

HiDATA webinar:
Data Science in the Post-Covid World
11 May 2021



Data science in emergency



UiO : University of Oslo

@freeges

frigessi@medisin.uio.no

Norwegian strategy during Covid-19 emergency

- **Keep control of the spread of the virus.**
- Control means:
 - the number of patients is manageable for the health system
 - the number of cases allows efficient testing, isolation, contact tracing and quarantine.
- Local interventions allowed and encouraged.



New confirmed cases of Covid-19 in Norway

Seven-day rolling average of new cases (per 100k)

Major national restrictions announced





UiO : University of Oslo



NIPH COVID-19 modelling team



Birgitte Freiesleben de Blasio, Gunnar Øyvind Isaksson Rø, Alfonso Diz-Lois Palomares, Anja Brathen Kristoffersen ,
Francesco Di Ruscio, Solveig Engebretsen, Geir Storvik, Kenth Engø-Monsen.

Our tasks:

REQUESTS:

- Predict the number of covid-19 patients
- Predict number of patients requiring hospitalization, in particular ICU
- Estimate prevalence
- Estimate effective reproduction numbers

... at national and regional level

- National vaccination strategy
- National re-opening strategy

- 3 weeks
- 1 year

Data

- Total tested, total positive cases – excluding repeated tests
- Total imported positive cases
- Total new admitted patients to hospital
- Total new admitted to ICU, with and without ventilation
- Total deaths

All daily at municipal/hospital level.

- Basic data pipelines were not ready at start
- Overlapping registries not always coherent
- Seasonal and organisational irregularities

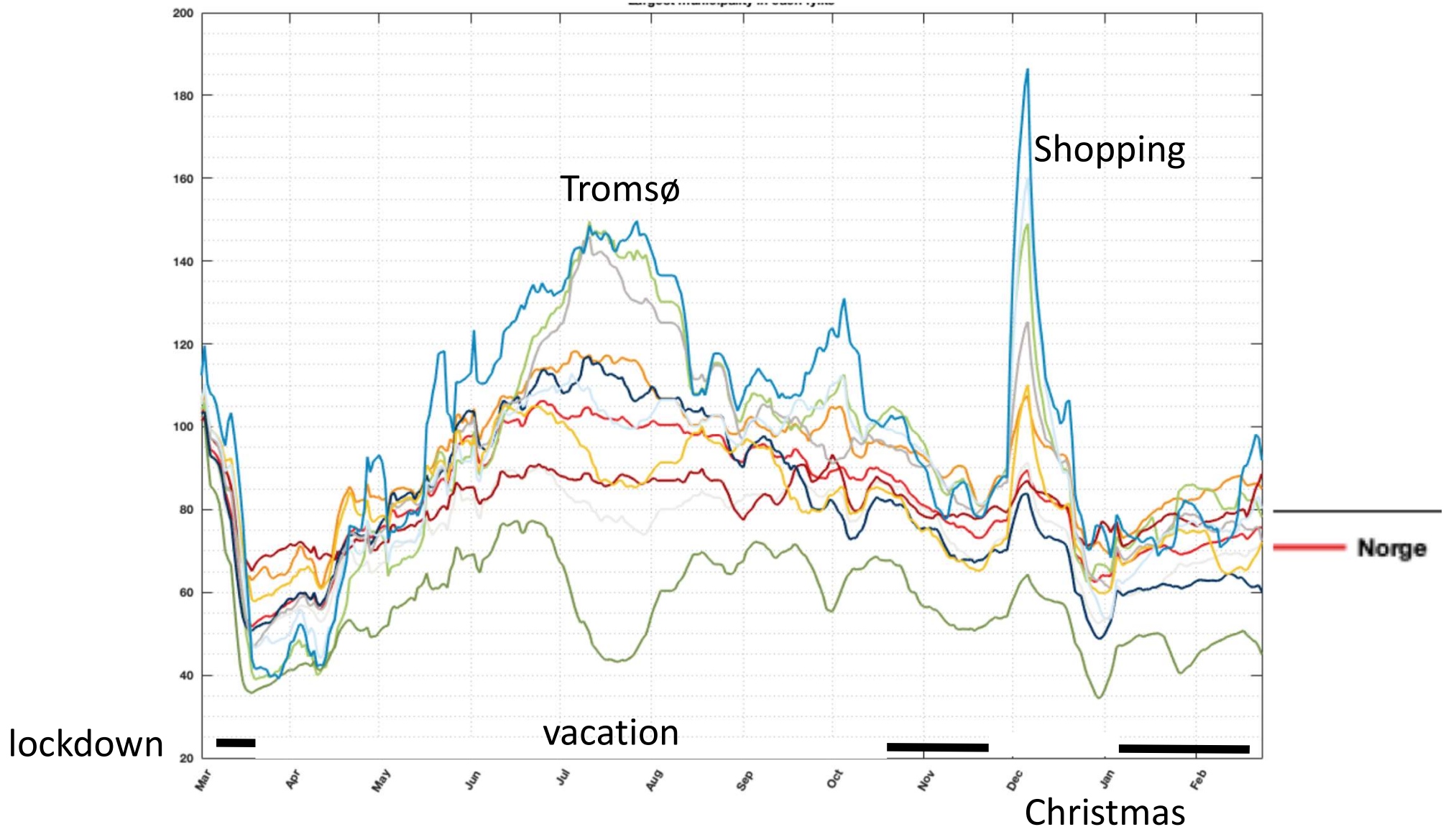
Mobility data

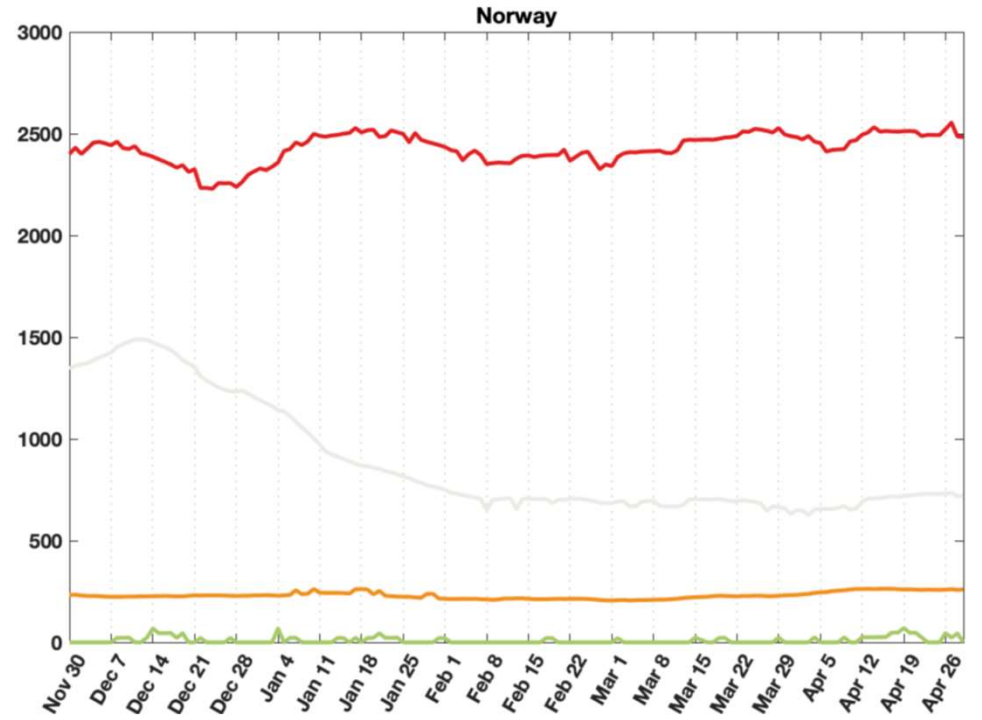
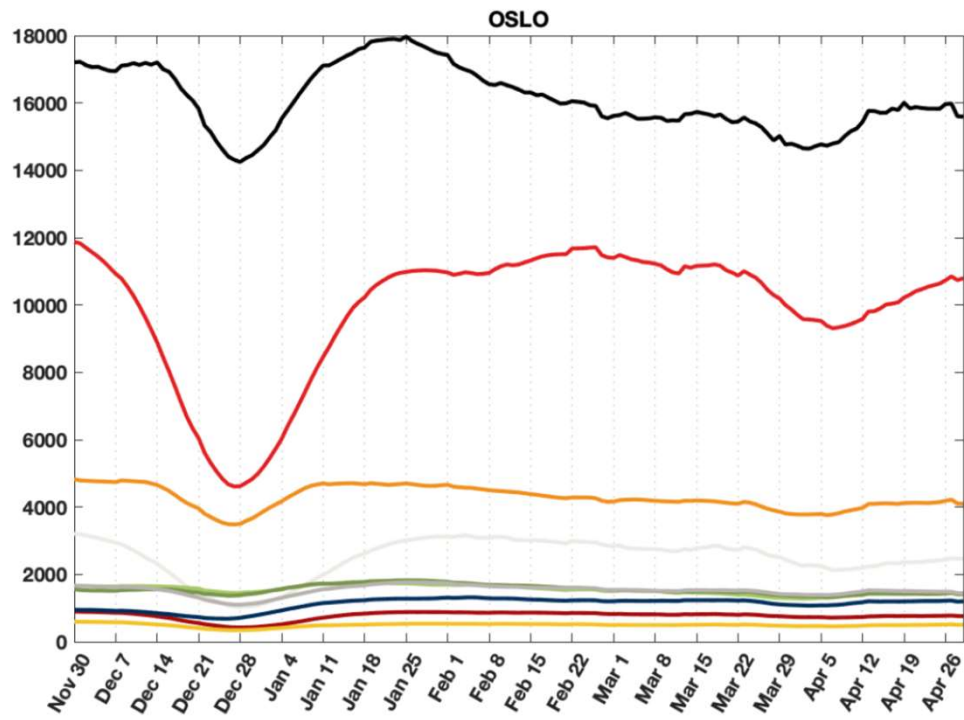


telenor mobile phone mobility, surrogate for real time mobility patterns

- Total movements between the 356 municipalities.
- 4 time intervals: 0-6, 6-12, 12-18, 18-24.
- Counts below 20 are censored.
- Telenor has ca. 48% market share – we scale up.

