

Predictive Modeling approach of COVID-19 propagation in France

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Paris France

Topics

- 1. Public data**
- 2. Evaluation of masks and social distancing**
- 3. Model calibration**
- 4. Simulation results**
- 5. Method: Covid multiscale rate modeling**
- 6. Main conclusions of simulations**

Public data

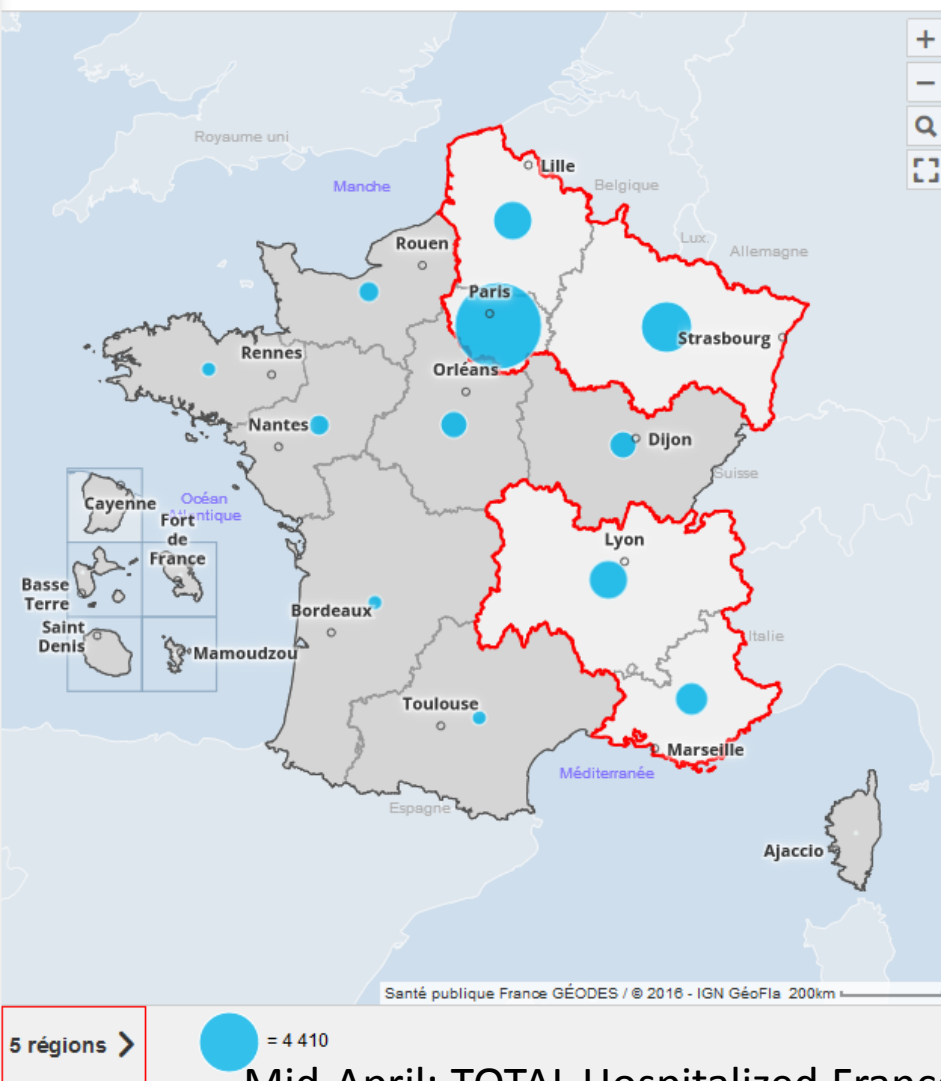
Focus on 5 Main regions

1 Nombre de personnes actuellement hospitalisées pour COVID-19 - tous âges 2020-05-06

ACTIONS

Nombre de personnes actuellement hospitalisées pour covid-19 - tous âges

ACTIONS



Chiffres-clés 2020-05-06

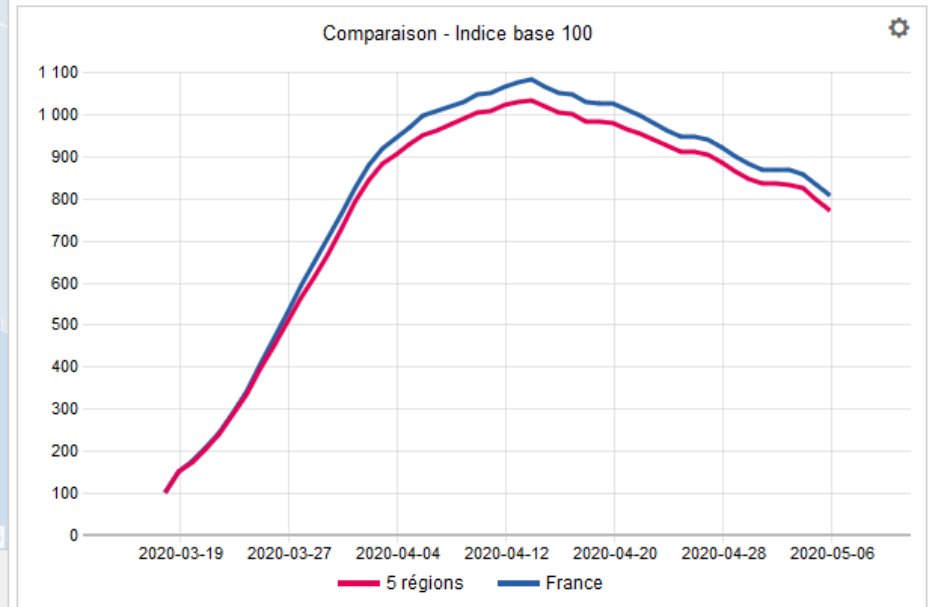
Statistique	France	5 régions
minimum	5 (Guyane - 03)	1 359 (Provence-Alpes-Côte d'Azur - 93)
maximum	9 930 (Île-de-France - 11)	9 930 (Île-de-France - 11)
moyenne	1 328	3 837
médiane	549	2 299
observations valides	18 sur 18	5 sur 5

France : 23 912

5 régions : 19 187

Graphiques et comparaisons

Évolution temporelle comparée



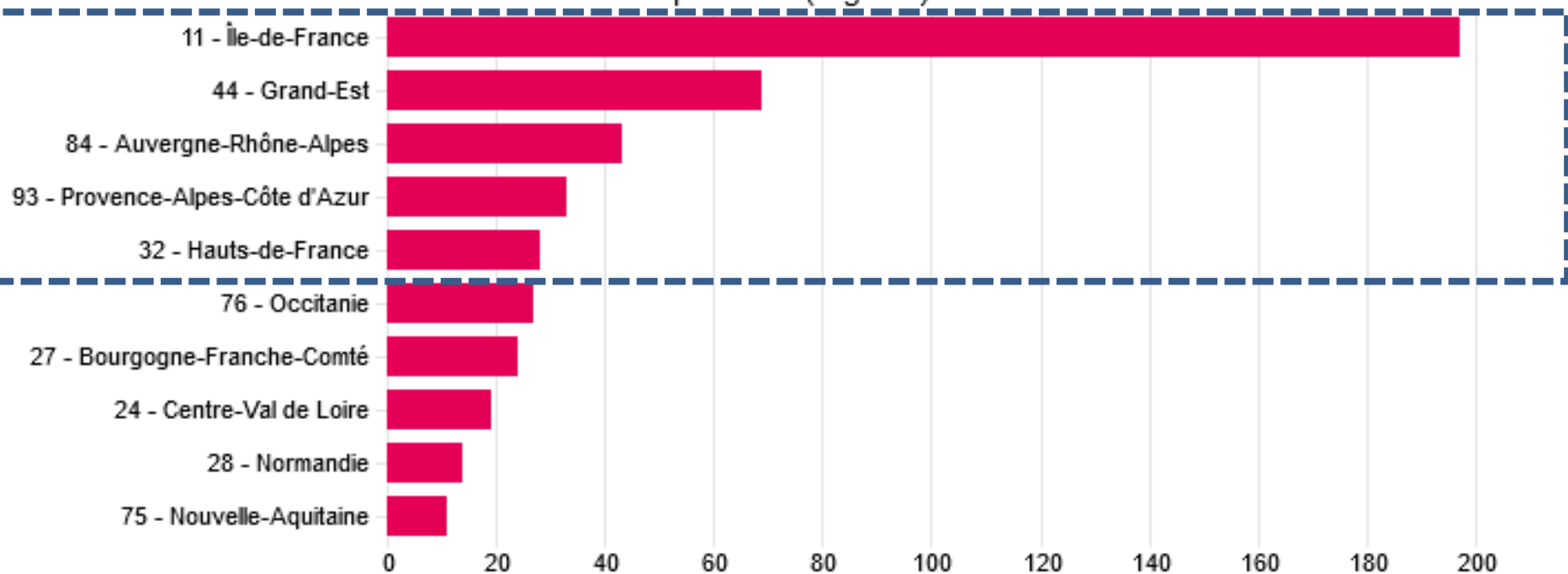
Mid-April: TOTAL Hospitalized France = 28254 vs 5 regions 23450 Fraction=0.8

Total population : 36.792 million/5 regions compared to 67 Millions Fraction=0.54

Conclusion: 1/2 of the population concentrate 80% of cases.

