Inference under Superspreading: Determinants of SARS-CoV-2 Transmission in Germany

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Outline

About me

Paper presentation

Motivation

Model

Challenges

Results

Short CV

- Studies of Mathematics, Statistics, and Economics
- PhD in Econometrics (causal inference, computational statistics, decision-theory)
- Statistical consultant (impact evaluation and survey research for World Bank)
- PostDoc in Psychometrics (Bayesian, longitudinal data, measurement, epidemiology)
- Visiting Professor in Econometrics

Experience

statistical programming

structural estimation

causal inference

survey measurements

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Inference under superspreading

Schmidt, P. W. (2024). Inference under superspreading: Determinants of SARS-CoV-2 transmission in Germany.

Statistics in Medicine, 43(10):1933-1954

Goal: Understand determinants of SARS-CoV-2 transmission

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- Method: Structural model on surveillance data with Bayesian inference
 - 1. Estimate infection dynamic based on cases
 - 2. Explain infection dynamic with covariates

- Seminal contributions that estimate effects of interventions:
 - Flaxman et al. (Nature, 2020)
 - ► Brauner et al. (Science, 2021)

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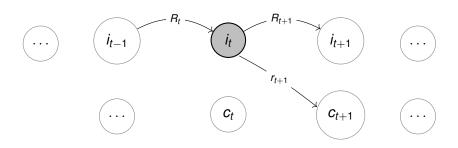
Is there anything to add?

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Is there anything to add?

- \times reporting delay \rightarrow symptom onset data
- \times superspreading \rightarrow model uncertainty
- × confounding of determinants → more covariates

- simplified: generation time and incubation fixed
- $ightharpoonup i_t$ infections at time t (unobserved)
- c_t cases with symptom onset at time t (observed)
- $ightharpoonup R_t$ reproductive number at time t
- $ightharpoonup r_t$ ascertainment rate at time t



► Transmission:

$$i_t \sim NB(\underline{i_{t-1}R_t},\underline{i_{t-1}}\Psi).$$

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Effect model:

$$R_t = R_0 \prod_{j=1}^J (1 + \beta_j x_{j,t}),$$

with covariates x_t .

Challenge: Superspreading

► Transmission:

$$i_t \sim NB(i_{t-1}R_t, i_{t-1}\Psi).$$

Challenge: Superspreading

Transmission:

$$i_t \sim NB(i_{t-1}R_t, i_{t-1}\Psi).$$

 Consistent with individual model of superspreading (Lloyd-Smith et al., 2005), where secondary infections are given by

$$NB(R_t, \Psi)$$
.

Assumption: Secondary infections independent given R_t.

Challenge: Superspreading

► Transmission model $i_t \sim NB(i_{t-1}R_t, i_{t-1}\Psi)$ is different from $i_t \sim NB(i_{t-1}R_t, \Psi)$.

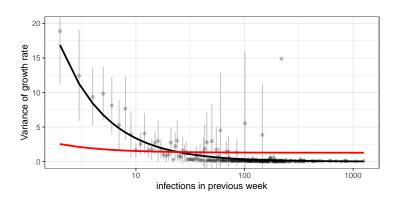


Figure: Mean estimate of empirical variance of weekly growth rates.

Challenge: Confounding

Previous studies use only non-pharmaceutical interventions (Brauner et al., 2021; Flaxman et al., 2020).

Challenge: Confounding

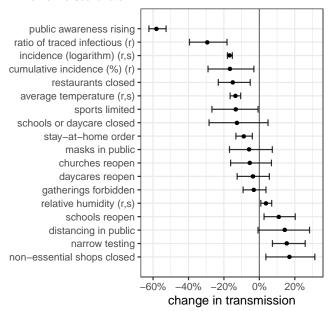
Previous studies use only non-pharmaceutical interventions (Brauner et al., 2021; Flaxman et al., 2020).

- This work further includes:
 - temperature, humidity
 - information on incidence
 - ratio of traced infectious
 - public awareness rising

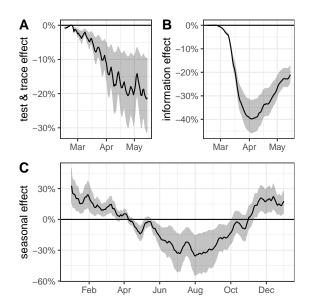
Data and Estimation

- German Covid-19 surveillance data by RKI with date of symptom onset
- DWD daily weather information
- Daily location-specific policy interventions from state legislative orders
- Estimation is based on cases in the 111 most impacted counties
- MCMC sampling implemented with JAGS (Plummer, 2019)

Intervention/covariate



Results: Total effects



Difficult task and many pitfalls remain unmentioned here.

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▶ Data on symptom onset can improve inference.

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Individual superspreading can be incorporated in models on aggregated cases.

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Data on symptom onset can improve inference.

Individual superspreading can be incorporated in models on aggregated cases.

Transmission driven by policy interventions, seasonality, and behavioral adaptation to local risk.

Conclusions of a non-epidemiologist

- Some opinions are supported by the data
 - ✓ Public awareness rising had impact
 - Behavior adapts to virus spread irrespective of policy interventions
 - Seasonal variation important driver
 - √ Test and trace had relatively strong impact

- Some opinions are not supported by the data
 - × Distancing/curfew/mask mandates had strong impact
 - X Only interventions cause economic and social costs

References I

- Brauner, J. M., Mindermann, S., Sharma, M., Johnston, D., Salvatier, J., Gavenčiak, T., Stephenson, A. B., Leech, G., Altman, G., Mikulik, V., et al. (2021). Inferring the effectiveness of government interventions against covid-19. *Science*, 371(6531).
- Flaxman, S., Mishra, S., Gandy, A., Unwin, H. J. T., Mellan, T. A., Coupland, H., Whittaker, C., Zhu, H., Berah, T., Eaton, J. W., et al. (2020). Estimating the effects of non-pharmaceutical interventions on COVID-19 in Europe. *Nature*, 584(7820):257–261.
- Lloyd-Smith, J. O., Schreiber, S. J., Kopp, P. E., and Getz, W. M. (2005). Superspreading and the effect of individual variation on disease emergence. *Nature*, 438(7066):355–359.
- Plummer, M. (2019). *rjags: Bayesian Graphical Models using MCMC*. R package version 4-10.



References II

Schmidt, P. W. (2024). Inference under superspreading: Determinants of SARS-CoV-2 transmission in Germany. *Statistics in Medicine*, 43(10):1933–1954.