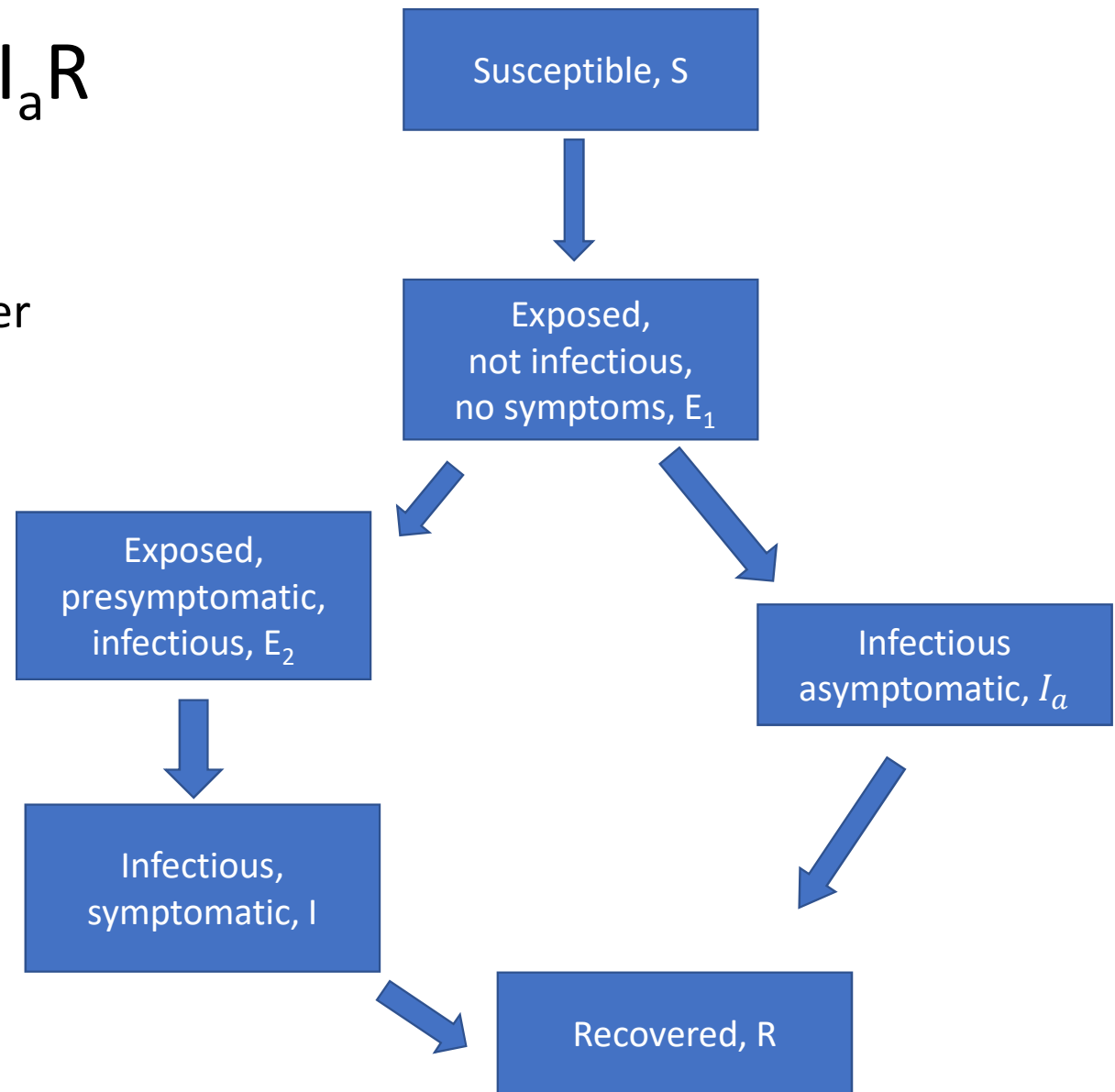
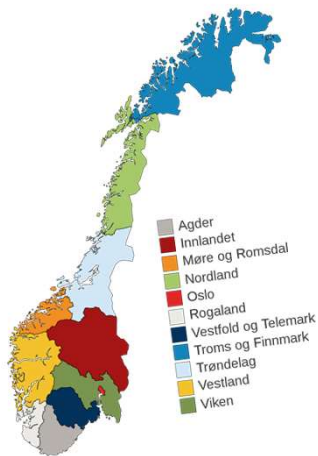
| Date, | Time, | From, | To, | Count |
|-----------|-------|-------|-------|-------|
| 20200224, | 12, | 1101, | 1101, | 8960 |
| 20200224, | 12, | 1101, | 1103, | 239 |
| 20200224, | 12, | 1101, | 1108, | 194 |
| 20200224, | 12, | 1101, | 1111, | 91 |



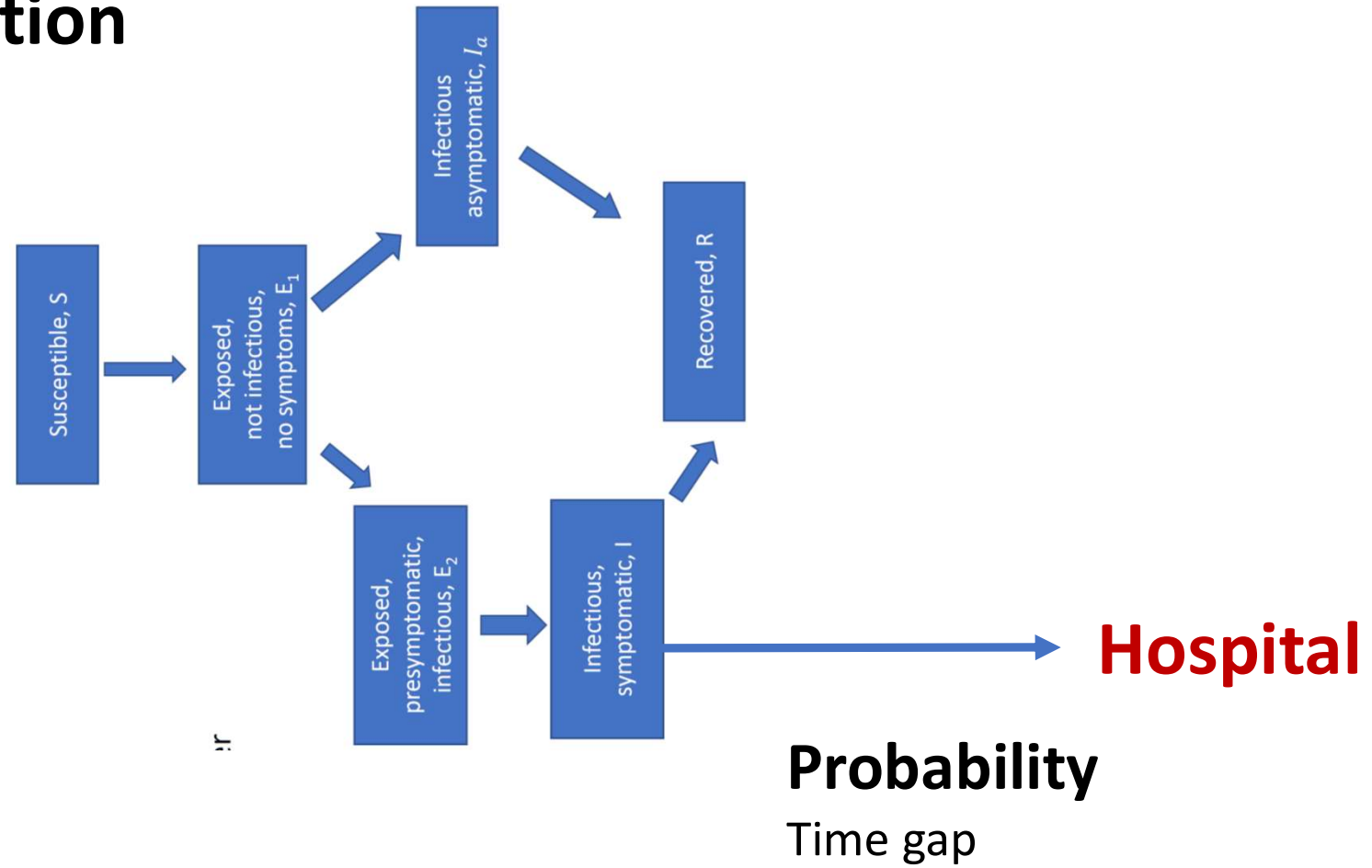


Regional stochastic SEI_aR

- One model per municipality (356)
- County level transmission parameter changing in time
- Other parameters at nation level



Hospitalisation





CORONAVIRUS

Estimating the burden of SARS-CoV-2 in France

Henrik Salje^{1,2,3*}, Cécile Tran Kiem^{1,4*}, Noémie Lefrancq¹, Noémie Courtejoie⁵, Paolo Bosetti¹, Juliette Paireau^{1,6}, Alessio Andronico¹, Nathanaël Hozé¹, Jehanne Richet⁵, Claire-Lise Dubost⁵, Yann Le Strat⁶, Justin Lessler³, Daniel Levy-Bruhl⁶, Arnaud Fontanet^{7,8}, Lulla Opatowski^{9,10}, Pierre-Yves Boelle¹¹, Simon Cauchemez^{1†}