Focus on 5 Main regions

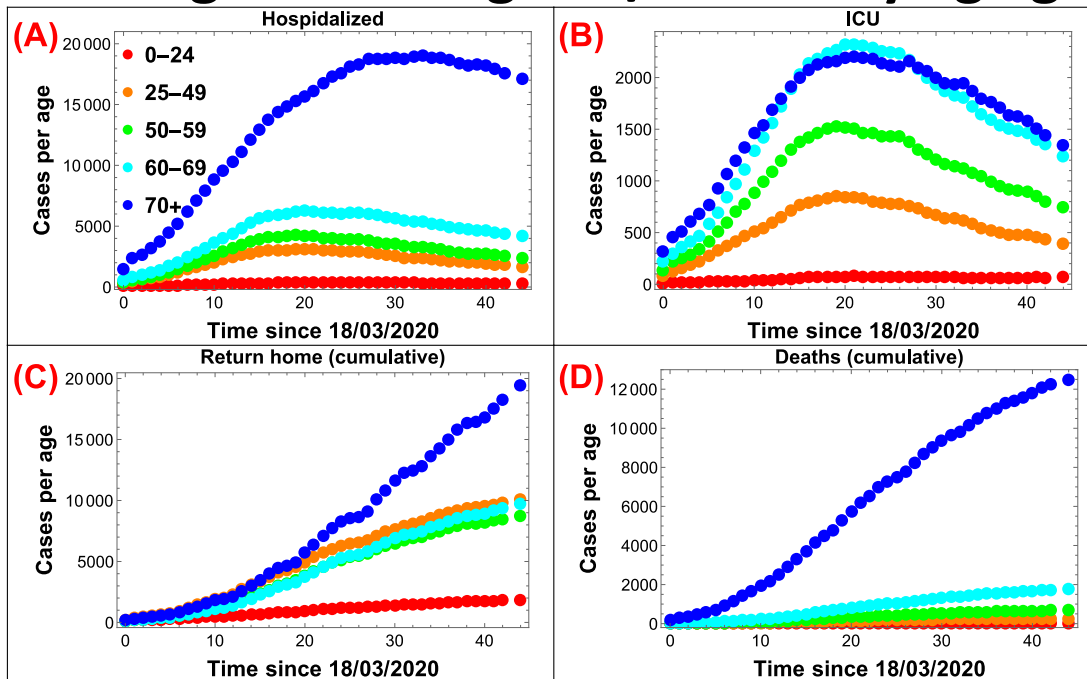
Nombre quotidien de nouvelles admissions en réanimation pour covid-19
les 10 premiers (régions)



Source : *Etablissements de santé déclarant des cas confirmés de COVID-19*

Monitoring 5 Main regions/France by age groups

France



Age group

0-24: group 1

25-49: group 2

50-59: group 3

60-69: group 4

>70: group 5

5 Regions

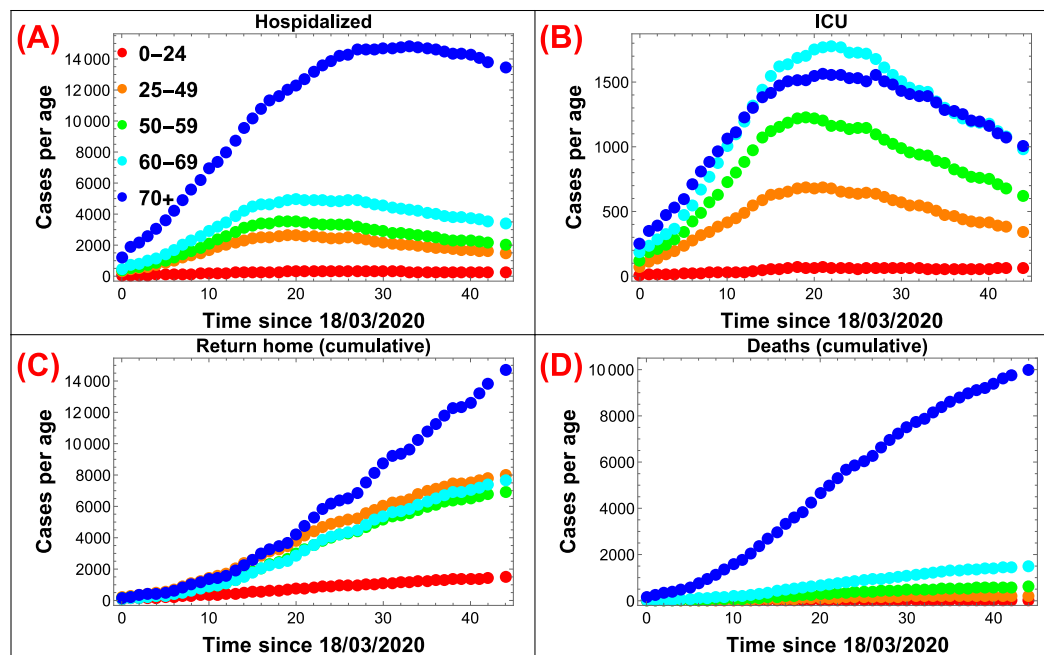
Île-de-France

Grand-Est

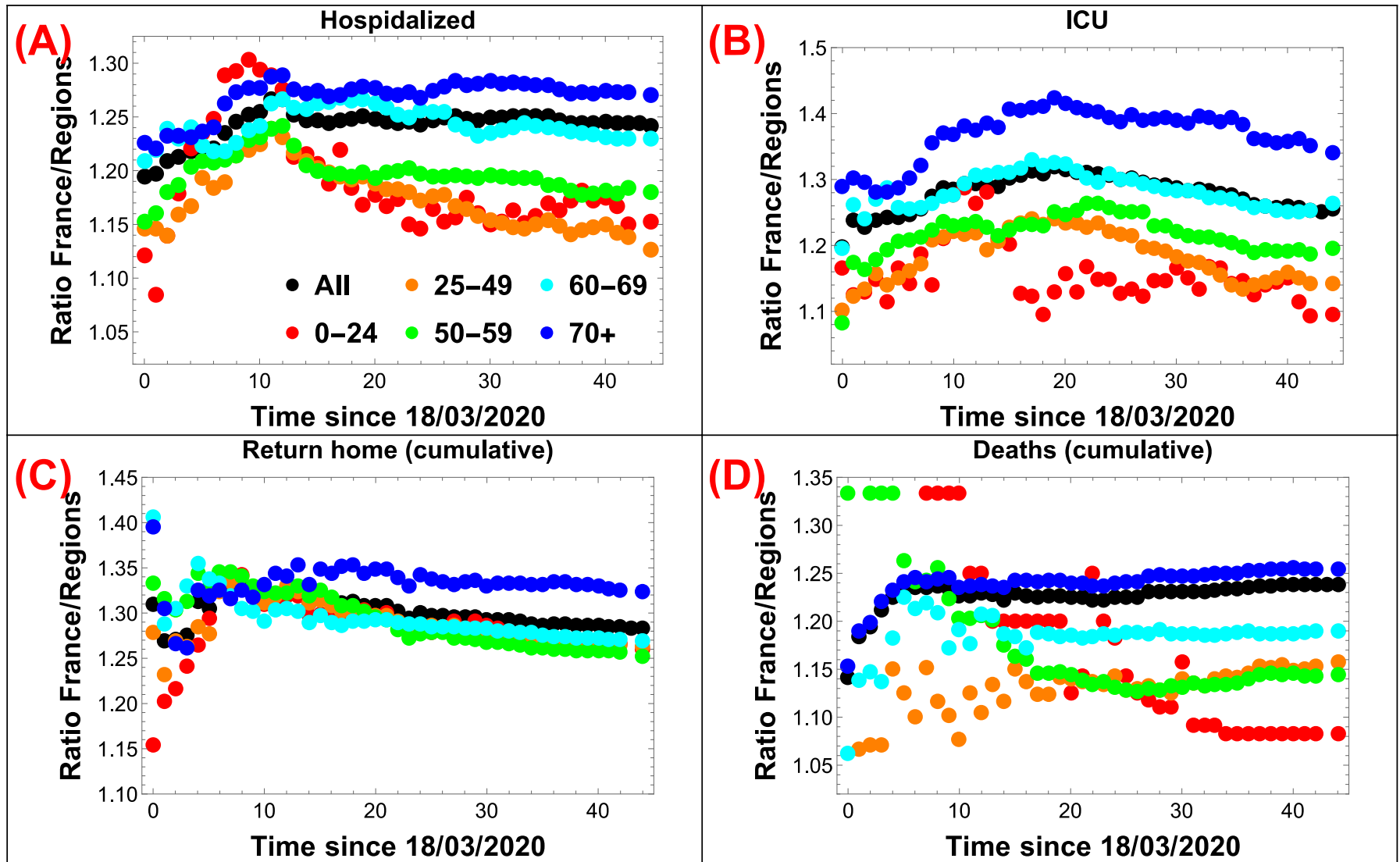
Auvergne-Rhône-Alpes

Hauts-de-France

Provence-Alpes-Côte d'Azur



Ratio France and 5 Regions (ratio of the curves in the previous slide)



Conclusion: ratio of various parameters is almost constant. We decided to focus on 5 most impacted regions and when possible extrapolate to France.

Construction of a multiscale rate Model

- Flexible compared to SIR and extension**
- Can account for changes in decisions occurring in hospitals**
- Can be updated in real time**

Model construction

0-24: group 1

25-49: group 2

50-59: group 3

60-69: group 4

>70: group 5

- Group 1—5: age group \rightarrow filled
- Infectious Category: 1—7: hospitalized, infected, recovered, etc...
- Time evolution from day $n \rightarrow n+1$
- Time after infection
- Time spent in each category.



