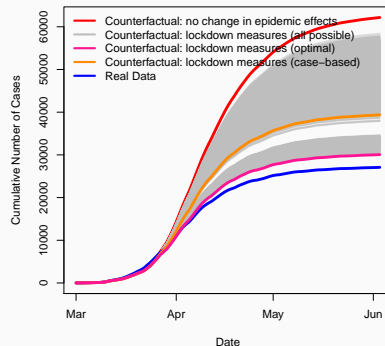


Optimal:

Westminster, City of London/Southwark/Lambeth

Partial lockdown

Locking down 3 regions on 2020-03-23



Optimal:

Westminster, City of London/Southwark/Lambeth

Note: Largest number of cases: Lambeth/Southwark/Brent
Largest (integrated) NIRF: Westminster, City of London/Camden/Tower Hamlets

Conclusion

We have embeded spatial dynamics in an empirical SIR model and found that:

1. Network externalities are first order
2. The national lockdown: *i*) reduced the spread by more than half, *ii*) was somehow late, and *iii*) further delay would have had extreme consequences.
3. “Case-based” lockdown policies appear suboptimal
4. Network externalities targeting, and partial lockdowns, seem very effective

Appendix

Estimation without controlling for test related variables

Model	(K)		(K [⊤])		(KK [⊤])		all		(K,K [⊤])	
	est.	se.	est.	se.	est.	se.	est.	se.	est.	se.
Autoregressive effect										
exp(γ_0)	0.381	0.033	0.456	0.034	0.424	0.032	0.365	0.036	0.344	0.038
exp(γ)	0.472	0.050	0.519	0.045	0.446	0.042	0.492	0.057	0.522	0.066
Network effect										
K:										
exp(ϕ_0)	0.563	0.066					0.496	0.091	0.592	0.071
exp(ϕ)	0.694	0.098					0.594	0.189	0.637	0.102
K [⊤] :										
exp(ϕ_0)			0.076	0.021			0.000	0.000	0.041	0.021
exp(ϕ)			0.598	0.175			0.000	0.000	0.203	0.424
KK [⊤] :										
exp(ϕ_0)					0.280	0.035	0.056	0.048		
exp(ϕ)					0.518	0.073	0.696	0.779		
pseudo-R ²	81.86%		80.90%		81.58%		81.89%		81.89%	