8 July 2020

Probability of hospitalization per age group

Table S1

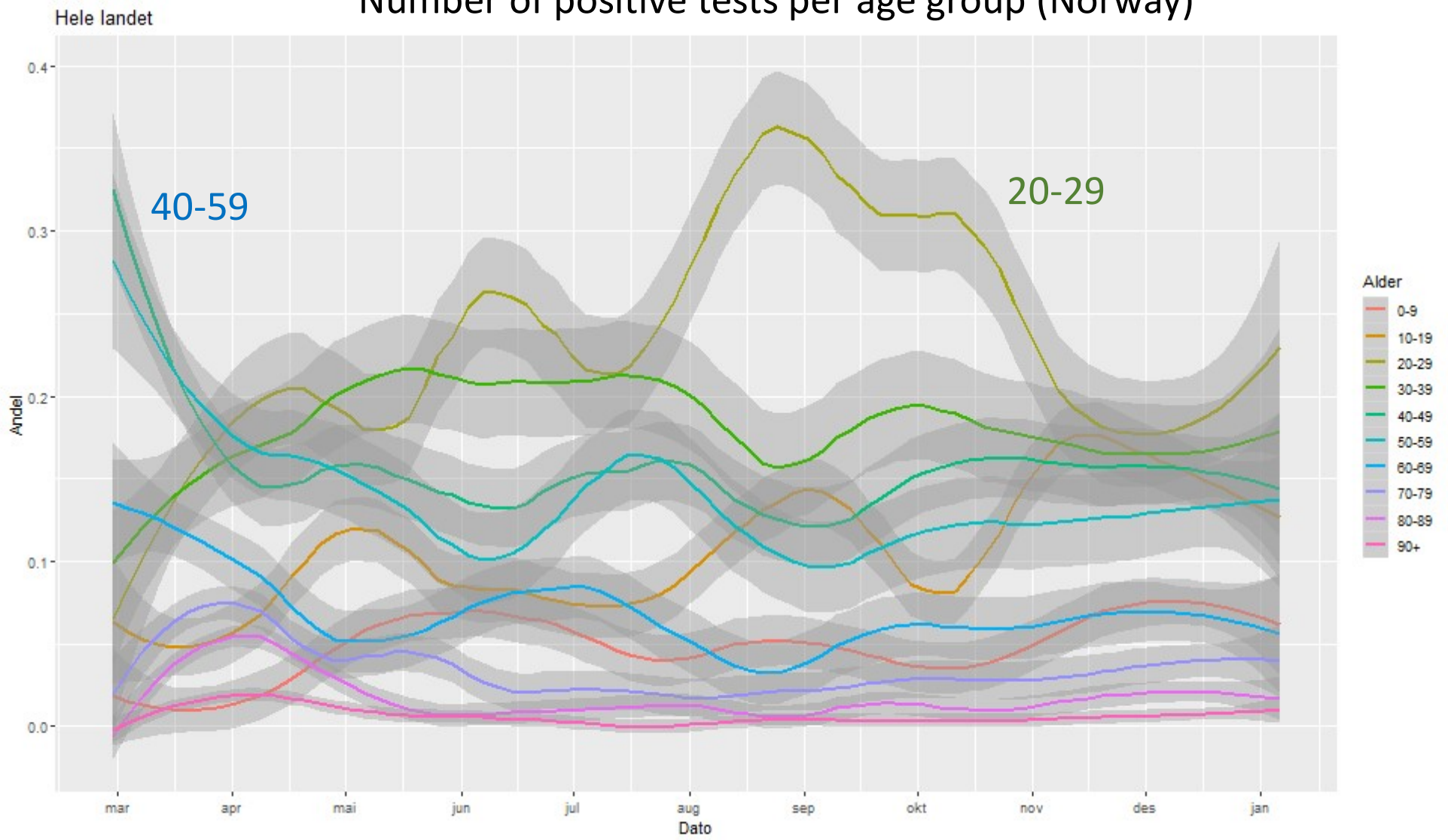
| Age group | Percent infections hospitalized | | | Percent of hospitalized cases that go to ICU | | |
|-----------|---------------------------------|---------------------|---------------------|--|---------------------|----------------------|
| | Male | Female | Mean | Male | Female | Mean |
| <20 | 0.2 (0.08-0.2) | 0.1 (0.07-0.2) | 0.1 (0.08-0.2) | 26.9 (23.1-31.1) | 16.7 (14.3-19.3) | 22.2 (19.1-25.7) |
| 20-29 | 0.6 (0.3- 0.9) | 0.5 (0.3-0.8) | 0.5 (0.3-0.8) | 14.0 (12.2-16.0) | 8.7 (7.5-9.9) | 11.6 (10.1-13.2) |
| 30-39 | 1.2 (0.7-1.9) | 0.9 (0.5-1.5) | 1.1 (0.6-1.7) | 19.2 (17.6-20.9) | 11.9 (10.9-13.0) | 15.9 (14.5-17.3) |
| 40-49 | 1.6 (0.9-2.6) | 1.3 (0.7-2.1) | 1.4 (0.8-2.3) | 26.9 (25.4-28.4) | 16.6 (15.6-17.7) | 22.2 (21.0-23.5) |
| 50-59 | 3.2 (1.8-5.2) | 2.6 (1.5-4.2) | 2.9 (1.6-4.7) | 33.4 (32.0-34.8) | 20.7 (19.8-21.6) | 27.6 (26.5-28.7) |
| 60-69 | 6.7 (3.7-10.9) | 5.1 (2.9-8.3) | 5.8 (3.3-9.5) | 37.3 (36.0-38.6) | 23.1 (22.2-24.0) | 30.8 (29.8-31.8) |
| 70-79 | 11.0 (6.2-17.9) | 7.8 (4.4-12.8) | 9.3 (5.2-15.1) | 30.2 (29.1-31.3) | 18.7 (18.0-19.5) | 24.9 (24.1-25.8) |
| 80+ | 37.6 (21.1-61.3) | 19.3 (10.9-31.6) | 26.2 (14.8-42.7) | 6.8 (6.5-7.2) | 4.2 (4.0-4.5) | 5.6 (5.3-5.9) |
| Mean | 3.3 (1.8-5.3) | 2.6 (1.5-4.3) | 2.9 (1.7-4.8) | 23.1 (22.6-23.6) | 14.3 (13.9-14.7) | 19.0 (18.7-19.44) |

Table S1: Percent of infections that are hospitalized and end up in ICU by age and sex. Percentage of infections that are hospitalized and the percentage that end up in ICU, conditional on being hospitalized.

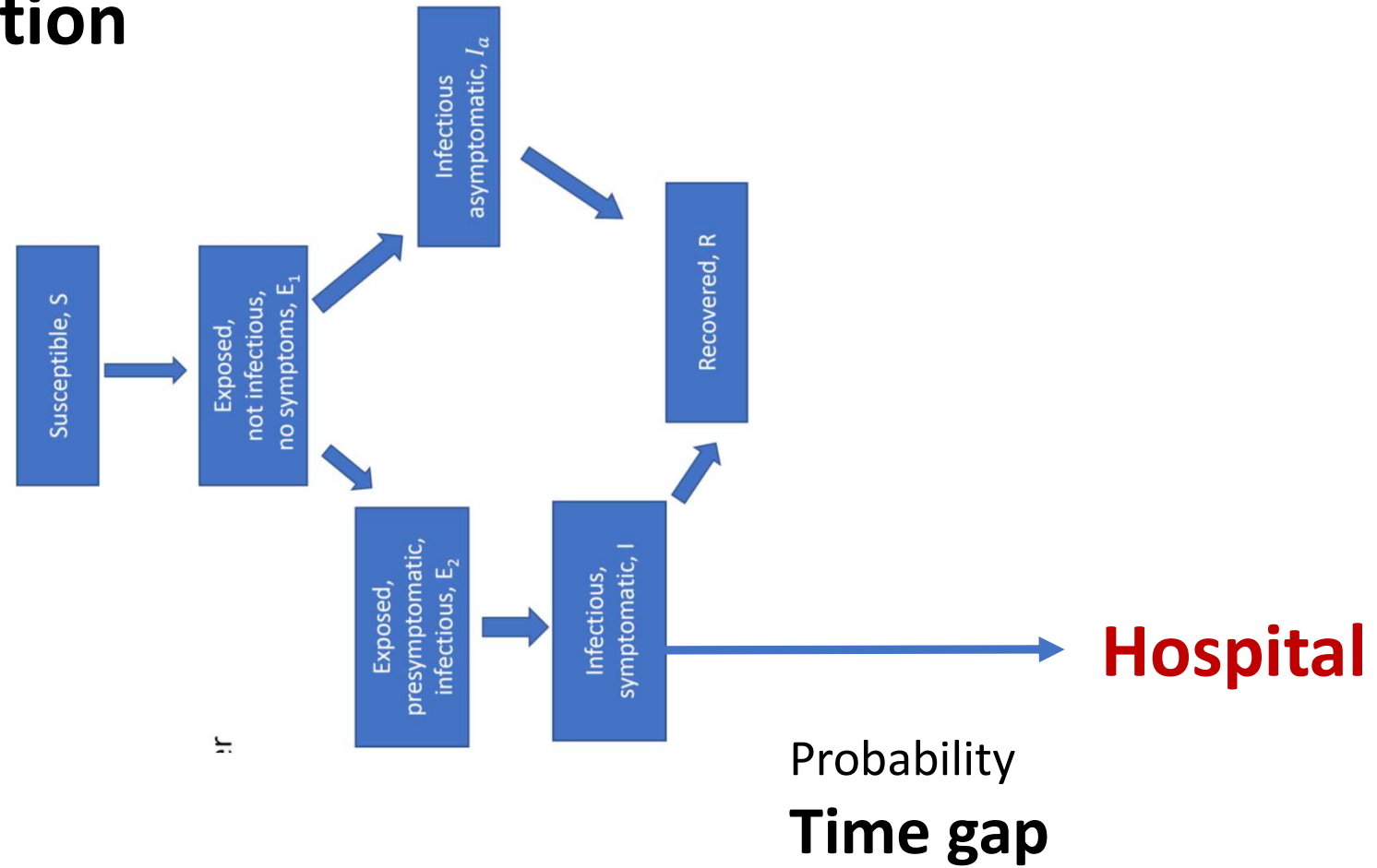
- Scaled to the **demography of each municipality** in Norway and corrected for percentage of elderly living in elderly homes.
- Assumes diseases burden follows demography
Nation average

| | |
|---------------|-------|
| 0-9 years | 0.2% |
| 10 - 19 years | 0.2% |
| 20 - 29 years | 0.6% |
| 30 - 39 years | 1.3% |
| 40 - 49 years | 1.7% |
| 50 - 59 years | 3.5% |
| 60 - 69 years | 7.1% |
| 70 - 79 years | 11.3% |
| 80+ years | 27% |

Number of positive tests per age group (Norway)

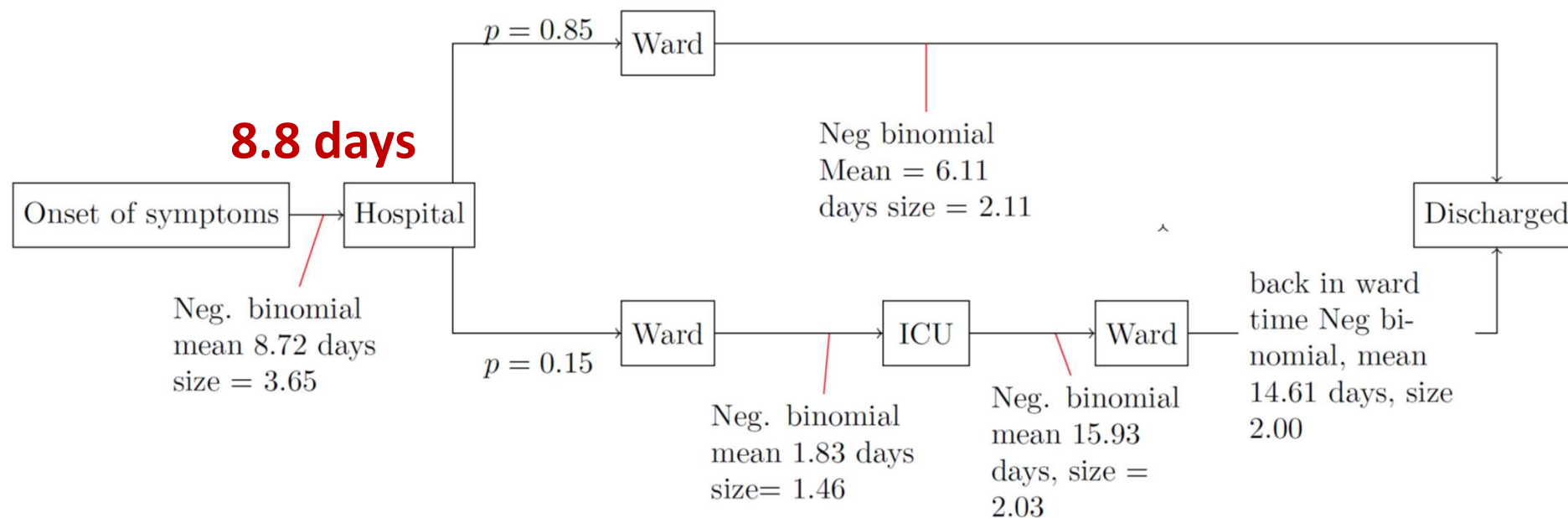


Hospitalisation



Hospitalisation lengths

Estimates from Norwegian registry data.