Ingredient of the model

- Discrete Dynamics

$$S(n+1|k) = S(n|k) - I_{new}(n|k)$$

$$I(n+1,1,1|k,1) = I_{new}(n|k)$$

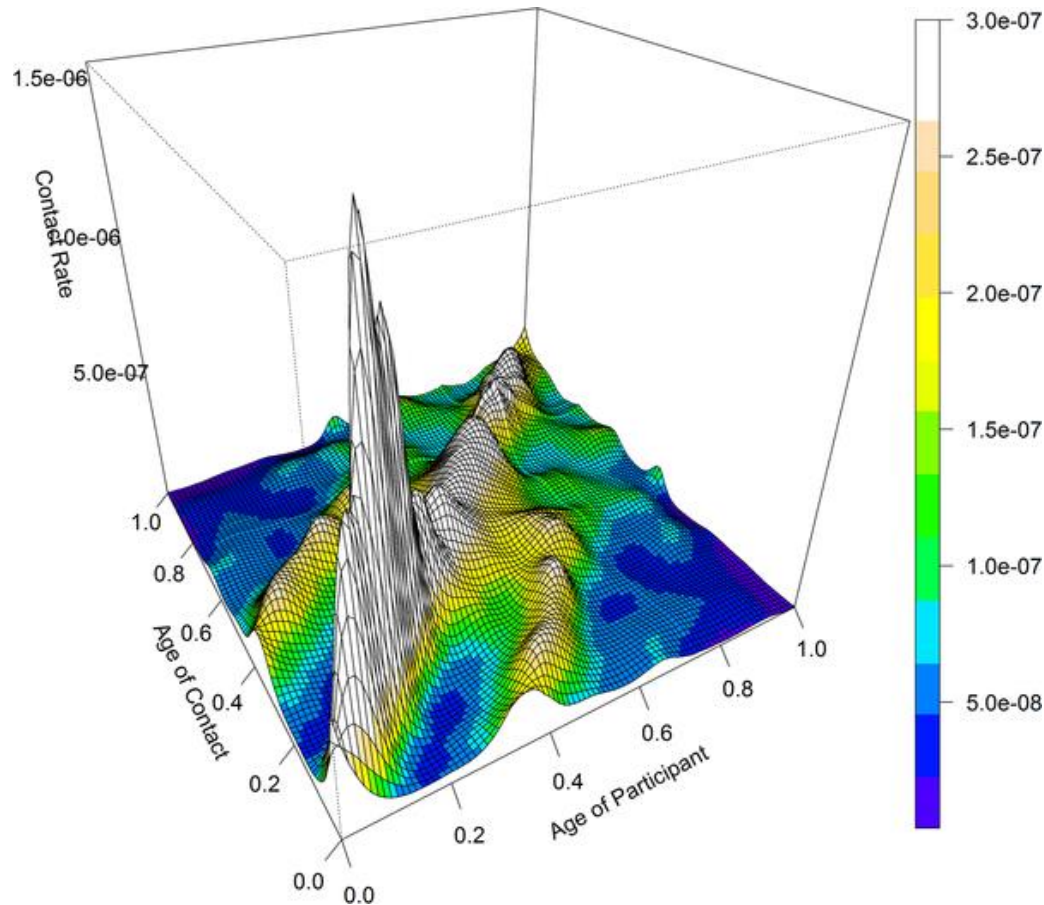
$$I(n+1,m+1,l+1|k,j) = p(n,m,l|k,j,j)\Sigma(n,m,l|k,j)$$

$$I(n+1,m+1,1|k,j) = \sum_{l,j' \neq j} p(n,m,l|k,j',j)\Sigma(n,m,l|k,j').$$

$S(n|k)$ Susceptible persons at day n in category k —group age

$I(n,m,l|k,j)$ Number of infected persons belonging to group k from category j at day n and at date m after infection and at day l in infection category j .
Newly infected person are in category 1.
Categories are hospitalized, infected, recovered, etc...

Contact matrices according to location



Reduced to 5*5 matrix

Béraud G, Kazmerczak S, Beutels P, Levy-Bruhl D, Lenne X, et al. (2015) The French Connection: The First Large Population-Based Contact Survey in France Relevant for the Spread of Infectious Diseases. PLOS ONE 10(7): e0133203.

<https://doi.org/10.1371/journal.pone.0133203>

<https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0133203>

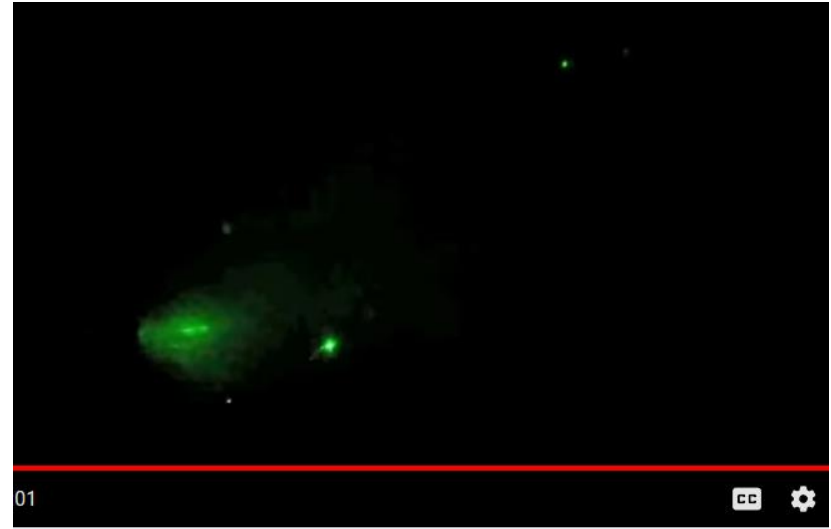
Effect of Mask



Wearing Mask



No Mask



With Mask

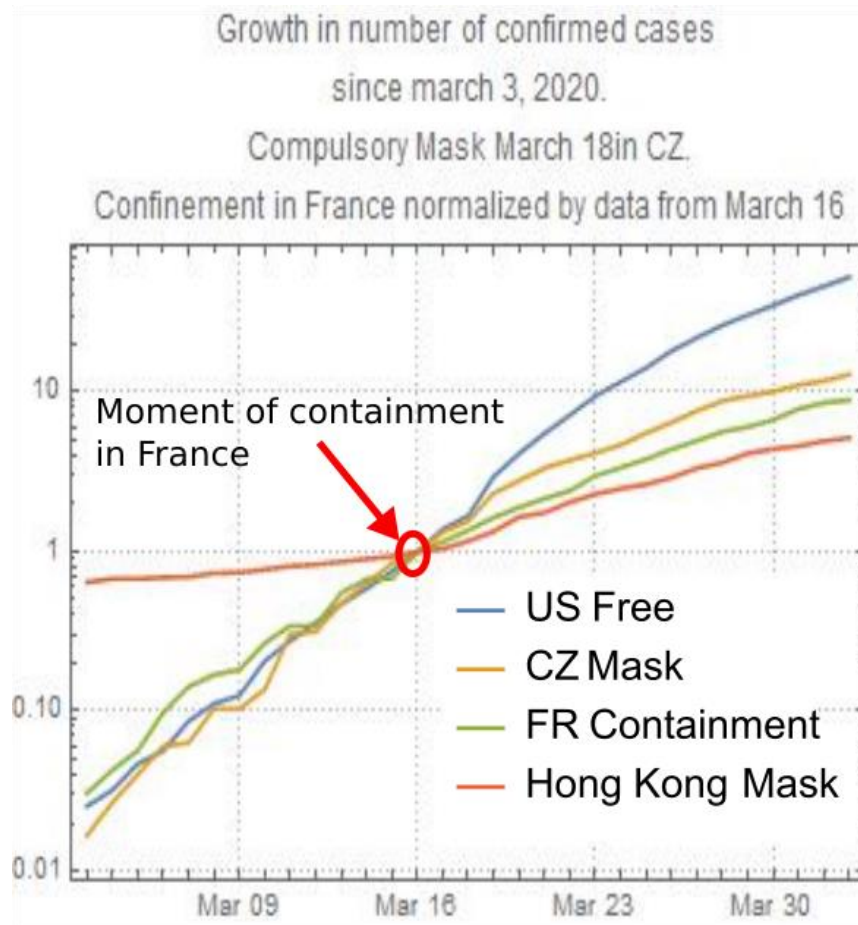
May7 2020

Anfinrud, P., Stadnytskyi, V., Bax, C. E., & Bax, A. (2020). Visualizing Speech-Generated Oral Fluid Droplets with Laser Light Scattering. *New England Journal of Medicine*, NEJMc2007800. <https://doi.org/10.1056/NEJMc2007800>

https://www.youtube.com/watch?time_continue=74&v=UNHgQq0BGLI&feature=emb_title

Conclusion: Masks reduce drastically projection

Wearing Masks: statistical evaluation



COVID-19 Prediction/Modeling/Discussion

<https://www.ibens.ens.fr/IMG/pdf/statistical-analysis2-0904.pdf>

Conclusion: Masks can stabilize the growth of the pandemic similar to confinement

Wearing Mask: reduction contact with fine and coarse particles

Definition: “coarse” >5 μm , “fine” <5 μm (Brownian particles)



Conclusion:

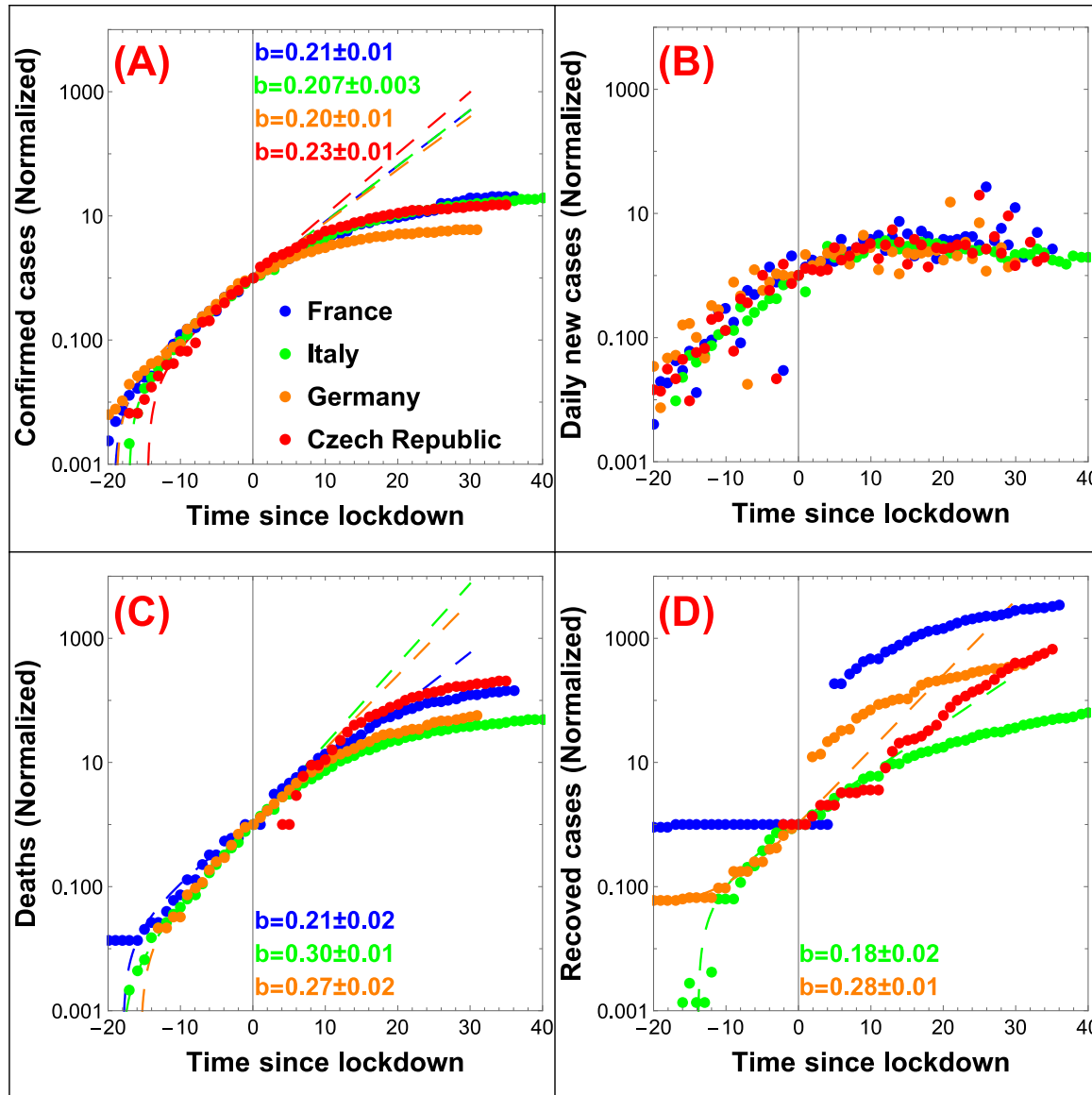
1-Fine particles contained 8.8 (95% CI 4.1 to 19) fold more viral copies than did coarse particles.

