





Testing and other interventions in social care settings

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Dept. of Mathematics, UK Health Security Agency, PROTECT and JUNIPER June 2022



mediar

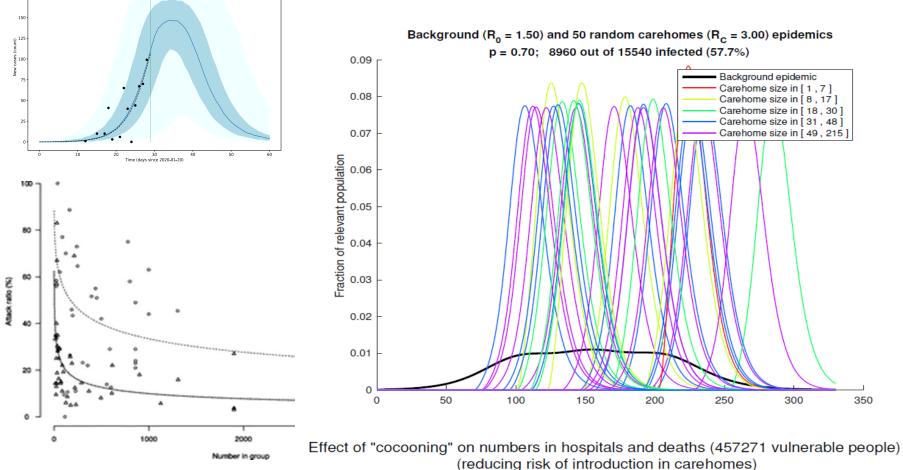
deterministic mode
 data
 2.5 - 97.5 quantile

25 - 75 quantile

200

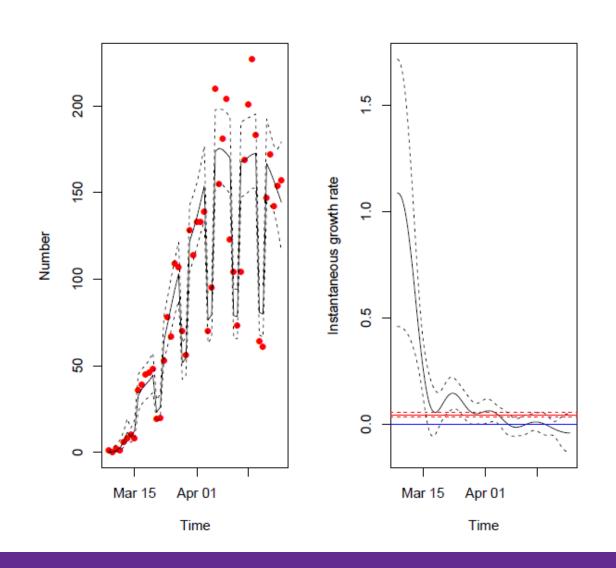
Enclosed Societies

- What did we know before?
 - Literature review for Flu.





Reported outbreaks over time

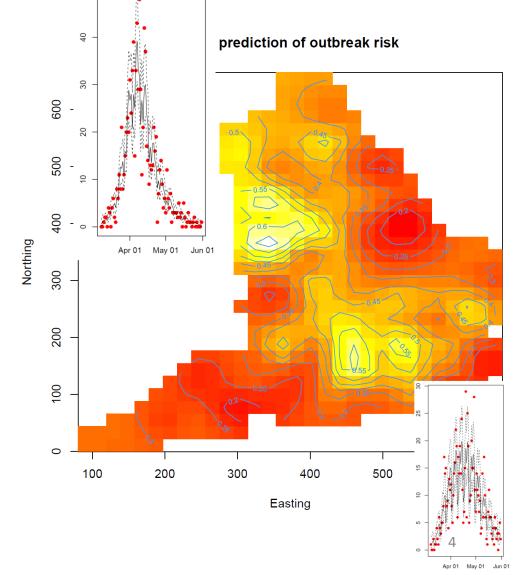


- There are
 N=15517 care
 homes in
 England
- Steady state of 190 outbreaks per day possible.
 - With a 5 day generation time and 4 generations of disease and 14 day observation then...
- P=0.41



