

JUNE

open-source individual-based epidemics model

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Outline

- 1 introduction: modelling epidemics
- 2 a first individual-based model
- 3 introducing JUNE
- 4 results for the first wave in England
- 5 a spin-off: Cox's Bazaar Refugee Operation
- 6 conclusions & outlook



June Almeida

discovered Corona Virus (1964)

introduction: modelling epidemics

the begin of epidemiology

- start of epidemiology as a science in early 20's

Kermack-McKendrick models, three papers from 1927-1935: W.O.Kermack & A.G.McKendrick, "A contribution to the mathematical theory of epidemics", Proc. R. Soc. Lond. A115 (1927) 700-721
reprinted in:
Kermack, W. and McKendrick, A. "Contributions to the mathematical theory of epidemics – I-III", Bulletin of Mathematical Biology. 53 (1-2) (1991) 33-118



- side-remark: about as old as quantum physics
- mathematical models developed in aftermath of Spanish Flu

50,000,000 dead worldwide (estimated IFR: 10%),
675,000 in US; 228,000 in UK; 426,600 in Germany

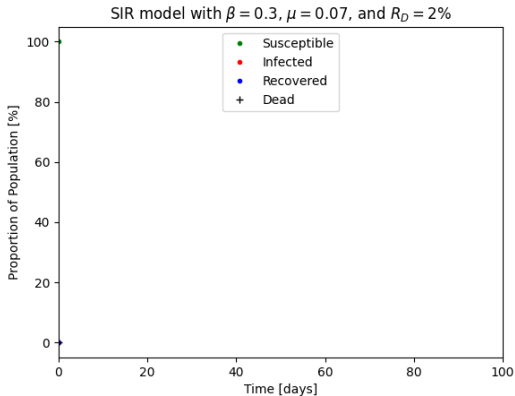
analytical models (SIR and friends)

- simple compartment models:
distribute population N in boxes according to infection status
- coupled rate equations for moving from one compartment to next
- simplest example: **S**usceptible-**I**nfected- **R**covered)

$$\begin{aligned}\frac{\partial S(t)}{\partial t} &= -\beta \frac{I(t)}{N} \cdot S(t) \\ \frac{\partial I(t)}{\partial t} &= +\beta \frac{I(t)}{N} \cdot S(t) - \mu I(t) \\ \frac{\partial R(t)}{\partial t} &= +\mu I(t) \\ N &= S(t) + I(t) + R(t) = \text{const.}\end{aligned}$$

- can introduce more compartments (“exposed”, “dead”, etc.)

example result for SIR model



improving SIR: social mixing

- add detail of population: age & gender
- this amounts to sub-boxes in one compartment, e.g. $S \rightarrow S = \sum_i S_i$
- modify interactions between boxes: social mixing matrices $\chi_{ij}^{(E)}$

count number of daily contacts of people i with people j in social environment E (home, work, school, ...)

- J.Mossong et al. (POLYMOD), PLoS Med 5 (2008), 3:e74
- P.Klepac et al. (BBC pandemics project),
<https://www.medrxiv.org/content/10.1101/2020.02.16.20023754v2>

- rate equations become, for example

$$\frac{\partial S_i(t)}{\partial t} = - \sum_E \beta^{(E)} \sum_j \chi_{ij}^{(E)} \frac{I_j(t)}{N_E} \cdot S_i(t),$$

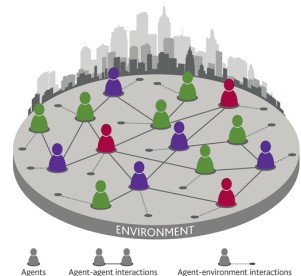
where $\beta^{(E)}$ parametrizes the infection probability in setting E

analytical models to simulation

- advantages of analytical models:
 - “just” differential equations, transparent dynamics
 - fast solutions/fits to existing developments
 - satisfactory capture of global trends
- obvious drawback of analytical models
 - population in “boxes” – insensitive to geography
 - deterministic models to describe statistical process
 - population densities only (“mean-field”):
outliers/super-spreader events/variations not captured
- ideal for short-term forecasting (often driven by machine learning),
not so good for understanding of inner dynamics
- solution: agent-based models

agent-based models: paradigm

- method for simulating complex systems
- individual agents represent actors:
 - individuals, organisations, etc.
 - interact with each other and environment according to fixed rules
- study impact of individual agent behaviour patterns on whole system
- core method of computational social sciences
frequent preconception: useful for qualitative analyses only,
impossible for quantitative real-world applications (too complex)
 - "mother of all ABM's":
Schelling, T.C., "*Dynamic Models of Segregation*", *Journal of Mathematical Sociology* 1 (1971) 143–186
 - some reviews:
Macy, M.W. and Willer, R., *Annual Review of Sociology* 28 (2002) 143–166;
MacNamee B. and Cunningham P., *International Journal of Intelligent Games and Simulation* 2(2003) 186–221;
Bruch E. and Atwell J. *Social Methods Research* 44 (2015) 186–221.