2-Surgical masks reduced viral copy numbers in the **fine fraction by 2.8 fold** (95% CI 1.5 to 5.2) and in the coarse fraction by **25 fold** (95% CI 3.5 to 180). Overall, masks produced a **3.4 fold** (95% CI 1.8 to 6.3) reduction in viral aerosol shedding

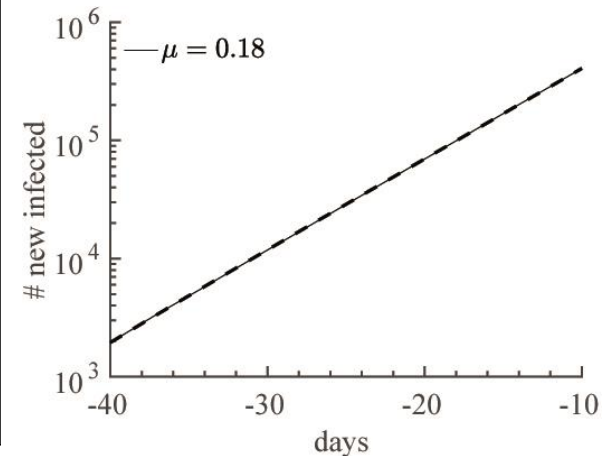
3-Hypothesis: *Mask could reduce the R_0 by 2.8: we tested a reduction between 2.5 and 3*

- **Method:** measured viral copy number using quantitative RT-PCR, and tested the fine-particle fraction for culturable virus.
- **Refs:** Milton, D. K., Fabian, M. P., Cowling, B. J., Grantham, M. L., McDevitt, J. J. (2013). Influenza virus aerosols in human exhaled breath: particle size, culturability, and effect of surgical masks. PLoS pathogens, 9(3).

Extracting parameters from data

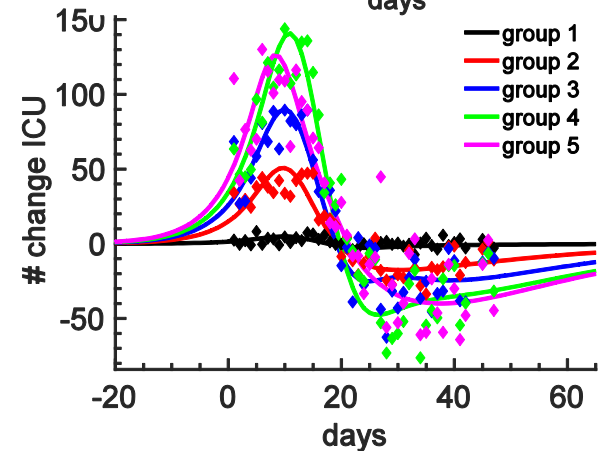
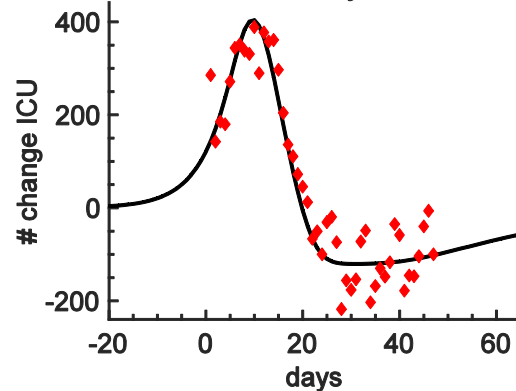
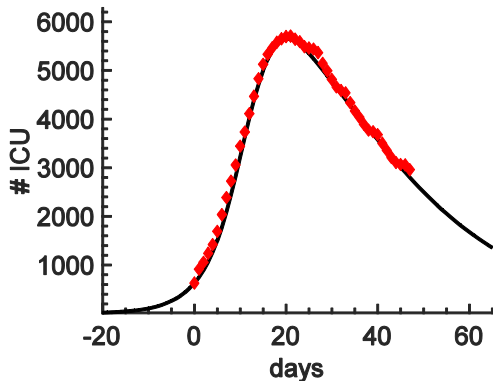
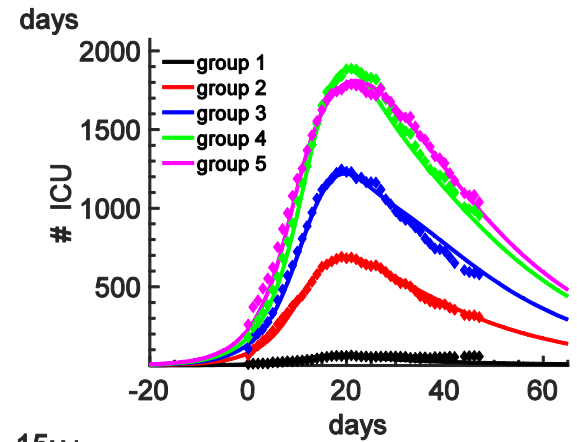
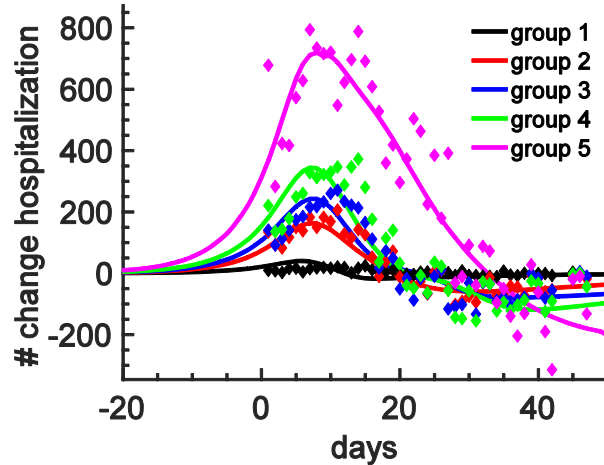
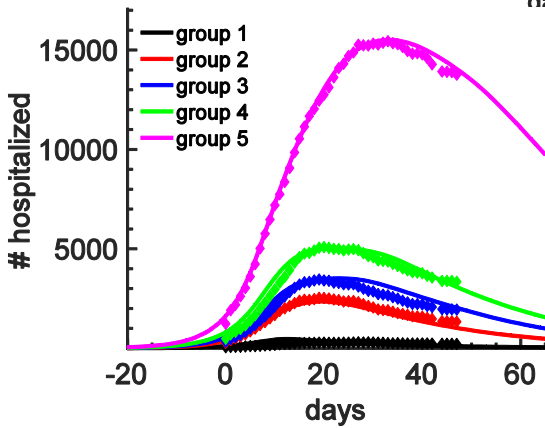
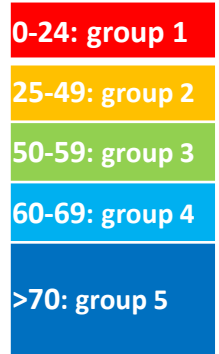
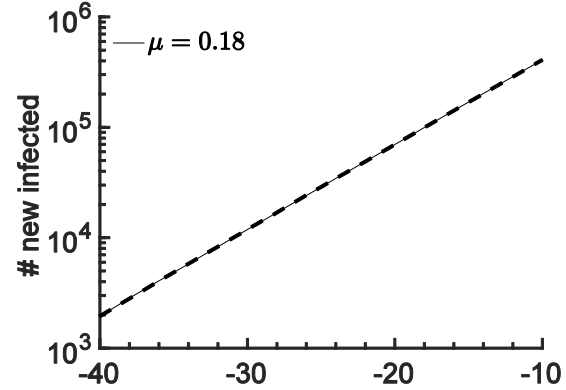
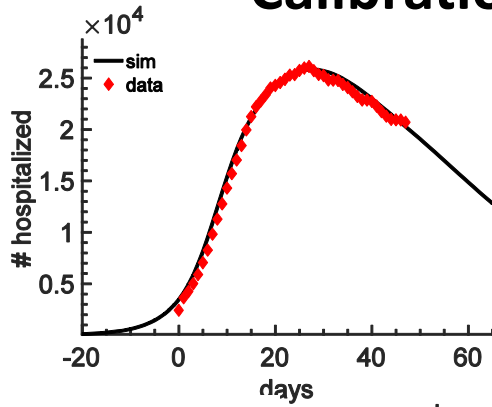


Lockdown:
 France: 17th March
 Italy: 9th March
 Germany: 22nd March
 Czech Republic: 18th March