Methods/Results - Spatial distribution

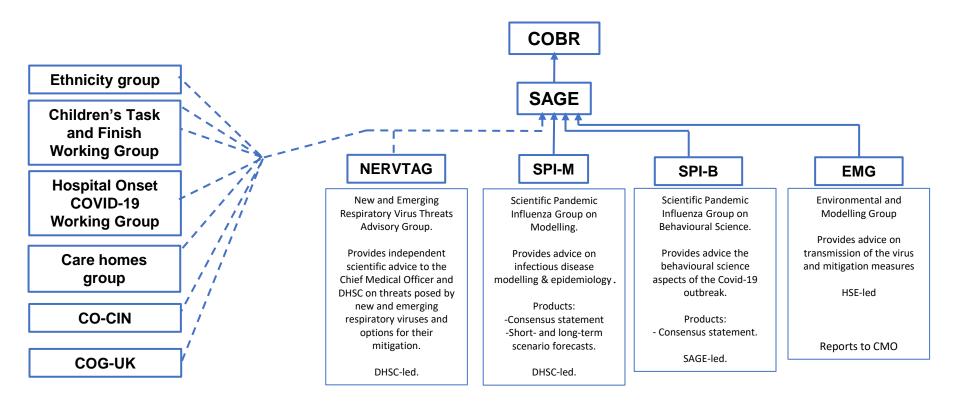
- Presence/absence of outbreaks
- Aims to support decision making of DPHs
 - Should they test care homes near
 - current outbreaks
 - Or randomly in space.
- 32% National average (at time)
- Use GAM (Gaussian Process) with binomial family





Groups feeding into SAGE

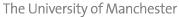
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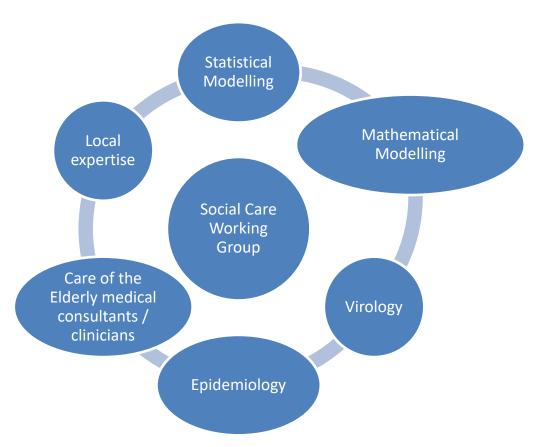






Sage Social Care Working Group – Core group & Expertise



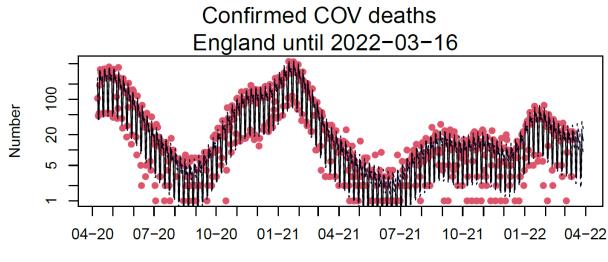


- April 2020: Group started as 'Sage Care Home Working Group'
- September 2020: Wider remit, clearly defined core members & new Terms of Reference.

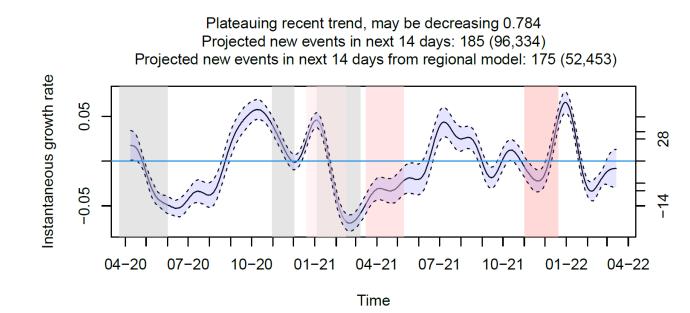


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Forecasting



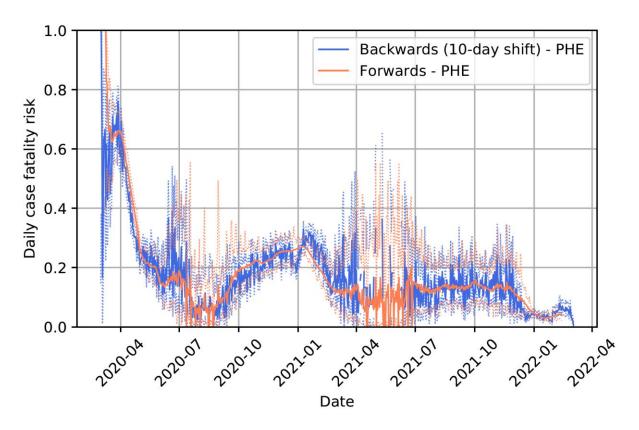
Time





Case fatality ratio within care homes

- Uses CQC death notifications and P1 and P2 positive tests matched to care home location and age
- Allows for delay from positive test to death
- Gives 14% CFR in 'stable' data period with appropriate age filters
- Highly variable