a simple example

(starting point of JUNE, I built it for my kids in March to explain how epidemics spread)

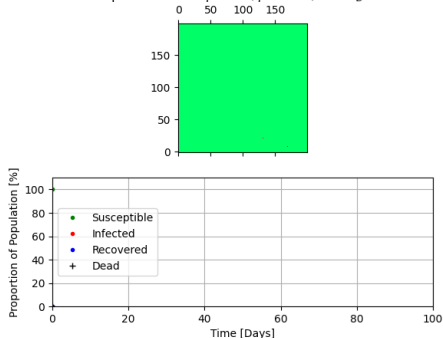
- agents fixed on grid (200×200)
- i infects susceptible s with

$$\mathcal{P}_{si} = \beta \exp [-(r_{si}^2 - r_0^2)/r_0^2]$$

with relative spatial distance r_{si}

- time to recovery or death distributed according to Gaussian
- observation(s):
 - similar to SIR solution
 - no herd immunity below 100%

A Simple Simulation $\beta = 0.5$, $\mu = 0.07$, and $R_D = 5\%$



a first individual-based model

agent-based models for epidemics: construction principle

- track disease progression through each individual
(thus populations become highly heterogeneous by health status during simulations),
- track contacts of individuals in social environments and geographies
- add explicit rules for disease transmission and impact
- agents are individuals: → **individual-based model**
- need to construct:

population & social environment
+
disease transmission & health impact

most famous example: Imperial College model

N.Ferguson et al., "*Strategies for containing an emerging influenza pandemic in Southeast Asia*", Nature 437 (2005) 209-214
N.Ferguson et al., "*Strategies for mitigating an influenza pandemic*", Nature 442 (2006) 448-452

- population model:
 - distribute population in age & sex according to national distributions
 - distribute population according to national population density
- social environments:
 - four relevant environments: household, company, school, other
 - household, company, and school sizes and composition distributed according to nation-wide distribution
 - "other" captures all contacts in the closer or wider community:
 - venues such as: shops, gyms, pubs, . . .
 - activities such as visiting friends & relatives, travel, . . .
 - distribute them according to a $1/(r + r_0)$ -law w.r.t. place of residence

- one underlying assumption:
transmission in equal parts in household, work/school, and other

(I am not sure in how far they changed this)

- disease transmission probability in interval Δt :

$$\mathcal{P}_{si}(t, t + \Delta t) = 1 - \exp \left[-\psi_s \beta^{(E)} \frac{\sum_i I_i(t)}{N_E^{\eta_E}} \Delta t \right]$$

- time-dependent infectiousness $I_i(t)$, and
- susceptibility ψ_s , and
- normalisation exponent η_E to capture different contact frequencies in environments E

- outcome of infection from continuously updated public health data

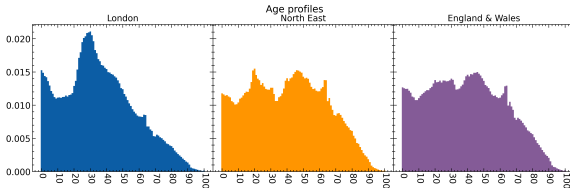
(they started as soon as the first data came out of Wuhan)

JUNE

motivation: why granularity matters

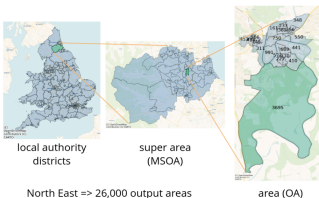
- impact of COVID=19 highly age-dependent
→ **need geographical granularity for regional planning**

(coincidence: Durham hosts & maintains England & Wales census data of past decades)



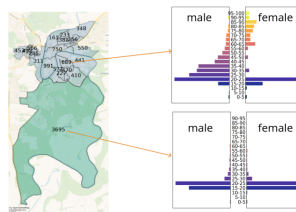
inputs: demographics

- last census (2011)
(data freely available from Office for National Statistics)
- hierarchical data structure