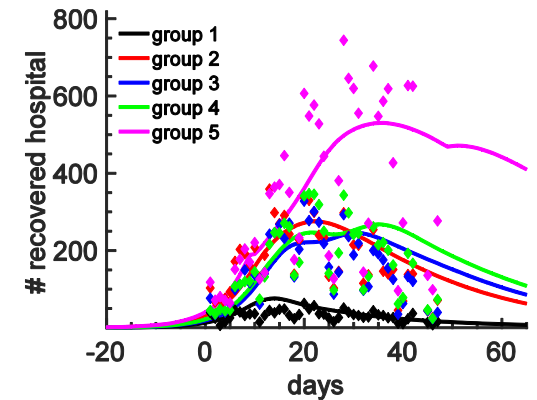
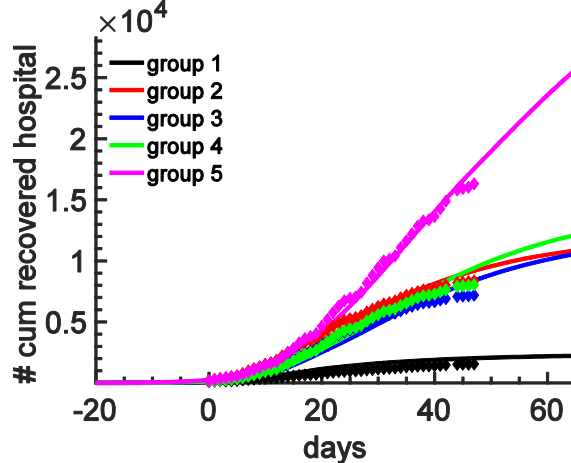
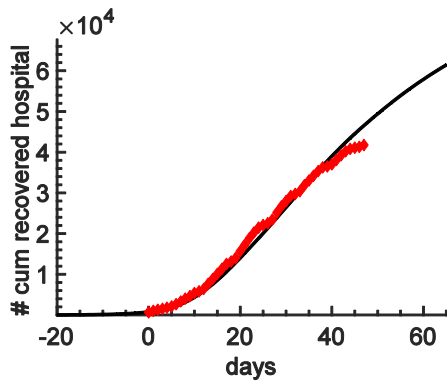
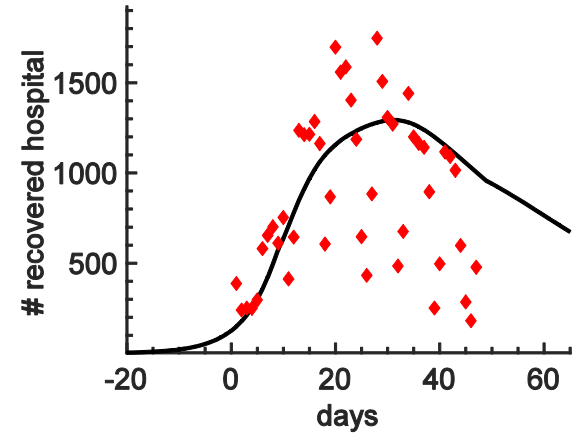
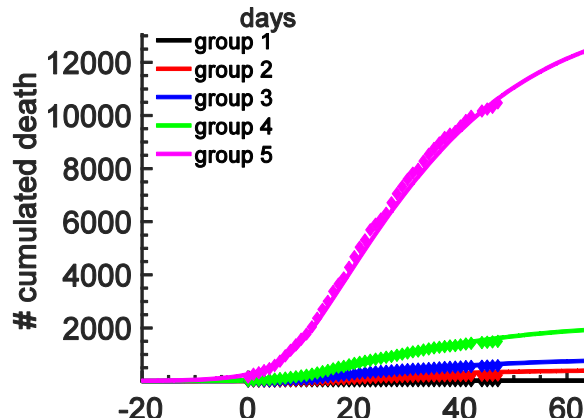
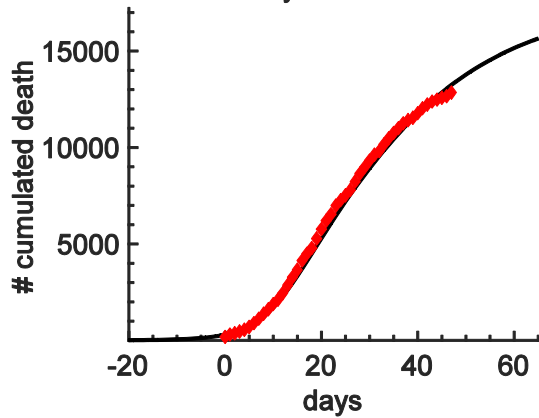
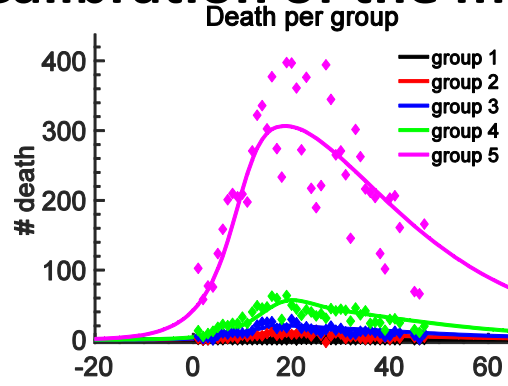
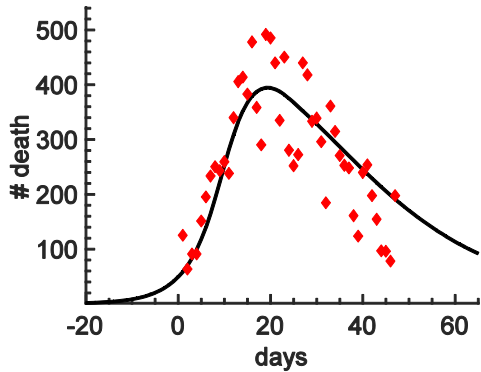
Rate of confirmed cases: exponential growth $f(t) \sim \exp(bt)$ $b=0.2$

Calibration of the model to the data



Conclusion: model accounts for the initial exponential growth, fit calibration at instantaneous time, cumulative (total fluxes)

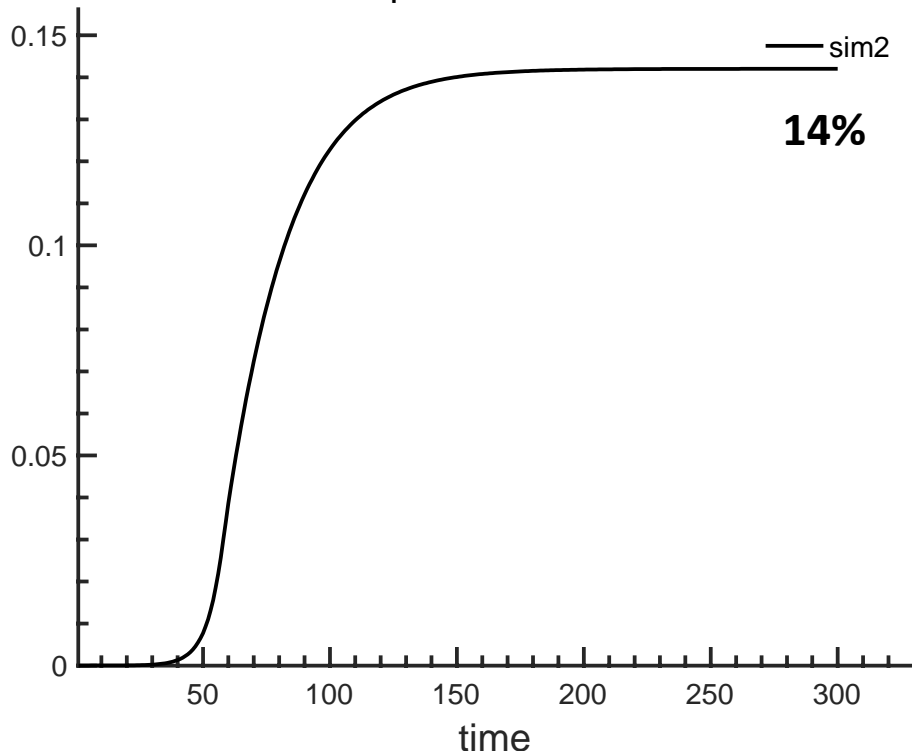
Calibration of the model to the data



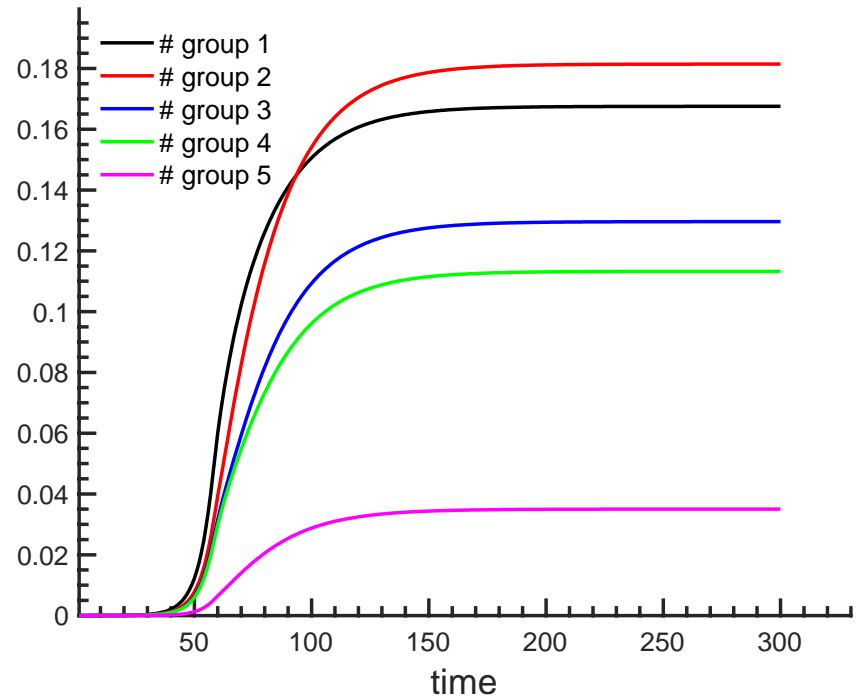
Conclusion: the model accounts for various variables (per age group) simultaneously during a period before and during confinement for several category and age groups.