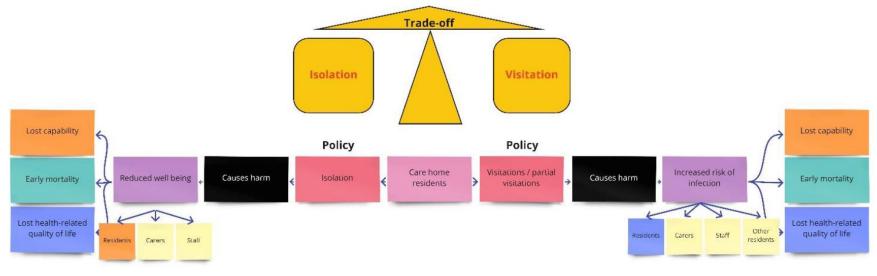






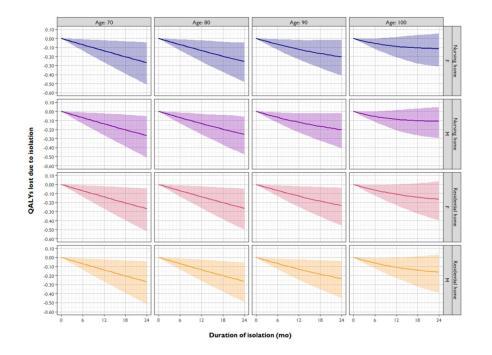
How do we create a visitor policy that is safe but enables residents to see their family? <u>Alexander Thompson</u>

- Potential harm caused by isolation
- Homes in different areas of high / low prevalence
- Each individual and family member may have a different view on safety vs quality of life.





Quality adjusted Life Years...

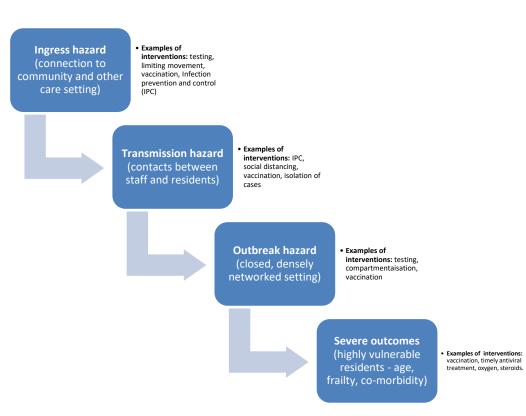


But does the QALY capture necessary impacts on wider wellbeing and health (ASCOT).