Up to 31 May 2020

Based on > 1500 hospitalisation cases, taking censoring into account.

Oslo

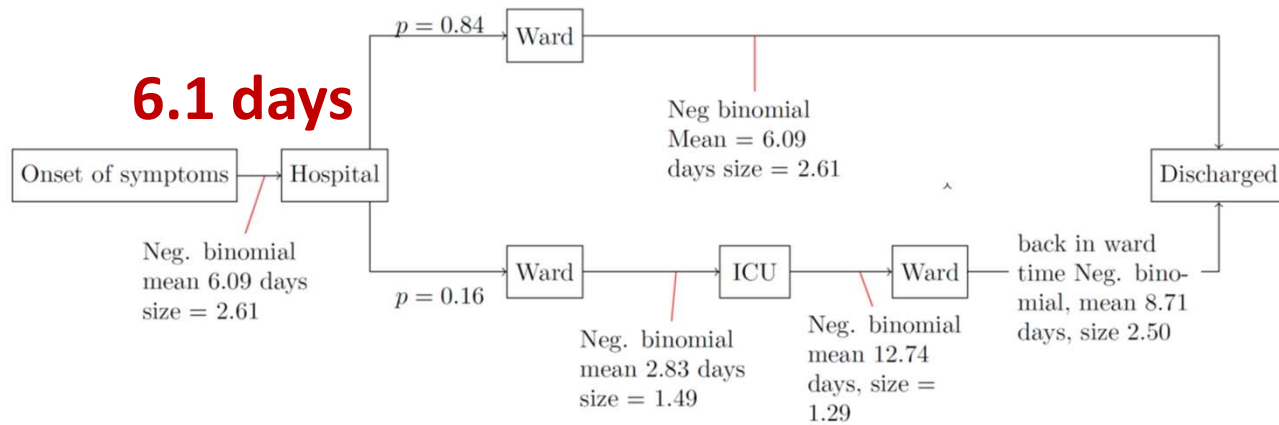


Figure 33: Hospital assumptions and parameters used between 1 January 2021 and 1 March 2021 for those living in Oslo

not Oslo

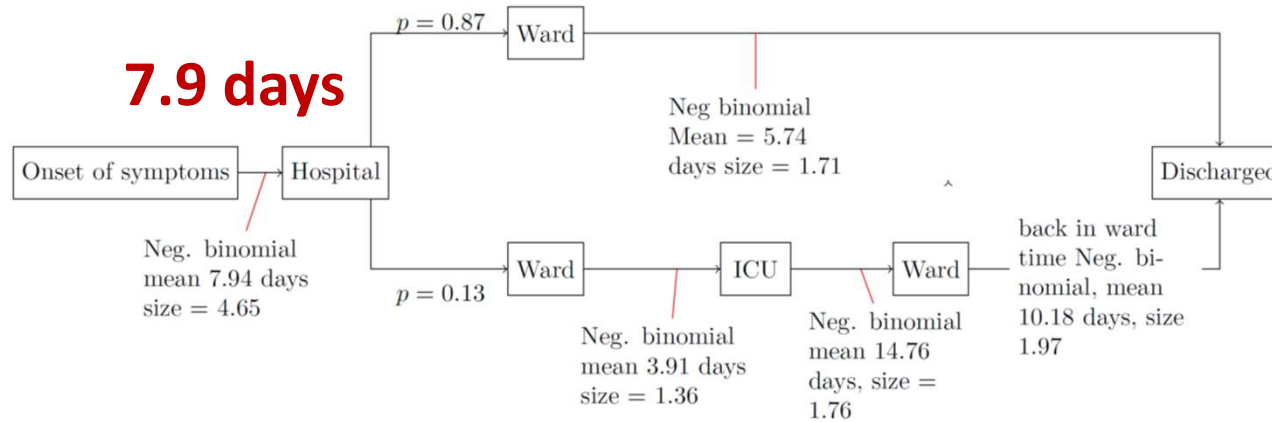
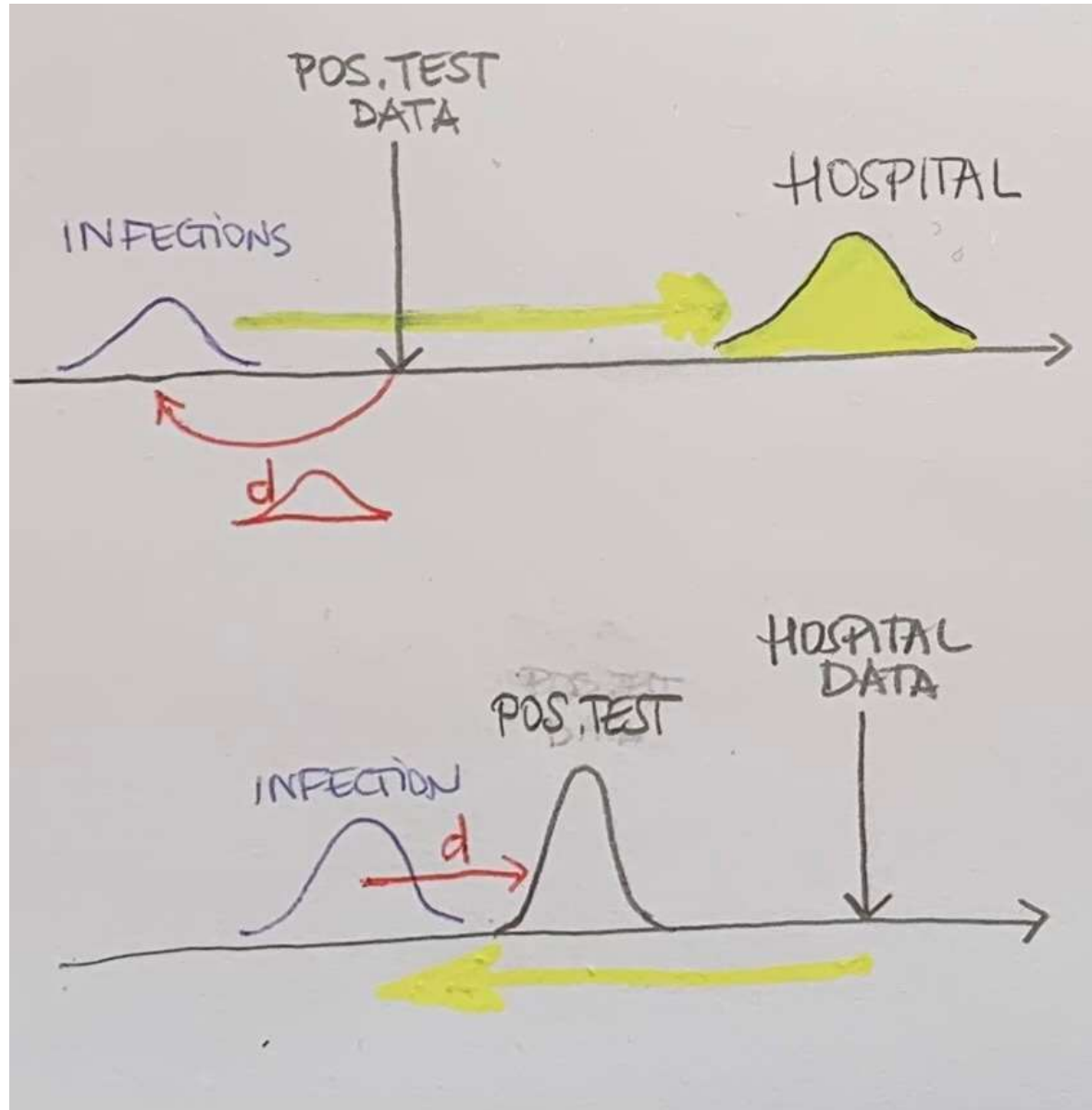


Figure 34: Hospital assumptions and parameters used between 1 January 2021 and 1 March 2021 for those not living in Oslo