No deconfinement scenario: simulations for 300 days with

Proportion infected



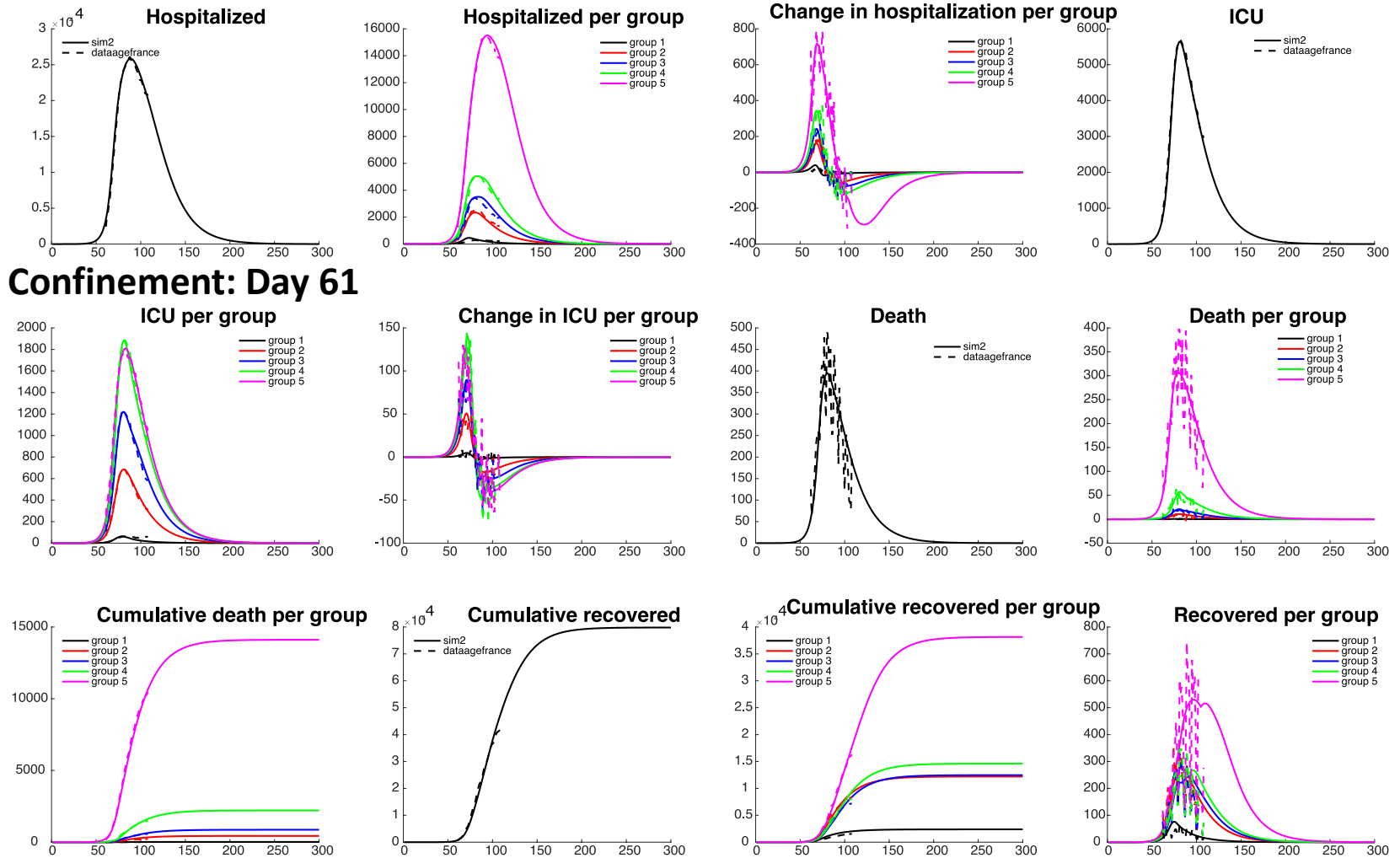
Fraction infected



Conclusion:

1. Small fraction of infected
2. Large variability among age group
3. Latter fluctuation can recreate quickly a pandemic: no collective immunity acquired.

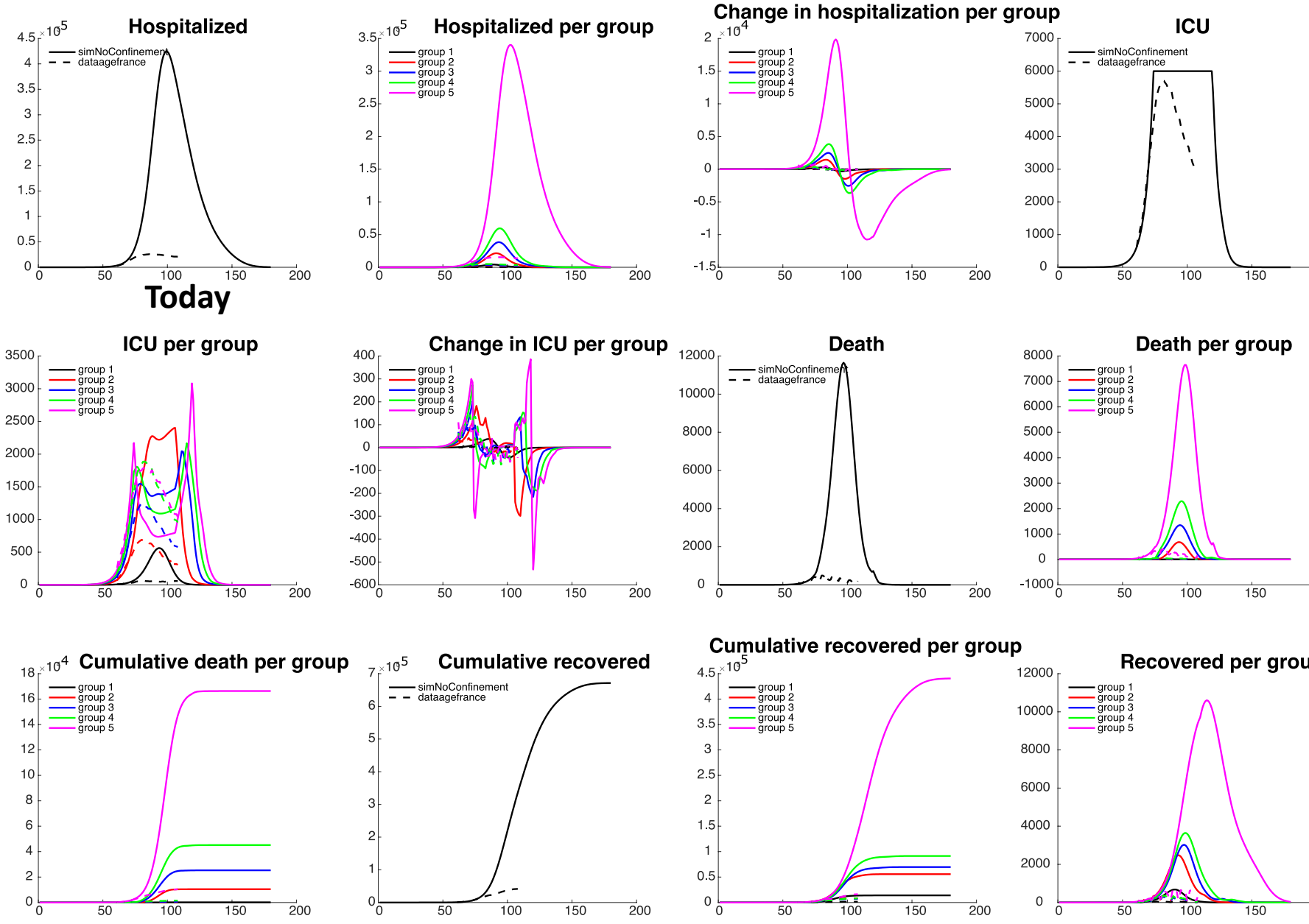
No deconfinement scenario: Simulations for 300 days with



Confinement: Day 61

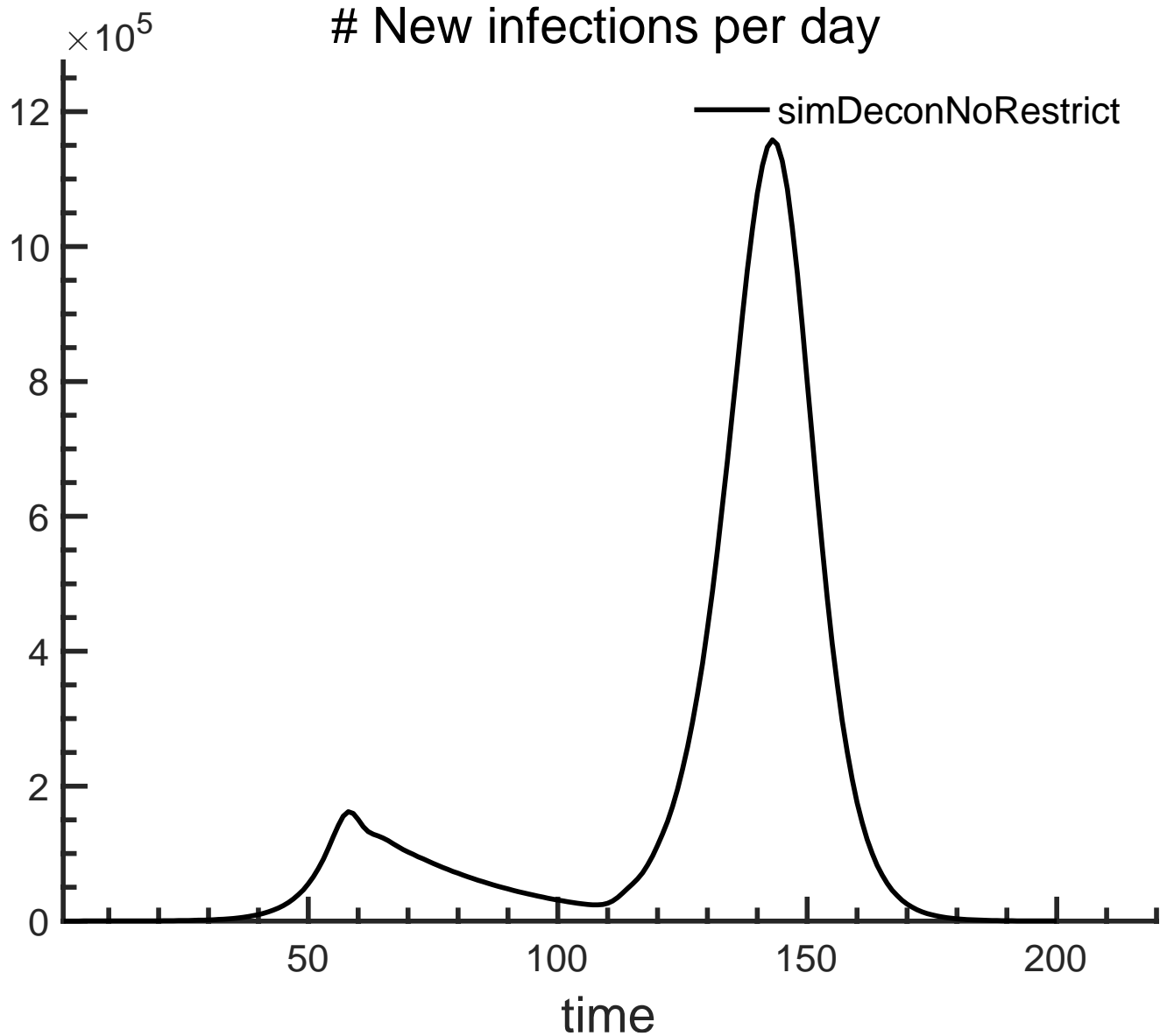
Conclusion: pandemic is fully controlled