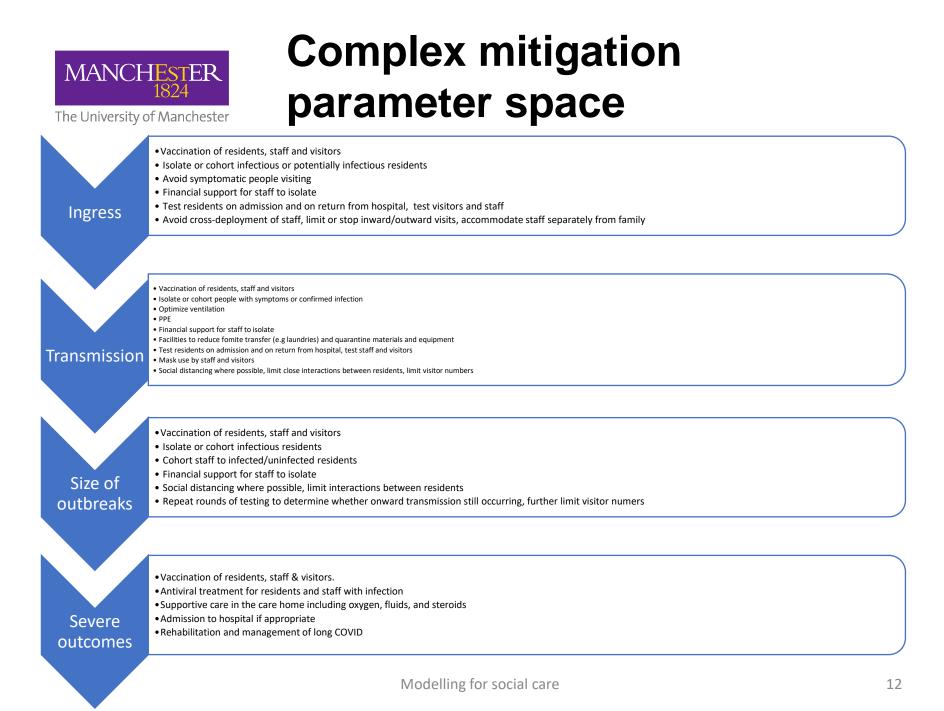
Need to integrate with epidemic models



Hazard profiling

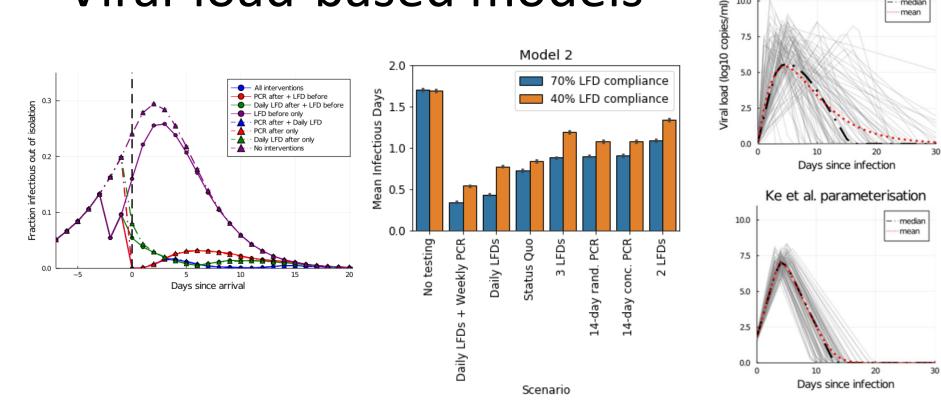


Modelling for social care





Viral-load-based models



Model 2 (Kissler et al. data)

median

mean

10.0

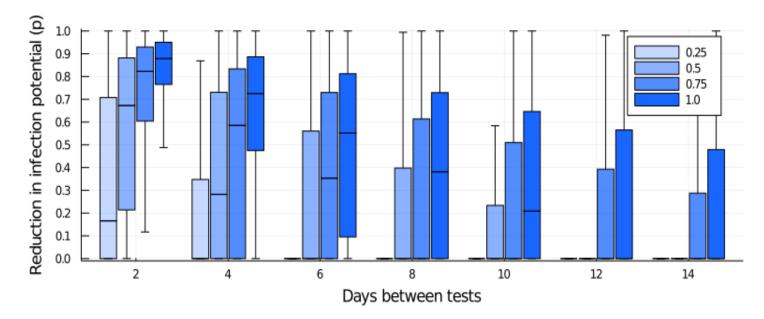
Ke, R et al. 'Daily sampling of early SARS-CoV-2 infection reveals substantial heterogeneity in infectiousness', Medrxiv 2021. Kissler, S. M. et al PLOS Biology 19(7), e3001333. 2021







- Behaviour and acceptance is critical to efficacy for testing.
- Sensitivity and specificity vary over time

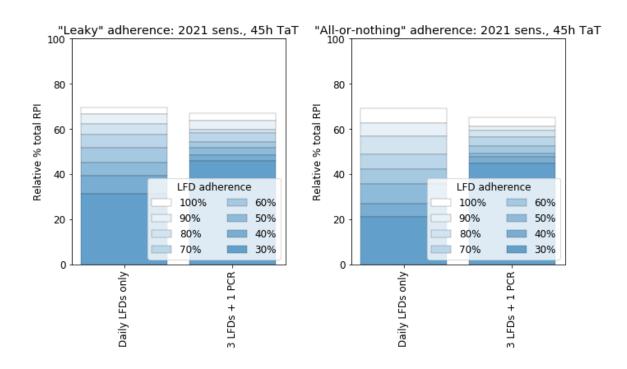




The role of adherence

Type of adherence matters: Leaky: Everyone takes next test with same probability AoN: Fraction of people do all tests, fraction do none