From 1 Jan 2021

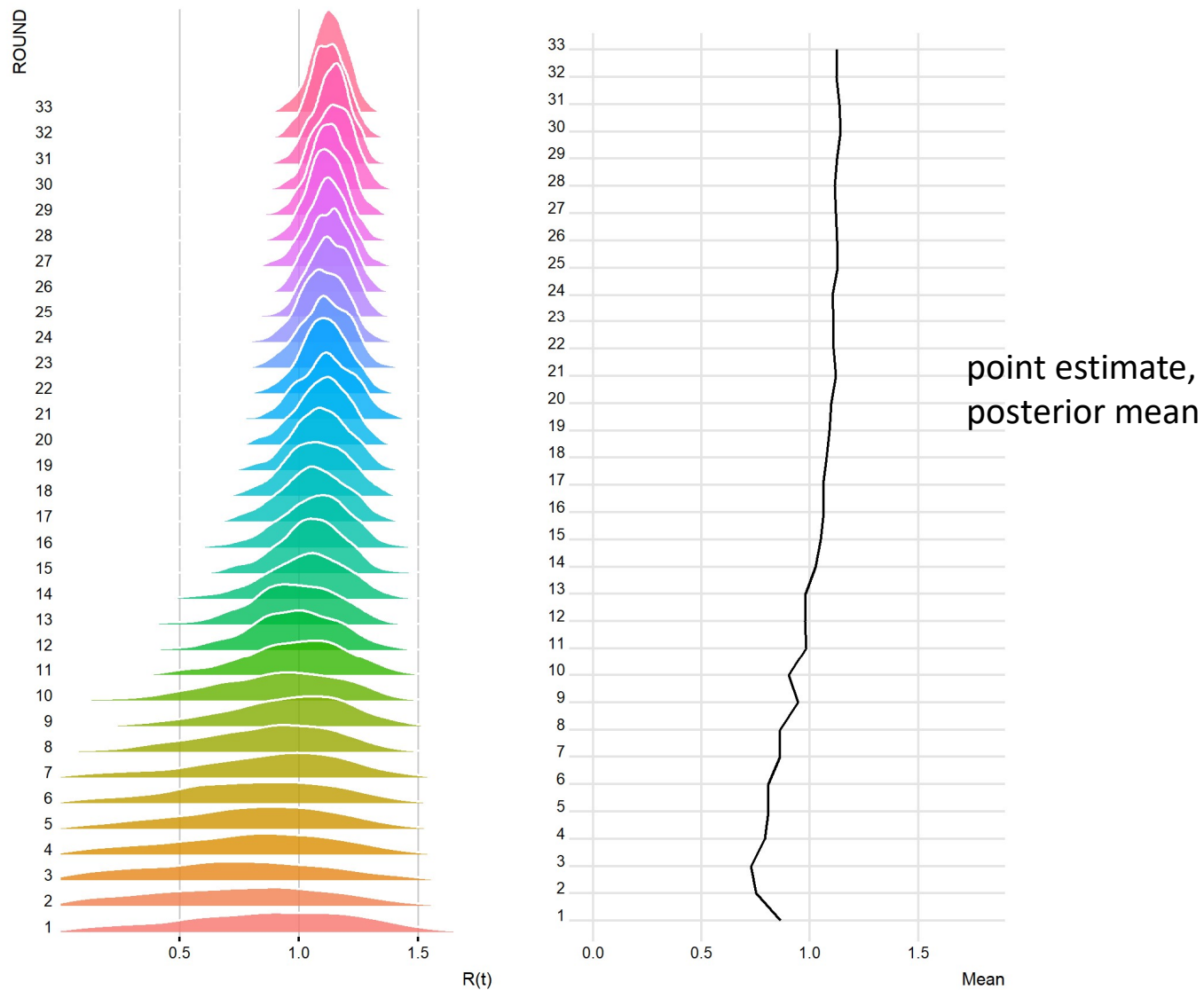
Lack of coherence
between
test and
hospital data



Splitted sequential ABC
to estimate parameters
with uncertainty

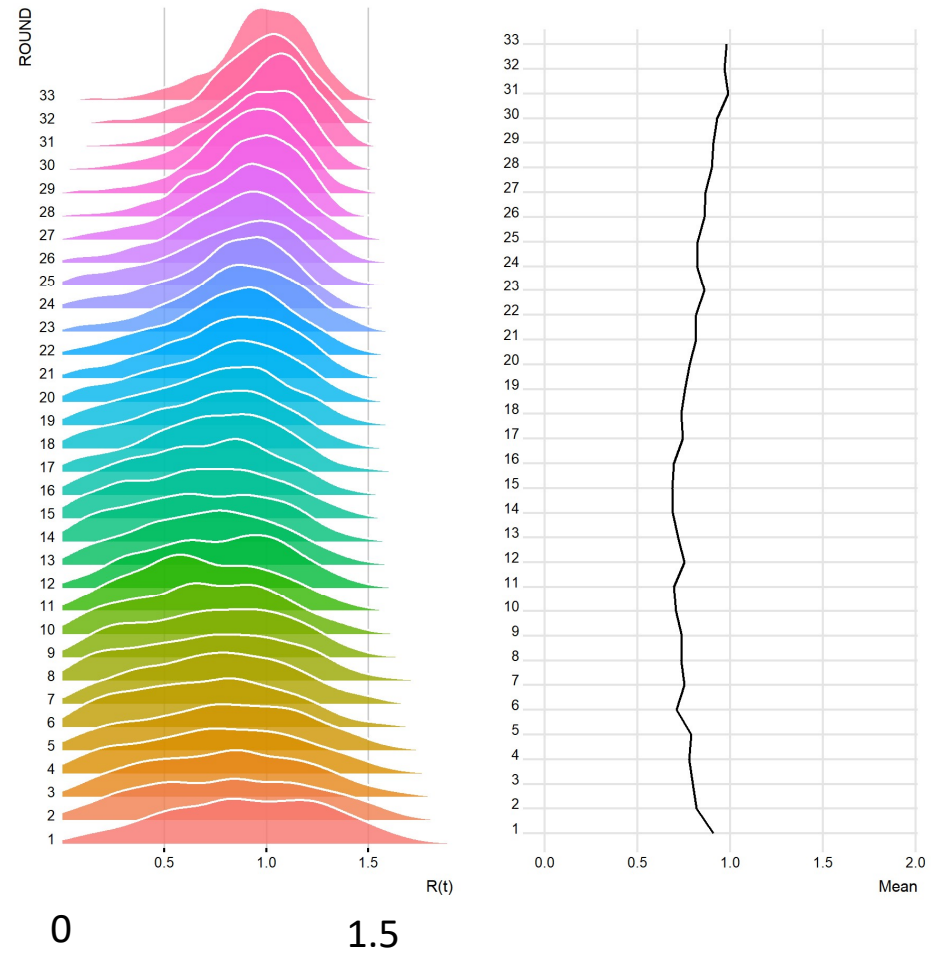
Iterative algorithm:
ca. 2 days
on 2000 cores