Scenario: Had the population not have been confined: simulations for 200 days



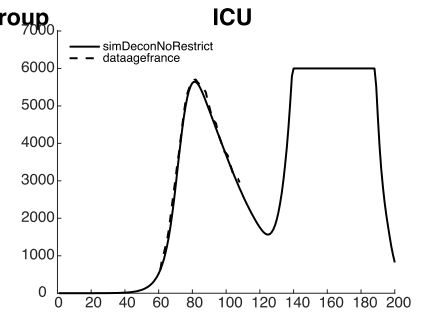
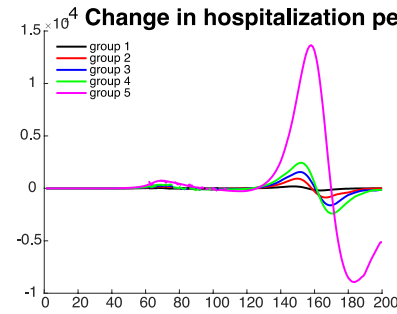
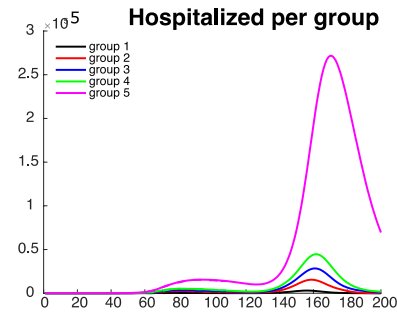
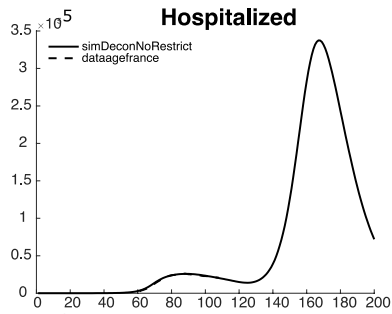
Conclusion: major consequence on destabilization of death and healthcare system

Return to normal social interactions as before confinement: Deconfinement occurring May 11 and

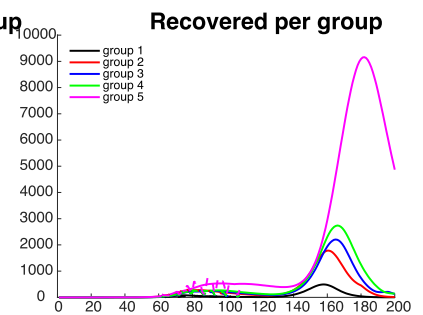
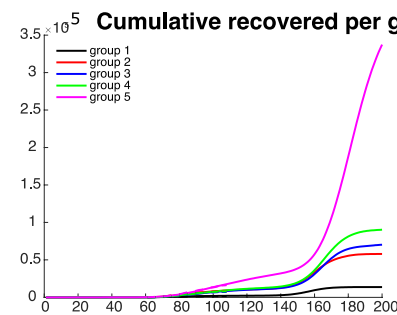
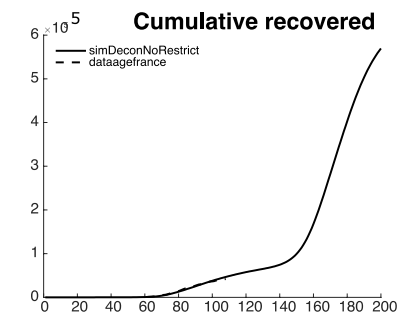
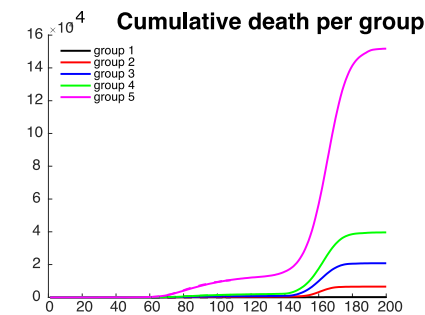
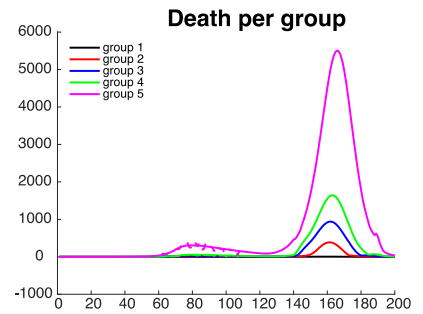
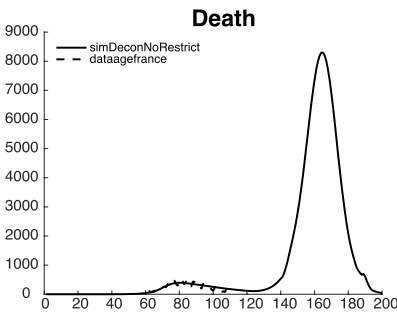
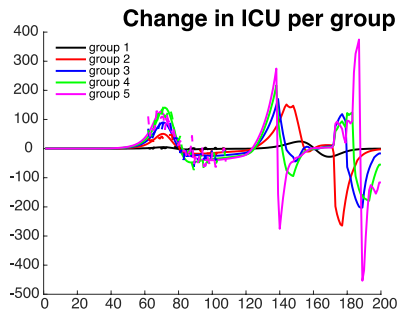
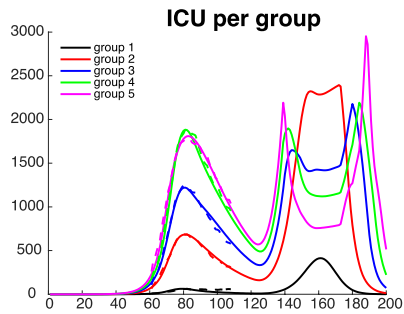


Conclusion: at the peak, the number of infection per day is 6 times the one observed in April

Deconfinement occurring May 11 and return to normal social interaction as before confinement