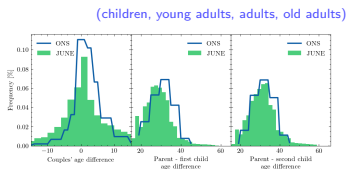
- OA's with ~ 250 residents, with similar characteristics

- build virtual population in OA: age, gender, ethnicity, deprivation index
- example: Durham



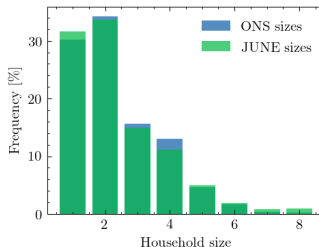
virtual households

- correct compositions important: primary place for infections
- household composition in 20 categories at OA level
- further test: interplay with social mixing
- example: North-East England



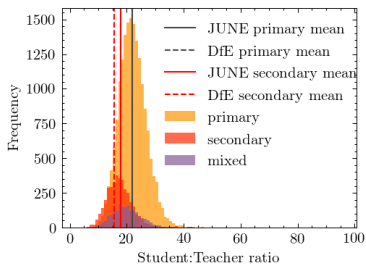
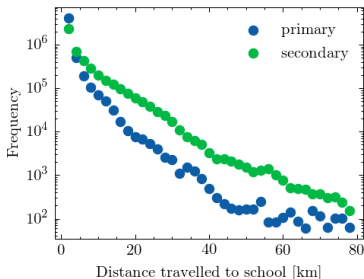
- also: communal facilities

(carehomes)



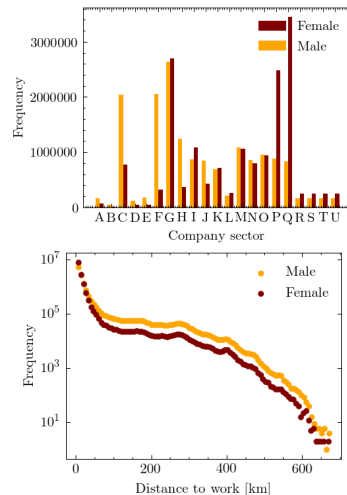
virtual schools

- information about schools: age range of children and locations
- send kids to nearest age-appropriate structure
- could modulate this with school sizes, if necessary



work & virtual companies

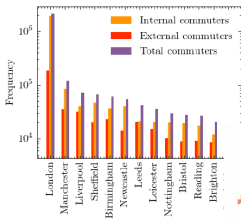
- workers and companies in ~ 20 macro-sectors at MSOA level
- know age/sex distribution in sectors nation wide
- distribute workers over companies (we know their sizes in bins)
(construct a big origin-destination matrix & optimise)
- information about commute mode: public vs. private



commuting patterns

- distinguish commuting into and within metropolitan areas

(external vs. internal commuters)

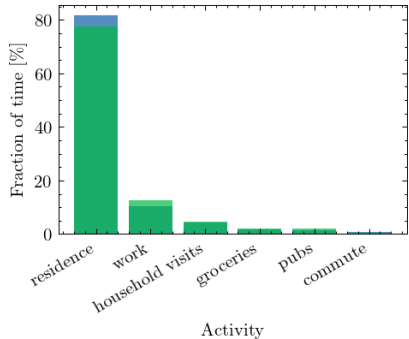
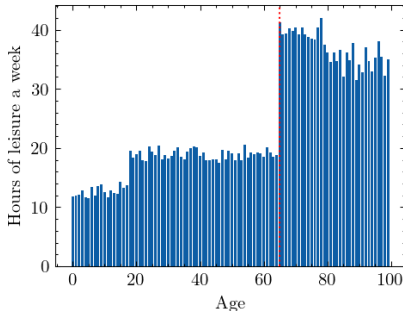


simulating daily structure

- JUNE allows flexible daily routines, separation of weekday/weekend
- time spent on activities known from ONS surveys

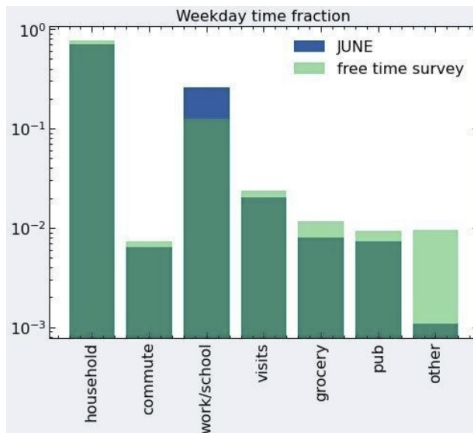
(this changes under lock-down)