effective reproduction number, now, Oslo

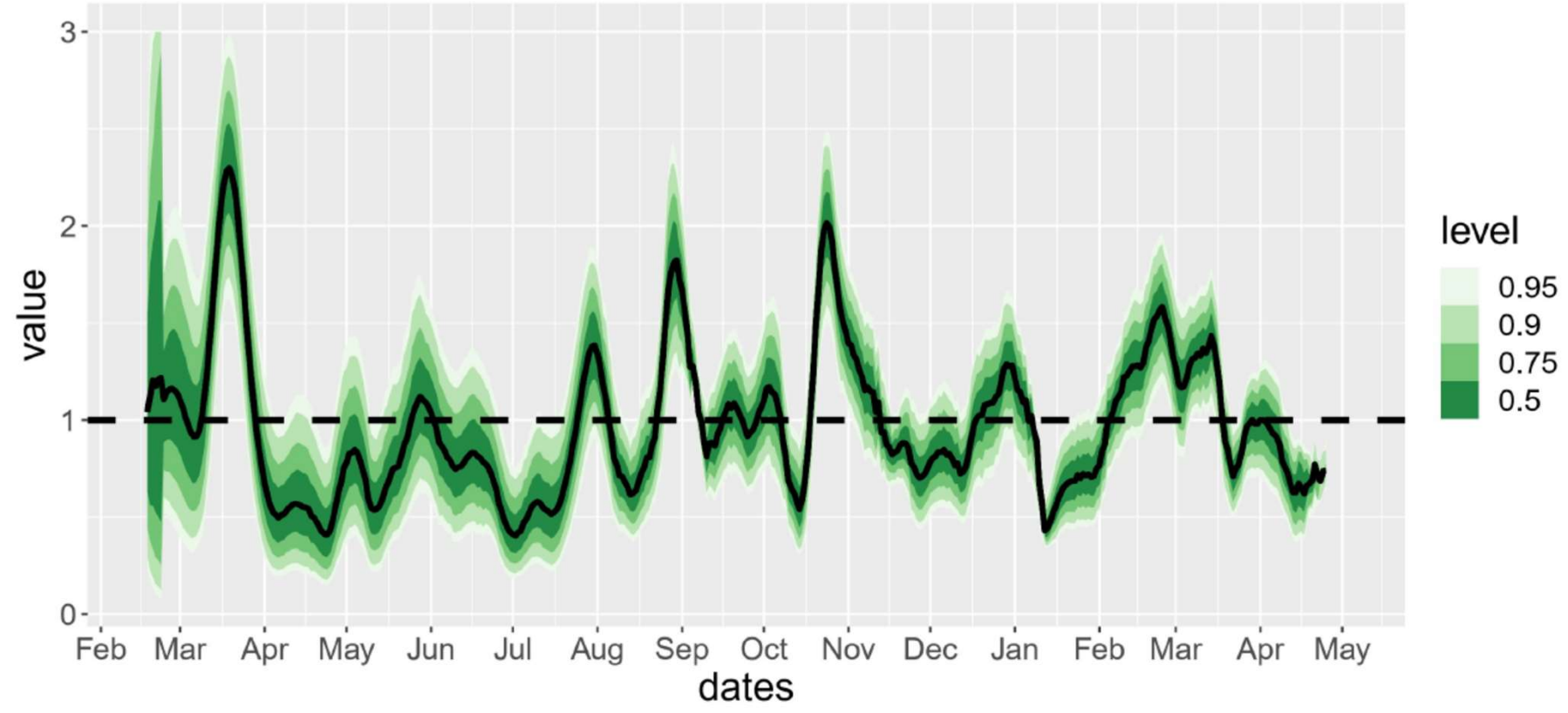


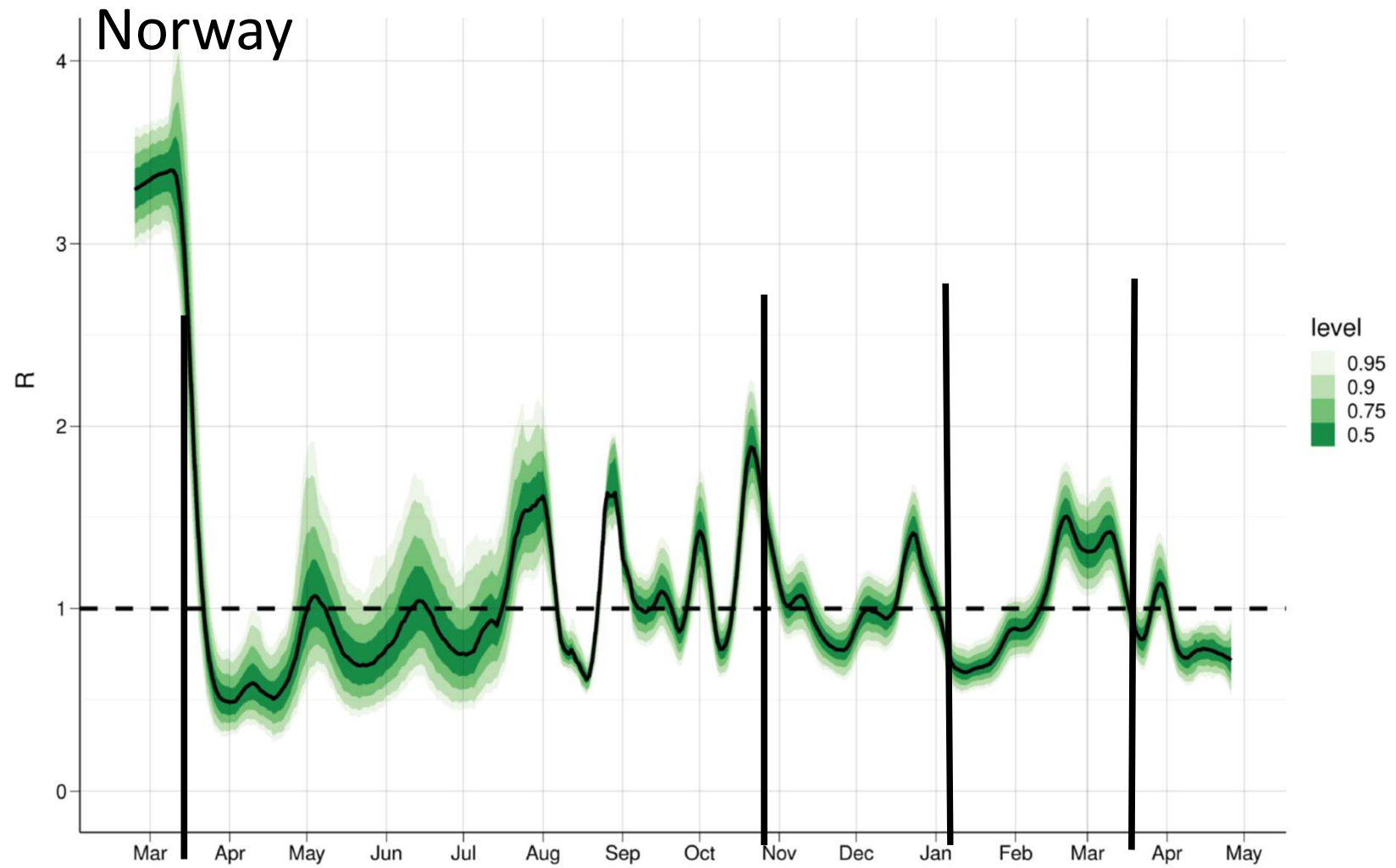
Some counties with very uncertain estimates.

effective reproduction number, Rogaland



Oslo





(12 mar – 20 apr)

(28 oct – 2 dec)

National long-term scenarios with vaccination plans and future interventions

1. PMAJ Pfizer, Moderna, AstraZeneca, Janssen
2. PMJ Pfizer, Moderna, Janssen
3. PM Pfizer, Moderna

- Vaccine deliveries follow a realistic scenario.
- The roll-out accounts for regional prioritization, +20% to Oslo, Moss, Sarpsborg Fredrikstad.
- Vaccine uptake: 90% in all age groups and full adherence to the vaccination schedule.
- Vaccine efficacy for Pfizer and Moderna: 1. dose 60%, 2. dose 90%, for others: 60%, 67%.
- With and without seasonal effects.
- No waning immunity after infection or vaccination.

- The initial conditions in each municipalities follows the last estimates of the regional effective reproduction number.
- Number of vaccinations made by today is known.
- The long term scenario results are based on 100 simulations and accounts for stochasticity within the model.

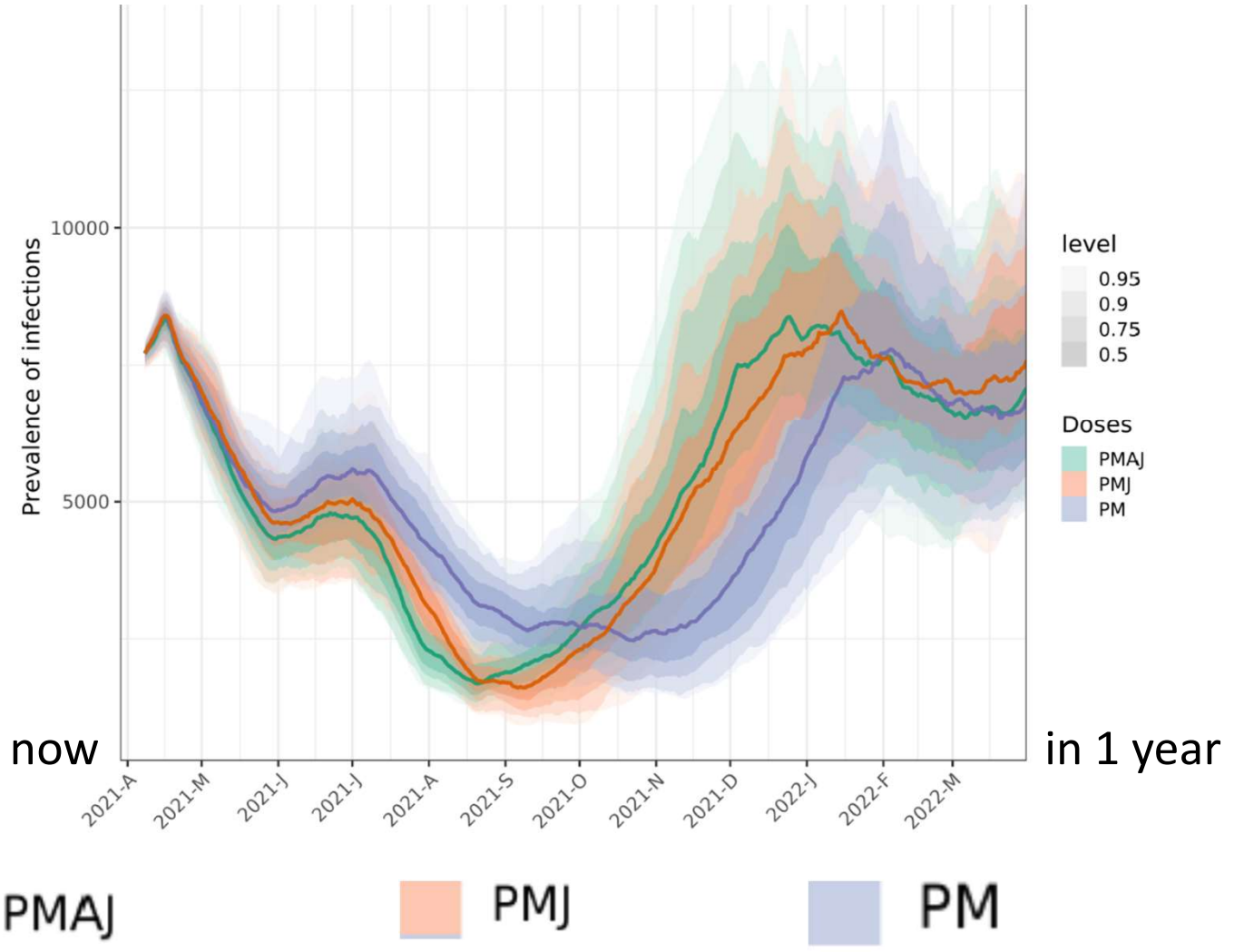
FUTURE interventions:

The government actively controls the reopening of the society:

- When more than 200 hospital beds are occupied, interventions are started that lower the reproduction number to 0.8.
- When less than 50 hospital beds are occupied, interventions are relaxed and the reproduction number increases to 1.2.

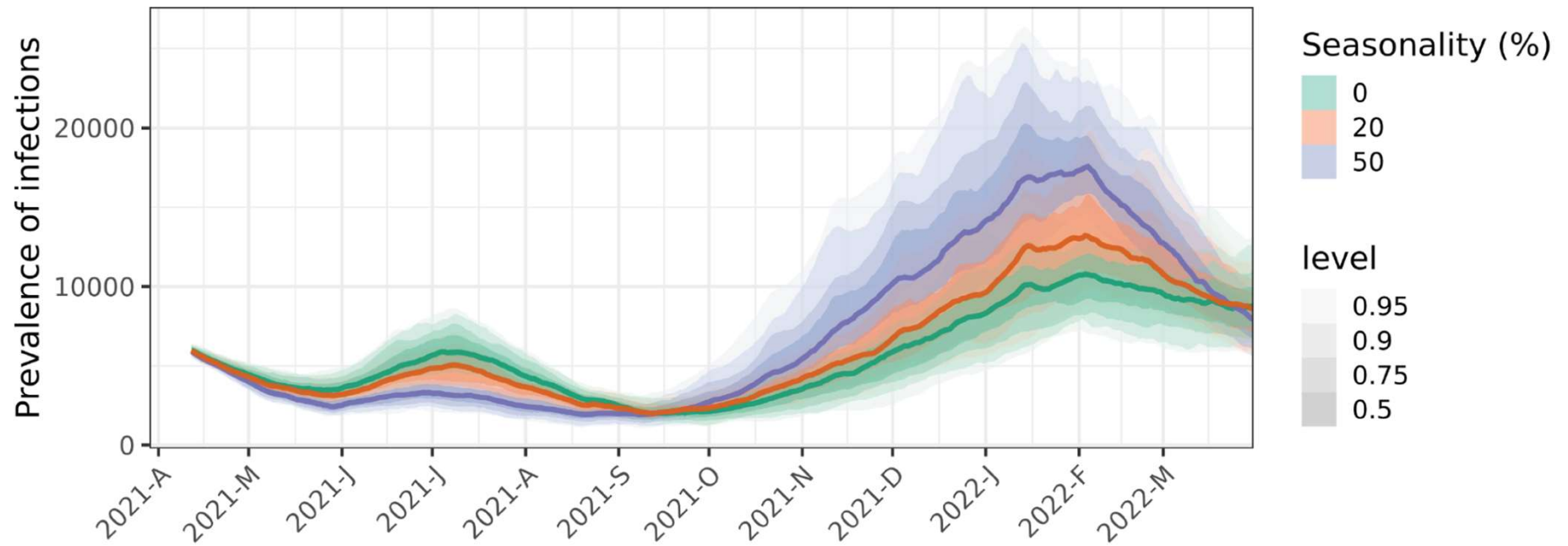
The number of hospital admissions is evaluated every three weeks

Scenario as per end of March



| | | | |
|-----------------------|------------------------|------------------------|------------------------|
| Infections | 374314 (330838-417789) | 373283 (334684-411881) | 356206 (320438-391974) |
| Hospitalisations | 8237 (7469-9004) | 8117 (7450-8785) | 7467 (6955-7979) |
| Ventilator treatments | 631 (569-692) | 617 (563-671) | 561 (520-601) |

Scenario only PM, with seasonal effect as per end of April



Science in emergency

Boris Johnson, 12 March press conference



“It is inevitable that most people will get the disease, so we should let the epidemic proceed to allow 60% of the population to become infected and build **herd immunity**.”