Strategies with high frequency are most affected by this difference







patient No.1st 63410

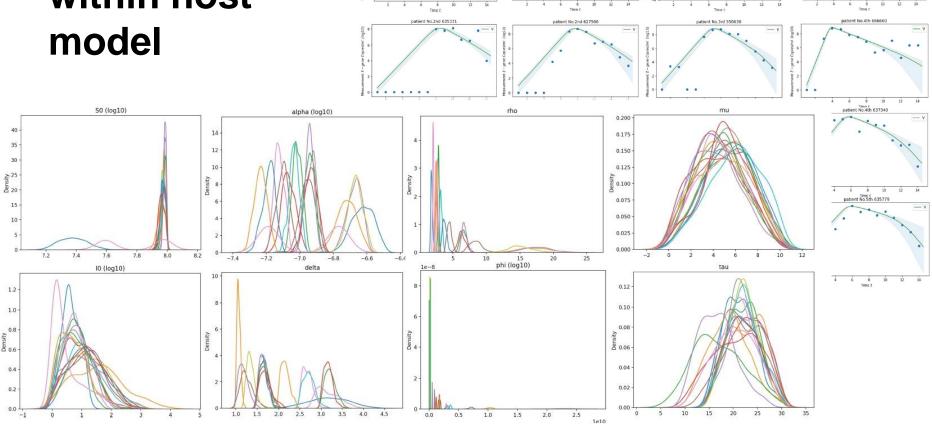


atient No 2nd 635729

credit interval for simplified model.

patient No.1st 63549

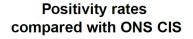
Results of within host model

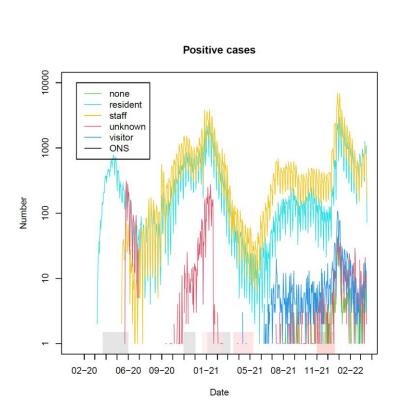


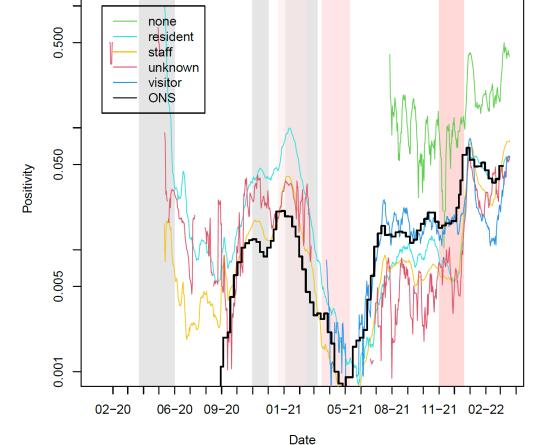


Correlation with ONS CIS

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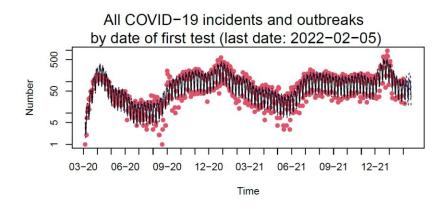


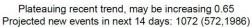


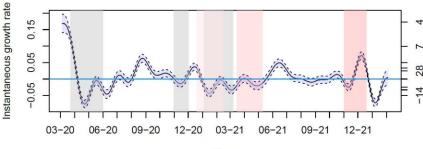


Transmission

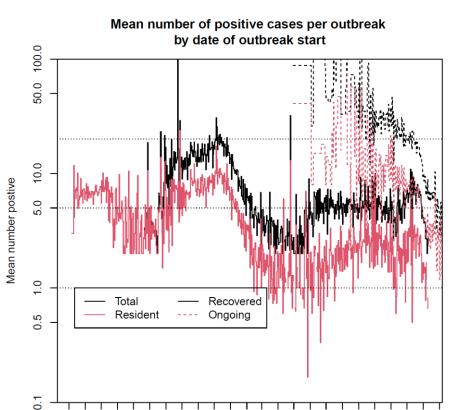
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12-20

03-21

dailymerge\$date

06-21

09-21

Modelling for social care

03-20

06

20

09-20

12-21



Summary

- Data was weak...
- ... got better but not perfect. Currently few negatives reported.
- Hard to evaluate specific interventions
- Eager to build in future research and look at acceptability
- Testing residents may not be viable
- Home care and other settings critical.
- Staff data linked to workplaces adds value.
- Timescales for reporting are short!

MANCHESTER Acknowledgements

- Data: UKHSA, DHSC and CQC
- University of Manchester COVID-19 Modelling group (co-leads: Thomas House + Lorenzo Pellis; Luke Webb for spatial modelling; Chris Overton for CFR; Heather Riley extending model; Alex Thompson for health economics work; Carl Whitfield testing effiacy)
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- SAGE Care Home Sub-group (Steve Willner, secretariat; Eamonn O'Moore/Jenny Harries, co-chair)

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