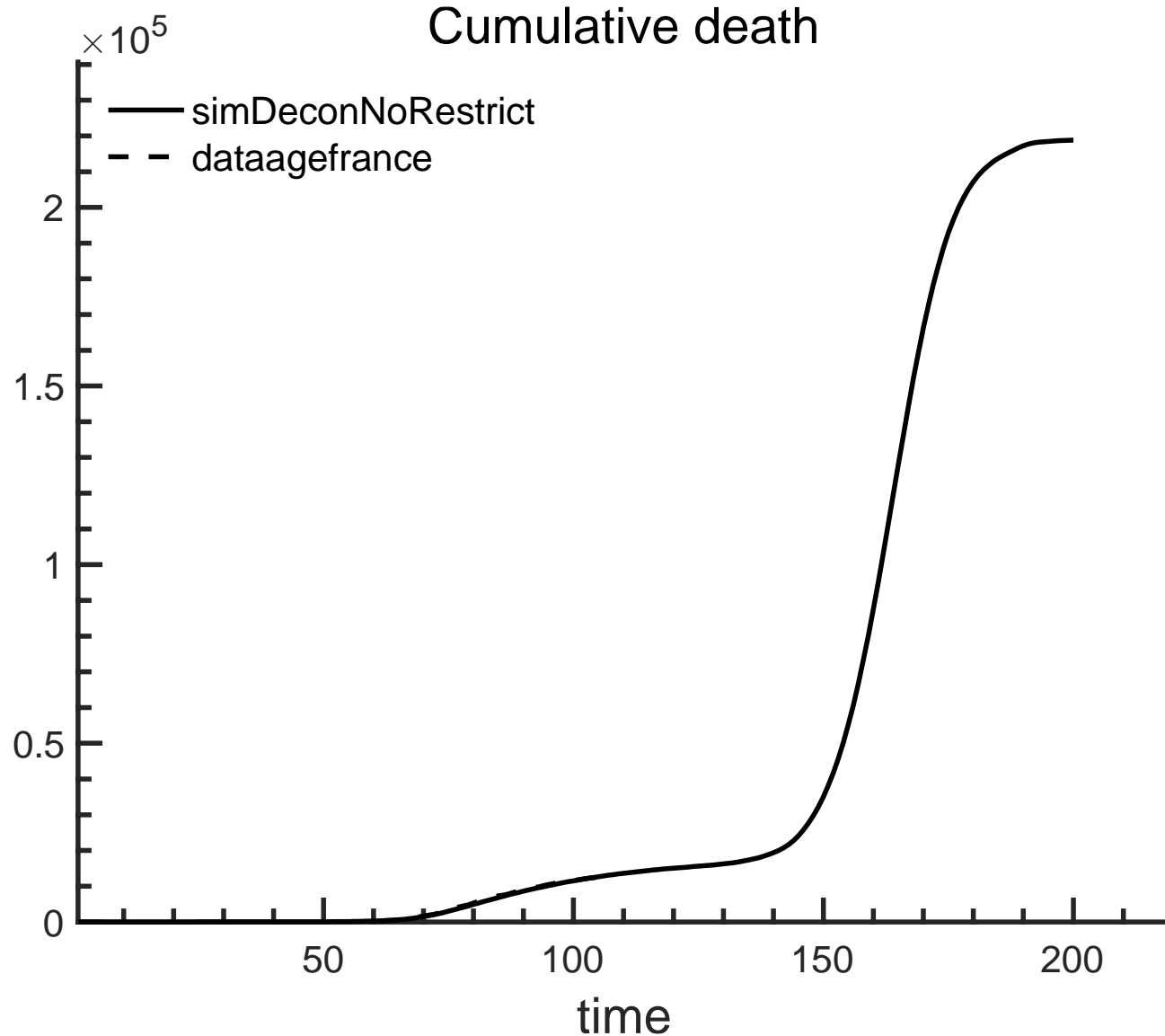


Confinement: Day 61



Conclusion: ICU saturated for 50 days, with a need 320 000 hospitalizations at the peak

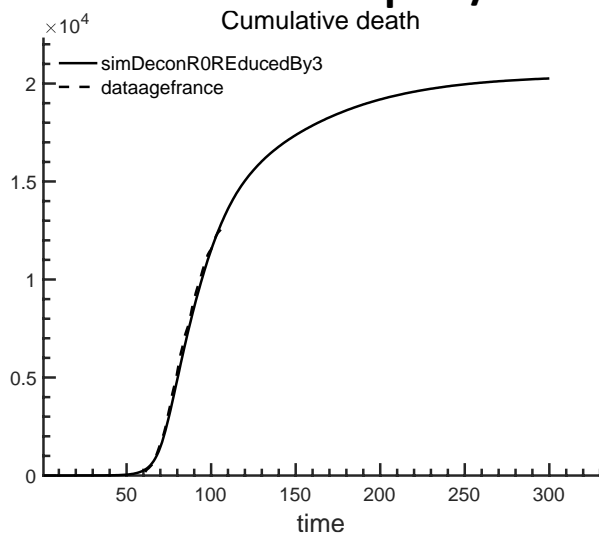
Deconfinement occurring May 11 and return to normal social interaction as before confinement



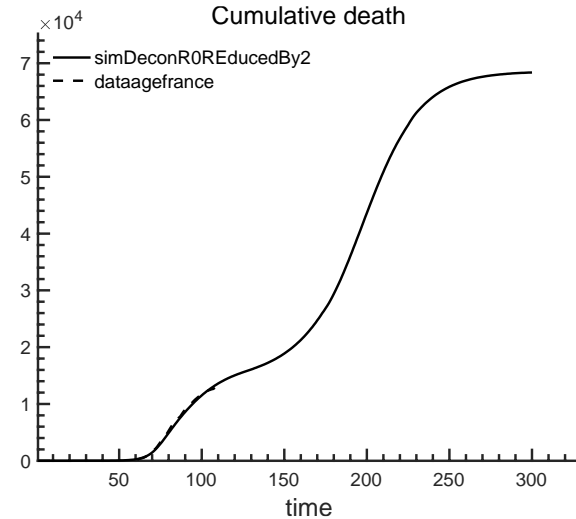
Conclusion: 220 000 deaths for the 5 regions after 200 days

Deconfinement occurring May 11 with social contact reduced

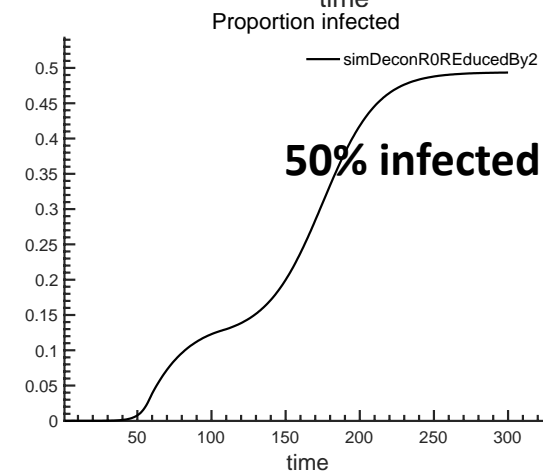
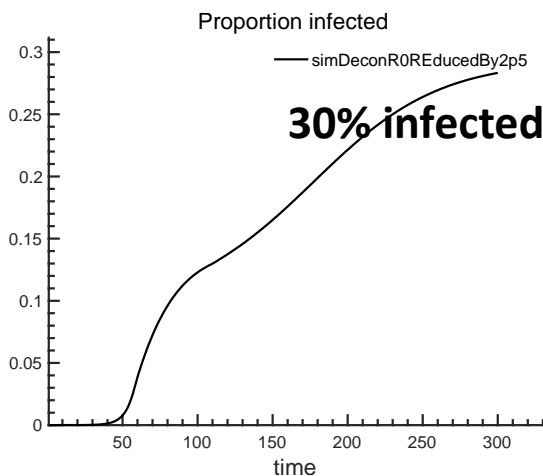
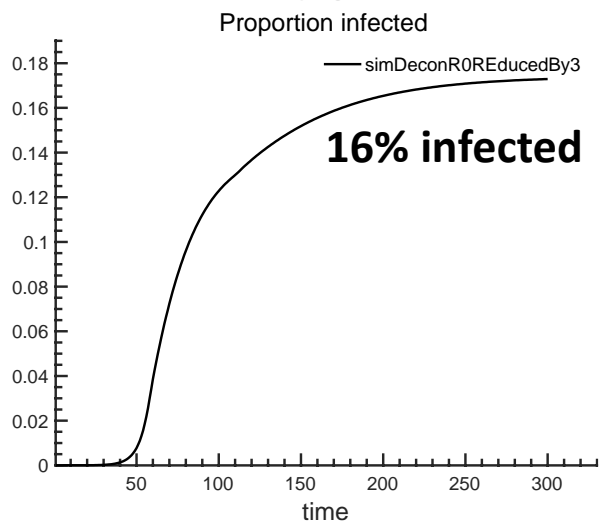
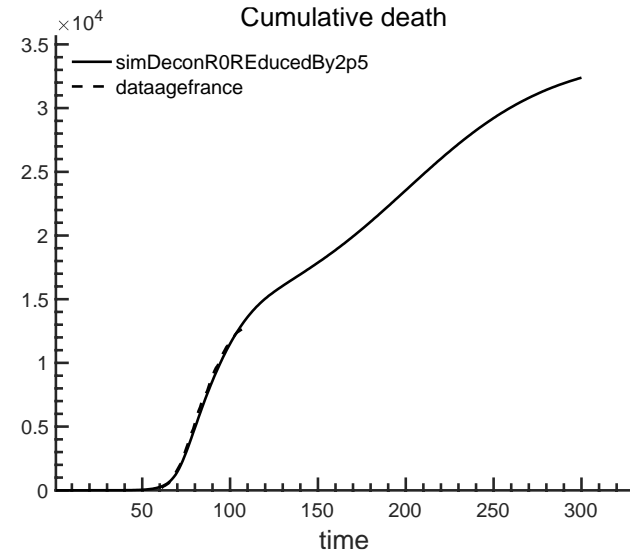
Measures to keep R0/3



keep R0/2.5



Keep R0/2



Conclusion: after 300 days,

total number of deaths (5 regions): R0/3the 21 000 (in hospital)

R0/2.5 35,000

R0/2 70.000

Main conclusions of the model

Main conclusion of the model

- **At time of deconfinement:** 13% of the 5 most critical regions: ~4.6 Millions will be infected by Monday May 11th 2020
- **Scenario of no-deconfinement:** 240 days after confinement: 14% of people will be infected but the pandemic could persist at low noise.
- **Scenario If NO confinement were made:** 12000 deaths/day at the peak. A total of 250, 000 deaths in the 5 regions and probably 450,000 around the country. 87% of the population would have been infection. ICU would have been saturated for 50 days.
 - group-age 0-24: 95% would be infected with 120 deaths/(5 regions)
 - group-age >70: 40% infected with 160 000 total death/(5 regions)

Main conclusion of the model

- **Deconfinement with no restriction: a large second peak is expected mid-July.** 220 000 deaths for the 5 regions after 200 days. ICU saturated for 50 days, with a need 320 000 hospitalizations at the peak
- **Deconfinement with restriction R_0 divided by 3:** No second peak: the pandemic is under control. after 300 days, the proportion of infected people tends to 18%. The total number of deaths (5 regions) is expected of 21000 in hospitals.
- **Deconfinement with restriction R_0 divided by 2.5:** 30% of the population is infected at day 300 with an increasing slope. : Cumulative death around 35000.

Main conclusion of the model

In a Deconfinement with restrictions, R_0 between 2.5 and 3 times less than initial R_0 need:

- Predict the dynamics of COVID-19: exponential phase: any fluctuation is largely amplified: Need a constant monitoring. Phase very sensitive to social interaction.
- To monitor continuously to keep any possible exponential explosion of the phase. Possibility to use the present model and hospitalized data
- Number of test per day should match the number of infected of the order of $\sim 100\ 000$.

Acknowledgements

- Thanks: Prod. D. Longrois (Bichat)
- Model can be adjusted for any other country or other French regions but requires careful adjustments.
- Software can be use to estimate R_0 and predict the change in the next few weeks.
- Softwares developed in
 - Matlab
 - Mathematica/Mapple
- Public sources from French government
 - <https://geodes.santepubliquefrance.fr/#c=news>
 - <https://www.data.gouv.fr/fr/datasets/donnees-hospitalieres-relatives-a-lepidemie-de-covid-19/>

END