- can translate into age-dependent probabilities for activities



validation: average time budget

- compare our simulation with data from ONS
- average time spent per day in different activities



social mixing matrices

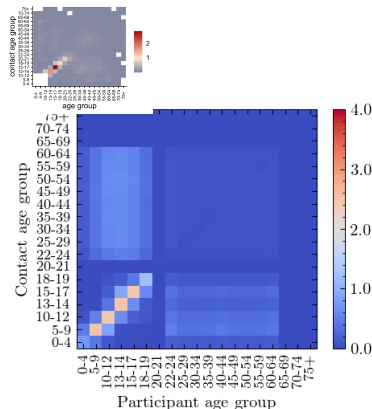
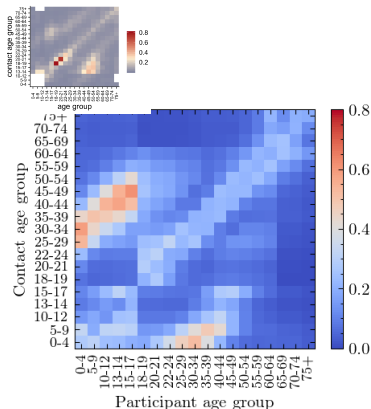
- move on to the environment for the interactions
- use social mixing matrices from POLYMOD and BBC Pandemics project
J.Mossong et al., PLoS Med 5(3) e74, <https://doi.org/10.1371/journal.pmed.0050074>;
P.Klepac et al., <https://www.medrxiv.org/content/10.1101/2020.02.16.20023754v2>
- denote number of contacts of person with age i with person of age j
- somewhat tricky format: averages over full population
→ have to normalise it to our fractured social environment
→ interplay with household sizes etc. (may have to be fitted?)

JUNE output

- example: household (left) & schools (right)
- broad agreement with input from surveys: interesting closure test

(BBC results as inlays)

(in JUNE contacts also depend on composition of environment)



disease transmission

- probabilistic process:

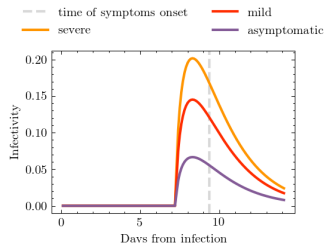
$$\mathcal{P}_{f|g}(t, t + \Delta t) = 1 - \exp \left[-\psi_s \beta_{si}^{(E,g)} \int_t^{t+\Delta t} dt' \sum_{i \in g} I_i(t') \right],$$

- ψ_s is susceptibility (reduced for children)
- $\beta_{si}^{(E,g)}$ is constant “closeness” in environment E , modulated by contact matrix:

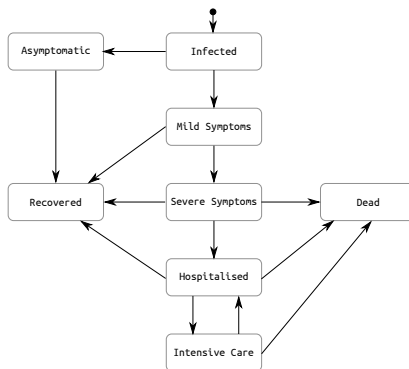
$$\beta_{si}^{(E,g)} = \beta^E \cdot \frac{\chi_{si}^{(g)}}{N_g}$$

with size of group N_g

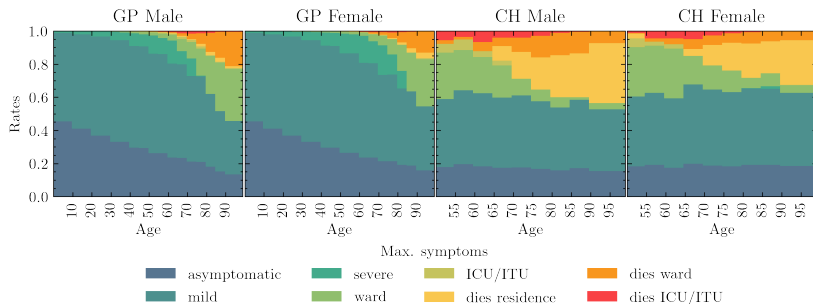
- $I_i(t')$ is infectiousness profile



outcomes of infection

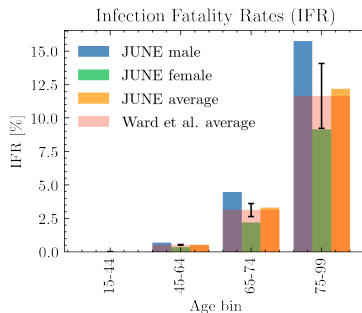
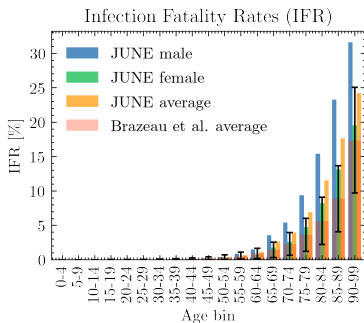


