

Demonstrates the utility of **network model frameworks** to capture heterogeneity of demographic attributes across worker roles and the individual nature of non-pharmaceutical interventions.

1. Motivation & aims

Globally, many countries have employed social distancing measures and nonpharmaceutical interventions (NPIs) to curb the spread of SARS-CoV-2 [1]. As part of a collective effort to protect public health by disrupting viral transmission, businesses also need to act appropriately by taking all reasonable measures to **minimise exposure** to coronavirus in workplaces and premises open to the public [2]. Adjustments in working practices can result in changes to the amount, duration, and/or proximity of interactions, thereby **altering the** dynamics of viral spread.

Study objectives:

- . Parameterise an **individual-based** network model of workers, stratified into work sectors, using a **data-driven** approach;
- ii.Study epidemic spread of SARS-CoV-2 amongst a population of workers and analyse the impact of **NPIs** targeted towards working practices.

2. Network model description

- Network model construction with nodes representing workers and connections representing contacts that can result in disease transmission in distinct settings: workplace, household and social (Fig. 1).
- Ran a SARS-CoV-2 outbreak upon the network, with transmission scaled according to setting and symptomatic status.

Simulation overview: •Population & time horizon: 10,000 workers, 365 days.

- •Simulation count: 1000 runs
- 50 distinct network realisations
- 20 runs per network realisation
- Initial disease state conditions:
- Ten individuals in an infectious state.
- All other individuals began in a susceptible state.
- •Default worker pattern: Assumed all workers had the same working pattern of five days at the workplace (Monday-Friday) and two days off (Saturday and Sunday).
- •Intervention implementation: Assumed that all NPIs implemented from day 15, including isolation and test-and-trace (default assumption of 70% adherent).



A network modelling approach to assess non-pharmaceutical disease controls against SARS-CoV-2 in a worker population Zeeman Institute: SBIDER, School of Life Sciences & Mathematics Institute, University of Warwick, UK. Joint UNIversities Pandemic and Epidemiological Research, https://maths.org/juniper/.





5. Results: Adherence to test, trace & isolation

 Absence of sufficient adherence to non-pharmaceutical interventions increases the chance of SARS-CoV-2 spreading widely in the population (Fig. 6).



6. Possible model developments

- Augment model with age structure.
- Inclusion of part time workers.
- levels of cough and fever are high due to non-COVID-19 causes.
- parameterise the model framework.

The Social Contact Survey data are available from https://wrap.warwick.ac.uk/54273/. The code repository for the study is available at: https://github.com/EdMHill/covid19 worker network model. EH, BA, MK, LD and MT were supported by the Medical Research Council through the COVID-19 Rapid Response Rolling Call [grant number MR/V009761/1]; MK, LD and MT were supported by the Engineering and Physical Sciences Research Council through the MathSys CDT [grant number EP/S022244/1] and by UKRI through the JUNIPER modelling consortium [grant number MR/V038613/1]. The funders had no role in the \study design, data collection or analysis.

[1] Hale T, Angrist N, Cameron-Blake E, Hallas L, Kira B, Majumdar S, et al. Oxford COVID-19 Government Response Tracker, Blavatnik School of Government.; 2020. URL: www.bsg.ox.ac.uk/covidtracker. [2] UK Government. Working safely during coronavirus (COVID-19); 2020. URL: https://bit.ly/38hCjwG [3] Danon L, Read JM, Keeling MJ, House TA, Vernon MC. Social contact study [Dataset]; 2009. Available from: https://wrap.warwick.ac.uk/54273/. [4] Danon L, House TA, Read JM, Keeling MJ. Social encounter networks: collective properties and disease transmission. J R Soc Interface. 2012;**9**(76):2826–2833. [5] Danon L, Read JM, House TA, Vernon MC, Keeling MJ. Social encounter networks: characterizing Great Britain. Proc R Soc B Biol Sci. 2013;**280**(1765):20131037.





Fig. 6: Case and isolation summary statistics under differing levels of adherence to test, trace and isolate measures.



• Allow for clustering of individuals within individual workplaces.

• Explore sensitivity to alternative epidemiological and intervention assumptions, e.g. presence of other respiratory infections and impact on test capacity when

• Application to **other countries**, given availability of associated data to

Acknowledgements

References