16 March 2020

16 March 2020

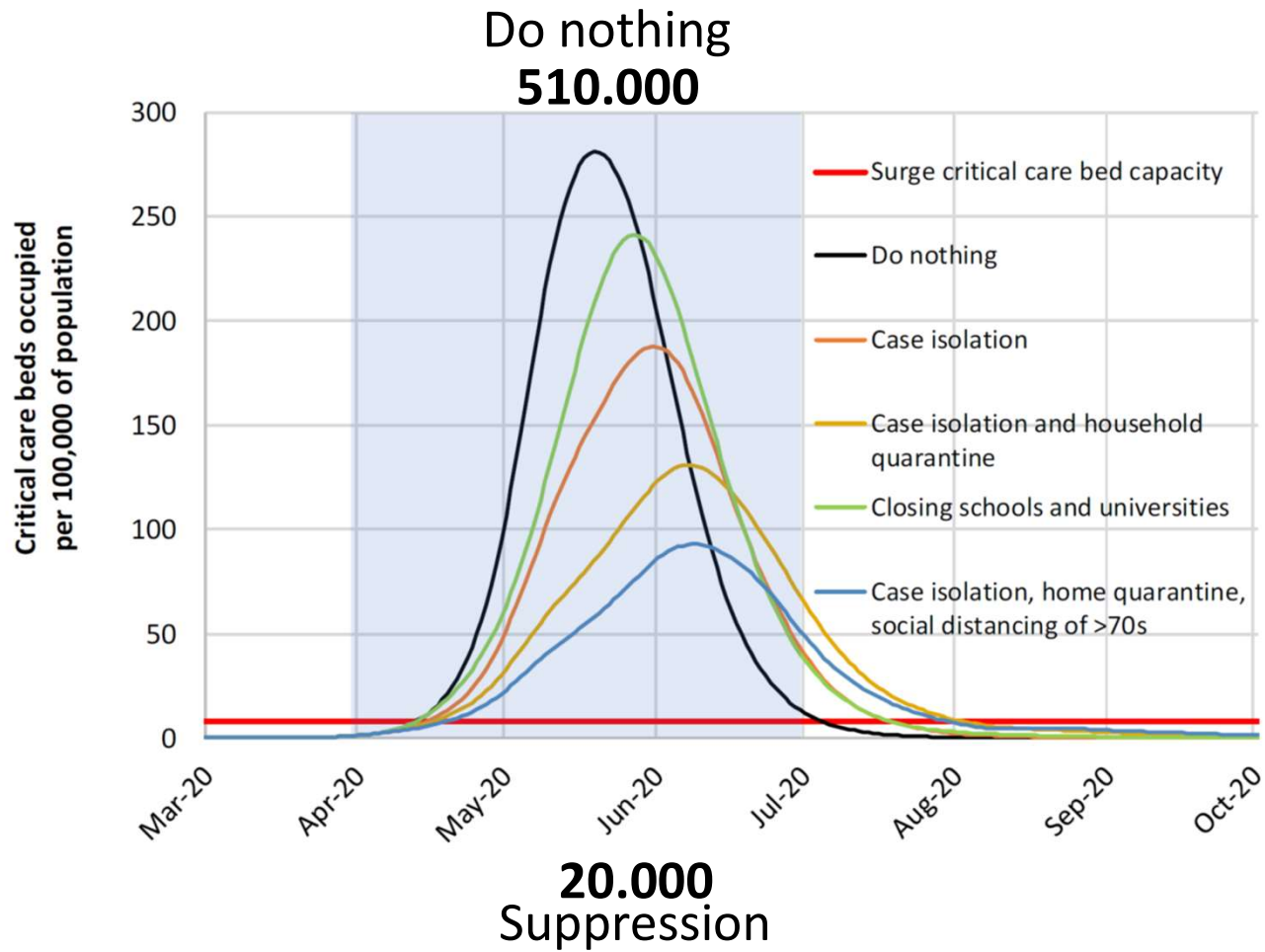
Imperial College COVID-19 Response Team

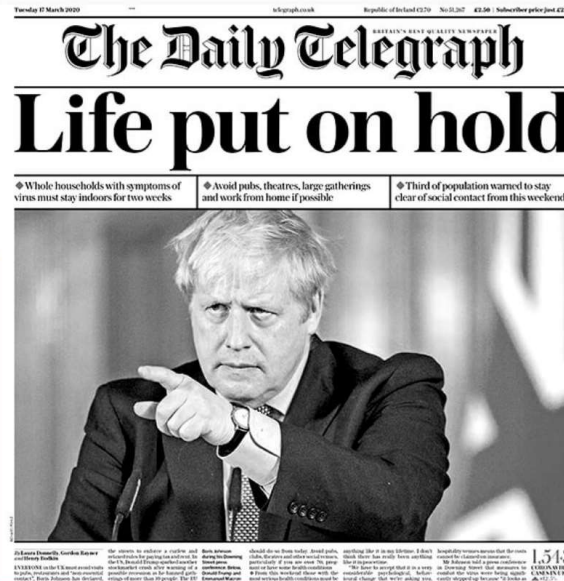
Report 9: Impact of non-pharmaceutical interventions (NPIs) to reduce COVID-19 mortality and healthcare demand

Neil M Ferguson, Daniel Laydon, Gemma Nedjati-Gilani, Natsuko Imai, Kylie Ainslie, Marc Baguelin, Sangeeta Bhatia, Adhiratha Boonyasiri, Zulma Cucunubá, Gina Cuomo-Dannenburg, Amy Dighe, Ilaria Dorigatti, Han Fu, Katy Gaythorpe, Will Green, Arran Hamlet, Wes Hinsley, Lucy C Okell, Sabine van Elsland, Hayley Thompson, Robert Verity, Erik Volz, Haowei Wang, Yuanrong Wang, Patrick GT Walker, Caroline Walters, Peter Winskill, Charles Whittaker, Christl A Donnelly, Steven Riley, Azra C Ghani.

On behalf of the Imperial College COVID-19 Response Team

A stochastic, spatially structured, individual-based discrete time simulation model.





Mar
17


Vallance says 20,000 Covid deaths a "good outcome"

Sir Patrick Vallance tells MPs that **keeping the total number of Covid-19 deaths under 20,000 would be "a**


Mar
23

UK lockdown comes into effect


Johnson announces people may only leave their homes for **strictly limited reasons** and gives the police powers to enforce the rules. The furlough scheme introduced by

 **Imperial College** ✓
@imperialcollege

Neil Ferguson and Imperial **did not produce a model** for Sweden pointing to 85,000 deaths

 **Fraser Nelson** ✓ @FraserNelson · Sep 19, 2020
When Neil Ferguson's Imperial model pointed to 85,000 deaths, the Swedish authorities has enough in-house expertise to know the model was wrong. Anders Tegnell, its state epidemiologist, said...
[/article/why-bo...](#)


4:51 PM · Sep 20, 2020 · Twitter Web App

 **neil_ferguson** ✓ @neil_ferguson · Mar 26, 2020

2/4 **-This is not the case.** Indeed, if anything, our latest estimates suggest that the virus is slightly more transmissible than we previously thought. Our lethality estimates remain unchanged.

75 1.6K 5.1K

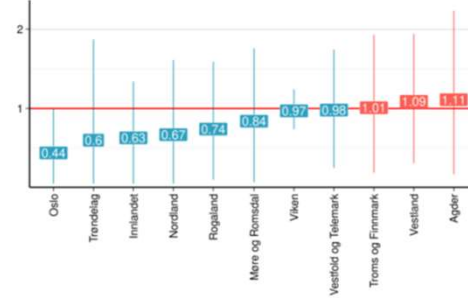
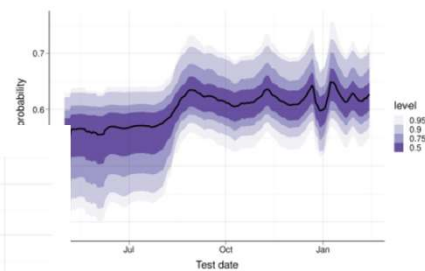
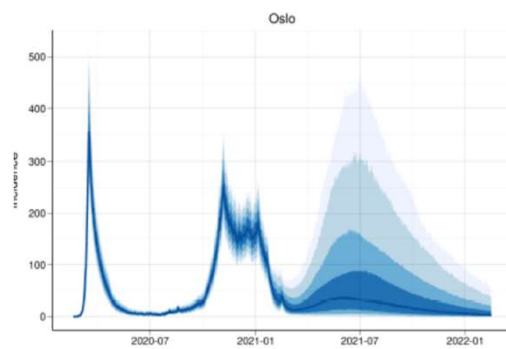
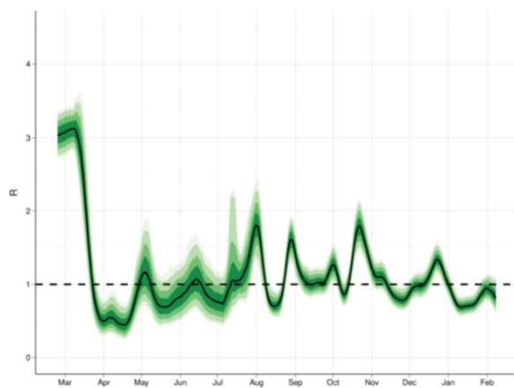
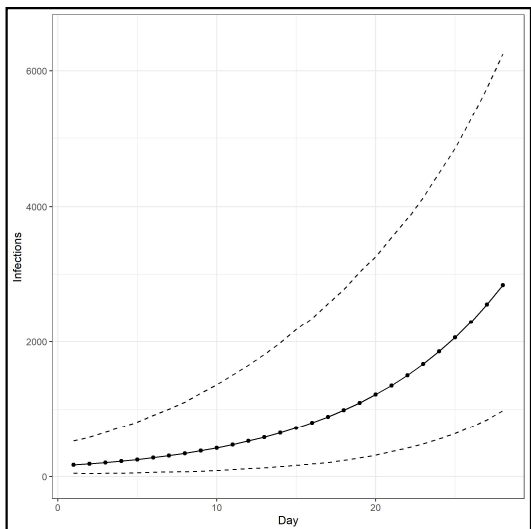
[Show this thread](#)

 **neil_ferguson** ✓ @neil_ferguson · Mar 27, 2020

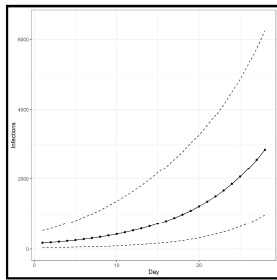
Let me be clear. This virus has a lethality substantially in excess of "seasonal" flu. Yes, up to half of those infected might not show symptoms. But that is accounted for in our estimates and always was. There is no credible data supporting the idea that 90% are asymptomatic.