- this was a tiring data-mining exercise with inconsistent and often contradictory data
- extra difficulty: include care homes (CH) vs. general population (GP)



validation

- compare infection fatality rate with other codes

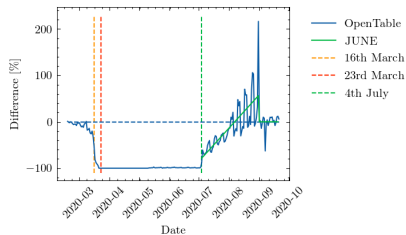
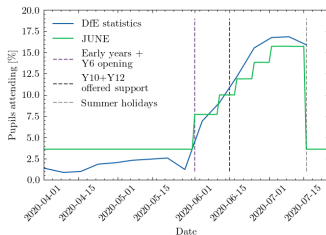


policies

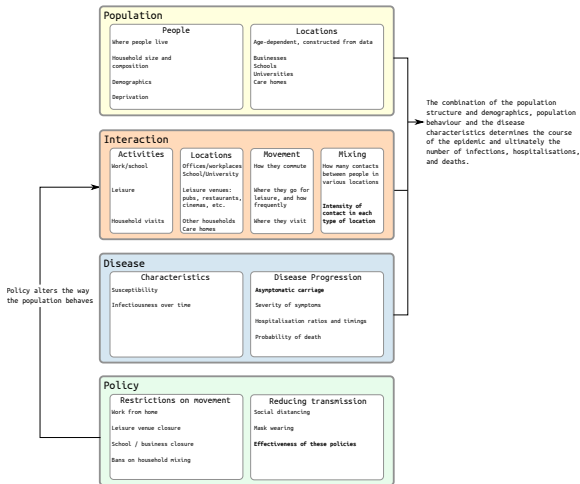
- below a list of relevant policies for results
- realised by modifications of β (e.g. social distancing) or by making activities unavailable

Date (dd/mm/yy)	Policy
12/03/20	case isolation at home
16/03/20	voluntary household quarantine, work from home, avoidance of leisure activities, encourage social distancing, stop all non-essential travel & contact, shielding of over 70's
20/03/20	closure of schools and universities
21/03/20	closure of leisure venues
11/05/20	multiple trips outside are allowed in England only
13/05/20	encourage to go back to work while distancing
01/06/20	meeting in groups of up to 6 outside allowed, shielding of over 70s relaxed, school reopening for Early Year and Year 6 students
13/06/20	'support bubbles' allowed
15/06/20	school reopening for Year 10 and 12 students for face-to-face support
04/07/20	leisure venues allowed to reopen, household-to-household visits permitted along with overnight stays
24/07/20	Mask wearing compulsory in grocery stores
01/08/20	shielding is paused, 'Eat Out to Help Out' scheme introduced
31/08/20	'Eat Out to Help Out' scheme ends
01/09/20	schools and Universities allowed to reopen, 'Rule of 6'
14/10/20	Tiered local lock-down system introduced

validation



JUNE simulation content - summary

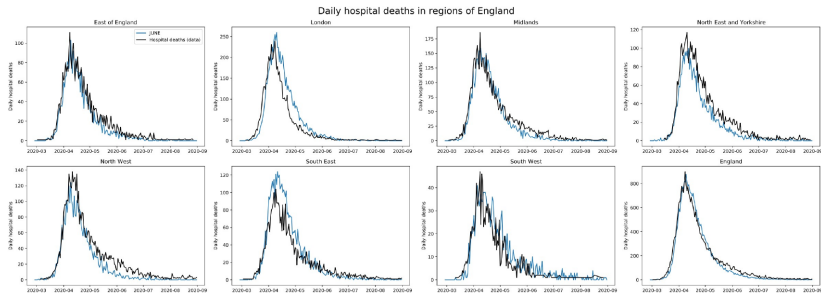


JUNE results for the first wave

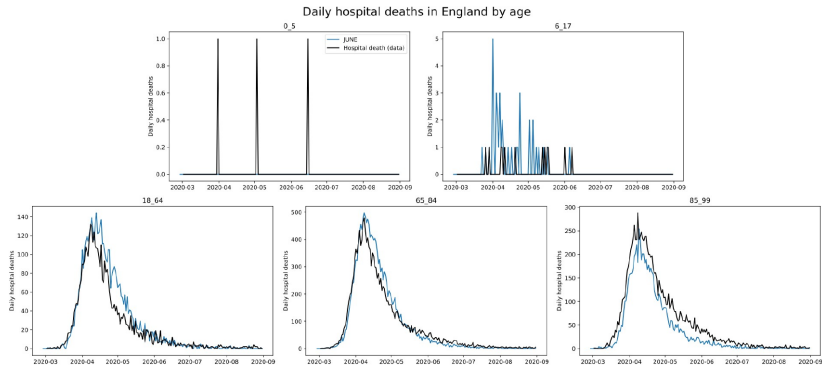
JUNE results

- below a selection of JUNE results for the 1st wave in England
- some aspects of the code changed in the meantime:
 - seeding of infections - must reorganise due to multiple CoVid streams
 - added gyms as leisure venues
 - take into account ethnicity in household compositions
 - added vaccination protocols
- refitting underway

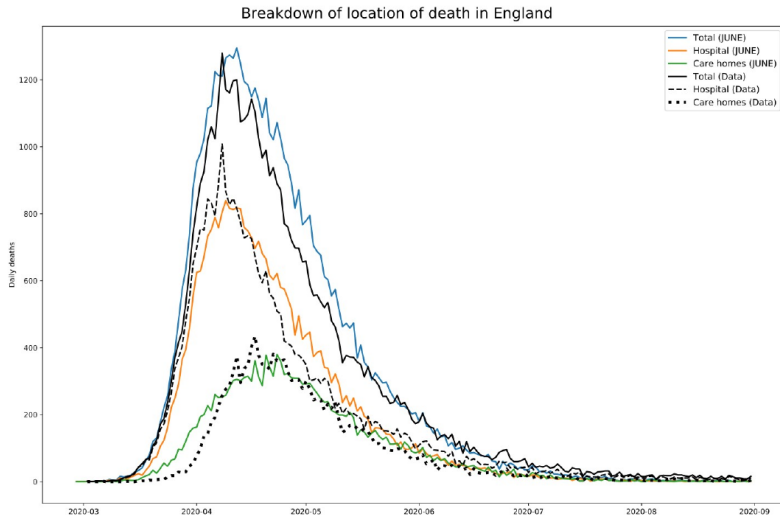
- 1st wave: deaths in hospitals - regional distribution



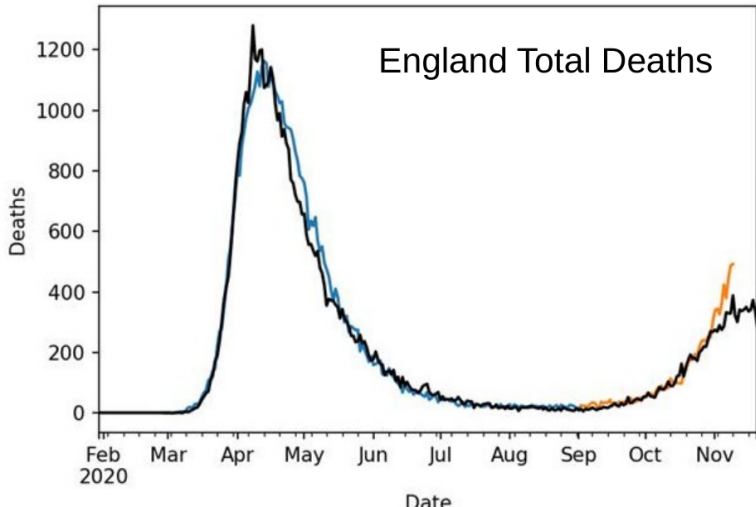
- 1st wave: deaths in hospitals - age distribution



- 1st wave: all deaths - distribution of location

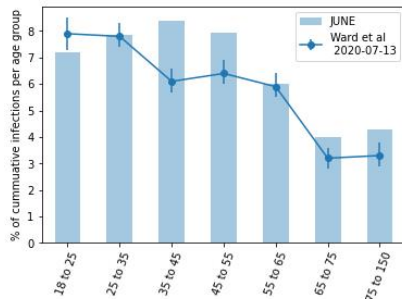


- extrapolation to second wave from September 1st
fitted parameters from first wave only!