296 801 1.6K

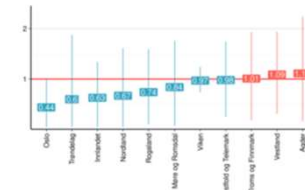
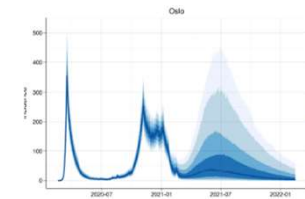
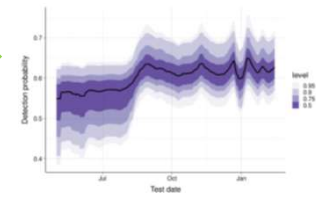
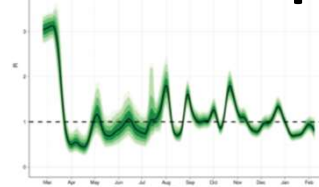
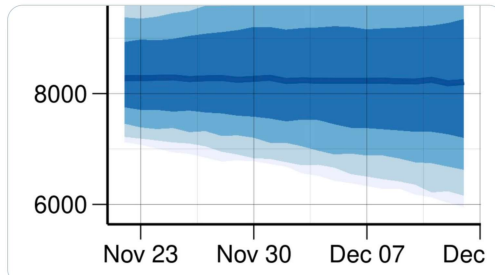
This online publication **has been corrected. The first corrected version first appeared at [thelancet.com/infection](https://www.thelancet.com/infection) on April 15, 2020 and the second on May 4, 2020**



Strongly, repeatedly influenced by the specialists and the public










Hallvard Sandberg @HalSandberg · 28 nov 2020
 Stemmer beregningene fra Folkehelseinstituttet, så har vi et problem. Her er antallet til en hver tid infiserte personer fram til 14. desember. Det er en flat linje. Det betyr at smitten ikke går ned. Da må vi enten stramme enda mer til, eller leve som nå og godta prisen.



CORONAVIRUSET →

Tall | Vaksinerte | Tiltak | Siste

| | | | | |
|---|--|---|--|---|
| Registrert smittet 286 Flat i går trend | Satte vaksinedoser 313 299 5,84 totalt per 100 | Innlagte 69 11 nå | Andel positive tester 1,3 % 0,2 siste u. | Døde 607 totalt |
|---|--|---|--|---|

|  Situational awareness and forecasting for Norway |  |  |  |  |  |  |
|--|--|---|--|--|--|---|
| <p style="text-align: center;"> FHI COVID-19 modelling team UPDATE NOTES-week 5 2021 </p> <p>Log of changes</p> <p>Here we list aspects of the model or of the input parameters which have changed compared to previous reports, and we explain the reason for these changes. Some changes will have big effects on some of our estimates.</p> <p>14 April: Hospitalisation risk: Our model requires the specification of the proportion of symptomatic and asymptomatic patients requiring hospitalisation. Previously we used estimates from Verity et al. (2020) based on Chinese data, adapted to the Norwegian demography, and to the reduced mobility of elderly patients living in elderly homes, and calibrated to carry a clearer signal of the infection signet with a delay, which we estimate to be around 35-600. The effect of this change is visible on the estimated cumulative number of infected individuals, which is now approximately 45,000. A further effect of this change is that the reproductive numbers are different, with R_0 larger and R_{eff} smaller than before, when we had a higher hospitalisation risk.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> | <p>We introduce a new changepoint in the R_0 from 20 April. This is the day of school resumption, and we will see if it is reflected in the data.</p> <p>Our model requires the specification of a required hospitalisation. Previously we used estimates from Verity et al. (2020) based on Chinese data, adapted to the Norwegian demography, and to the reduced mobility of elderly patients living in elderly homes, and calibrated to carry a clearer signal of the infection signet with a delay, which we estimate to be around 35-600. The effect of this change is visible on the estimated cumulative number of infected individuals, which is now approximately 45,000. A further effect of this change is that the reproductive numbers are different, with R_0 larger and R_{eff} smaller than before, when we had a higher hospitalisation risk.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> | <p>posterior probability of the time-irreversibility to start the uncertainty, which is the daily reproduction number probability that the reproduction rate will increase until June 26. They require amplification factor for all reported cases until the day before which is re-estimated every time a new case is reported.</p> <p>We introduce a new changepoint in the R_0 from 20 April. This is the day of school resumption, and we will see if it is reflected in the data.</p> <p>Our model requires the specification of a required hospitalisation. Previously we used estimates from Verity et al. (2020) based on Chinese data, adapted to the Norwegian demography, and to the reduced mobility of elderly patients living in elderly homes, and calibrated to carry a clearer signal of the infection signet with a delay, which we estimate to be around 35-600. The effect of this change is visible on the estimated cumulative number of infected individuals, which is now approximately 45,000. A further effect of this change is that the reproductive numbers are different, with R_0 larger and R_{eff} smaller than before, when we had a higher hospitalisation risk.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> | <p>of hospitalisation, from May 1 updated to the data on patients hospitalised in Norway.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> | <p>based only on demography. When reproduction number distribution and a new hospitalisation rates, which influence the after the first three weeks, we use rates from the model does not have age classes, and in winter. New projections include updated data.</p> <p>Approximately around the 21st of every month, acting from the start of the month.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> | <p>not list future changes which will happen.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> | <p>patients who have been infected. The amplified individuals estimates, are also added to the probabilities to seed cases to check if the age profile in the case update the probabilities.</p> <p>As we transitioned to using more updated estimates of the reproduction number, we have difficulties in convergence to run about 65 rounds, and for the very large working with several options to be doing for a long time, but when we simulate the next 1, 2, 3 and 2 weeks ago, with the new information.</p> <p>14 April: Change point for the reproductive number: On March 12, a number of contact restrictions were implemented. During that week 11, mobility was reduced significantly, and appears to stabilise on Monday March 16th. Between the 11th and 16th of March we expect a reduction of the reproduction rate. We model this change as a sudden jump from a first reproduction rate R_0 to a second and lower reproduction rate R_{eff}, through a change in the model parameter β. We have chosen Monday March 15 as the changepoint for the reproductive number because it gives the best fit to the hospitalisation data. If we move the changepoint to March 14, or assume a continuous linear reduction during week 11, the fit deteriorates. We also notice that the best changepoint depends on the assumed time between symptoms appearance and hospitalisation, which is assumed to have mean 8 days in this report. The optimal changepoint also depends on the assumed hospitalisation risk.</p> <p>20 April: Change in parameter estimation method: We use sequential ABC instead of iterative parameter calibration. Estimation of the reproduction numbers and of the amplification factor in the setting of the epidemic at the start is done using Approximate Bayesian Computation (ABC), as described in Engelen et al. (2020)¹. Sequential ABC avoids to calibrate R_0 first on part of the data and then, given the best value of each R_0, to find the best fitting R_{eff}, which might not lead to optimal estimation and is based on more ad-hoc choices. We also do not weigh the last part of the data more than the rest. Sequential ABC takes more time to run: therefore the daily report might use only the hospitalisation until yesterday.</p> <p><small>¹https://www.medrxiv.org/content/10.1101/2020.04.11.20055514</small></p> |



Christophe Fraser   @ChristoPhraser · 8h

I feel like we have a recurring problem on what to in time-periods when evidence is still being collected. Zero effect of vaccines on transmission is not a reasonable prior. Bayesian thinking seems highly appropriate. We can make temporary assumptions and update them.

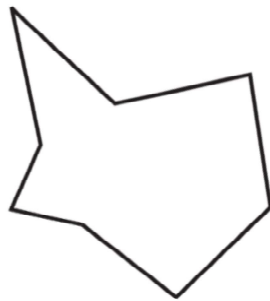
A new way of doing science?

Evidence-based science:

- available scientific knowledge and evidence from data lead to results which are “final given the current situation”.
- Results can be used *as-is*.
- Predictions will be checked in the future, and will be compared to reality.

Evidence-making science:

- models are always **starting points** and not the final truth;
- models, hypothesis, data **change**, also on the basis of needs and questions from the public, media and decision makers.
- Emergency requires **rapid results**. There is no time to curate data, to test code properly.
- How do you communicate to the public that your results might be wrong and are likely to change tomorrow?
- Scientists are not (always) **coherent** with themselves. Scientists must learn to say “I was wrong” and public must appreciate this, instead than getting more sceptical about science.
- **Uncertainty** means that model sets boundaries to describe the possible future. These boundaries create a space, a playground for decisions, where ethics, politics, economy, culture must be advoked.



https://www.fhi.no/contentassets/e6b5660fc35740c8bb2a32bfe0cc45d1/vedlegg/nasjonale-og-regionale-rapporter/national_regional_model_05may2021.pdf

Lindstrøm, J. C., Engebretsen, S., Kristoffersen, A. B., Rø, G. Ø. I., Palomares, A. D. L., Engø-Monsen, K., ... & de Blasio, B. F. (2021). Increased transmissibility of the B. 1.1. 7 SARS-CoV-2 variant: Evidence from contact tracing data in Oslo, January to February 2021. *medRxiv*.

Engebretsen, S., Engø-Monsen, K., Aleem, M. A., Gurley, E. S., Frigessi, A., & de Blasio, B. F. (2020). Time-aggregated mobile phone mobility data are sufficient for modelling influenza spread: the case of Bangladesh. *Journal of the Royal Society Interface*, 17(167).

Williams, G. A., Díez, S. M. U., Figueras, J., & Lessof, S. (2020). TRANSLATING EVIDENCE INTO POLICY DURING THE COVID-19 PANDEMIC: BRIDGING SCIENCE AND POLICY (AND POLITICS). *Eurohealth* 2020; 26(2).

Rhodes, Tim, and Kari Lancaster. "Mathematical models as public troubles in COVID-19 infection control: following the numbers." *Health Sociology Review* 29.2 (2020): 177-194.

Mangoni, A. A., & Erre, G. L. (2021). Translating evidence into practice during the COVID-19 pandemic: pitfalls and mileages. *Therapeutic Advances in Drug Safety*, 12.