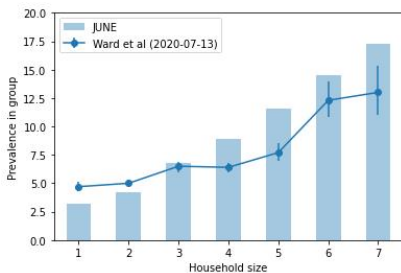


social imbalances

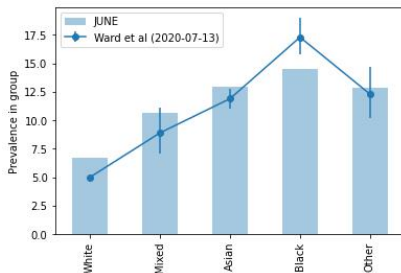
- look at cumulative infection rates until July 2020 in dependence on
 - age band
 - household size
 - ethnicity
- keep in mind: all imbalances only due to regional and sociological differences encoded in census data



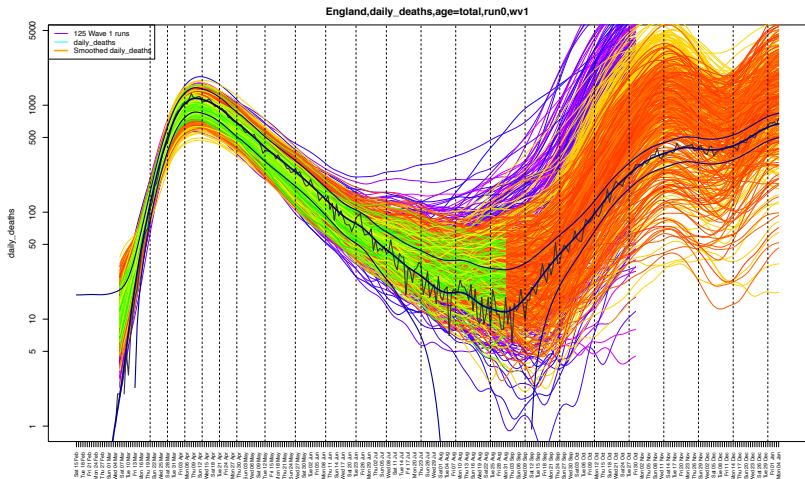
impact of household size



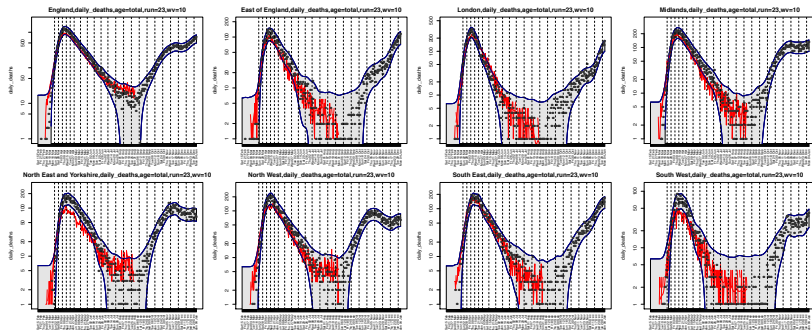
impact of ethnicity



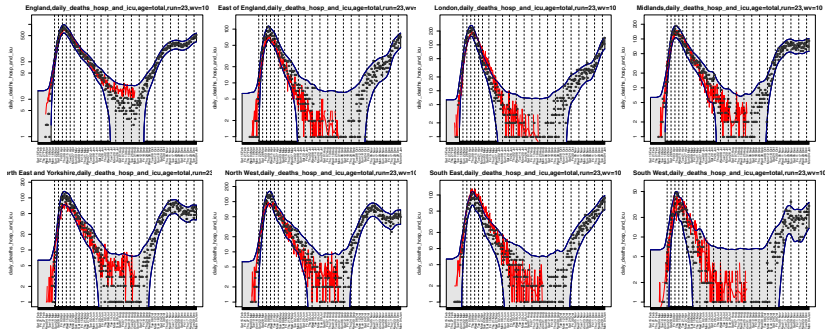
refitting snapshot



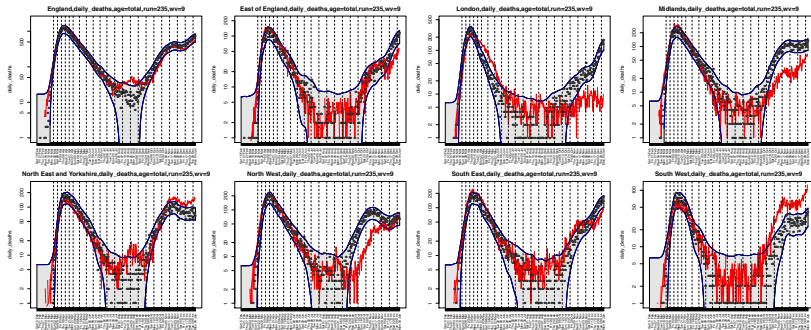
improved 1st wave: total deaths by region (example run)



improved 1st wave: deaths in hospital by region (example run)



first stab at 2nd wave (example run)



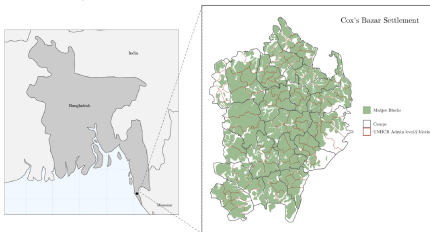
a spin-off:

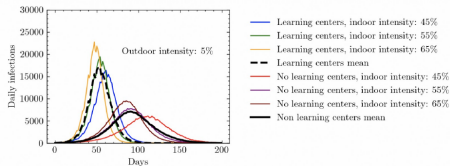
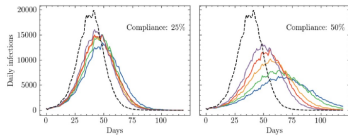
JUNE for Cox's Bazaar Refugee Operation

- one of our students (J.Bullock) has spent 6 months of placement with UN Global Pulse in New York
- ongoing contact, and he talked with them about our code
- as a result, three students collaborate with WHO, UN-Global Pulse, and IBM-MIT Watson AI lab to provide tools for scenario planning for Cox's Bazaar in Bangladesh

(huge refugee camp, Rohingya crisis in Myanmar)

- largest settlement in the world
- in some areas, the settlement is denser than New York City
- high risk of COVID transmission





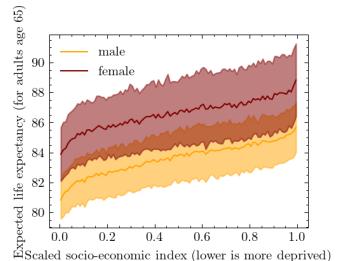
conclusions & outlook

summary

- epidemiology is an interesting field of science
- constructed an individual-based model with supreme granularity: demography, geography, sociology
- model informed operational planning of NHS:
 - early warning of second wave
 - understanding of transmission sociology
- code is highly flexible:
 - addition of new effects & policies relatively painless
 - adaptation to new environments: Cox's Bazaar
 - adaptation to Germany underway (M.Schott, U Mainz)
- challenge to widespread perception in computational sociology: more and better detail often helps

outlook

- continue decision support for NHS:
 - impact of mutations:
added Kent variant B.1.1.7
 - impact of vaccination protocols:
take-up, efficacy
 - new fits underway
(difficulty: Xmas mixing)
- follow-up studies on social imbalances:
ethnicity, IFR in dependence on
socio-economic deprivation
- use for historical studies:
Spanish Flu, Black Death



meet the JUNE team

- work started in March
- publication on medRxiv (<https://www.medrxiv.org/content/10.1101/2020.12.15.20248246v2>)
- very interdisciplinary research, programming and data crunching mainly driven by **PhD students** from Durham CDT

Joseph Bullock^{1,3}, **Carolina Cuesta-Lazaro**^{1,2}, **Arnau Quera-Bofarull**^{1,2},
Miguel de-Icaza-Lizaola^{1,2}, **Aidan Sedgewick**^{1,2}, **Henry Truong**^{1,3},
Aoife Curran^{1,2}, **Edward Elliott**^{1,2},
Richard Bower^{1,2}, Tristan Caulfield⁴, Kevin Fong⁴,
Ian Vernon⁵, Julian Willams⁶ & FK^{1,3}

- 1 Institute for Data Science Durham
- 2 Institute for Computational Cosmology Durham
- 3 Institute for Particle Physics Phenomenology Durham
- 4 University College London & NHS
- 5 Mathematical Sciences, Durham
- 6 Institute for Hazard, Risk & Resilience Durham