Estimate of the instantaneous reproduction number (R_t) of an ongoing epidemic incorporating nowcasting and back-projection corrections

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Basic reproduction number (R₀)

- R_o is the basic number (ratio) of reproduction and it characterizes the transmission potential of a disease
- It represents expected number of secondary cases produced by a typical primary case in an entirely susceptible population
- The R₀ of a disease is function of human social behavior and organization, as well as the innate biological characteristics of particular pathogens
 - Transmissibility
 - Contact rate
 - Infectious period
- Constant review
 - Measles: estimate: 12-18
 - Systematic review by Guerra et al (2017): 3-200 (!!)

Instantaneous (effective) reproduction number (R_t,R_{eff})

 R_t is the actual average number of secondary cases per primary case and it is usually a fraction of the R_0 :

$$R_t = R_0 x$$

where, in a simple model, \mathbf{x} is the frequency of susceptible individuals in a population at a given time \mathbf{t}

Why calculate R_{eff}?

It is commonly used to characterize pathogen transmissibility during an epidemic.

The monitoring of R over time provides feedback on the effectiveness of interventions and on the need to intensify control efforts.

To stop an epidemic, R needs to be persistently reduced to a level below 1.

Instantaneous (effective) reproduction number (R_t , R_{eff})

- R_{eff} > 1: number os new cases increases
- R_{eff} = 1: equilibrium of cases (endemic)
- R_{eff} < 1 : number os new cases decreases

 R_{eff} (and R_0) is not a rate! It is a number, a ratio.

Estimate the R_{eff} for a local community, where contacts/transmissions occur

How to estimate the R_{eff}

Ratio between the incidence number of new cases at a time t and the incidence number of new cases at

time *t-1*

day	Incidence number	Ratio N(t)/N(t-1)	
1	10	-	
2	12	1.2	
3	18	1.5	
4	26	1.4	
5	44	1.7	

However we must consider that the expected number of incidence cases at a given time t depends on the number of incidence cases in previous time steps t-s weighted an infectivity function \mathbf{w}_s

$$\mathbf{E}[I_t] = R_t \sum_{s=1}^t I_{t-s} w_s$$

Database and dates

There are 3 important dates for a given a case:

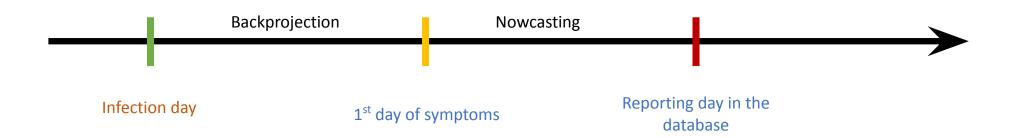


Brazilian national health surveillance of hospitalized individuals (SIVEP-Gripe) we have access to at least 2 of them:

- Date of 1st of symptoms
- Reporting date

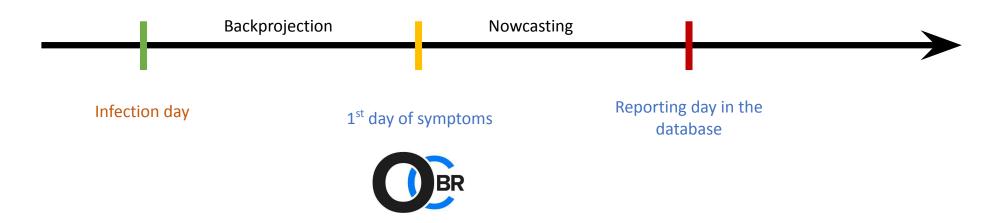
Delays!!

Correcting delays



Correcting delays

• Corrections:



Methods to estimate R_{eff}

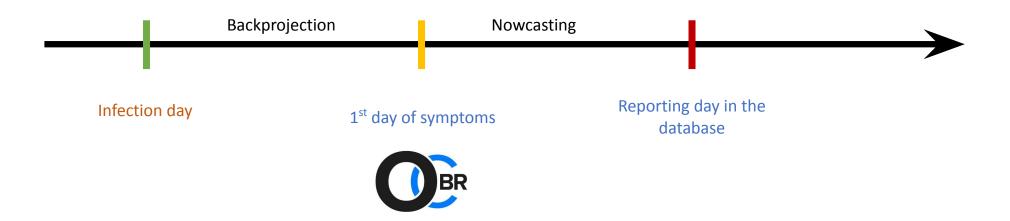
- Robert Koch Institute 2020
- Wallinga e Teunis 2004
- Bettencourt e Ribeiro 2008
- Cori et al 2013

Methods to estimate R_{eff}

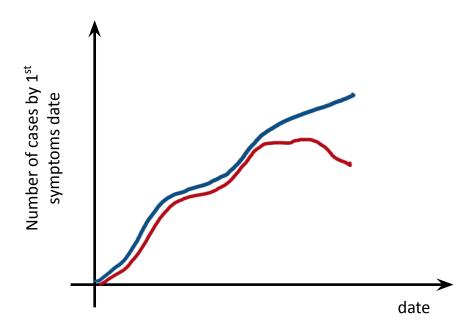
- Robert Koch Institute 2020
- Wallinga e Teunis 2004
- Bettencourt e Ribeiro 2008
- Cori et al 2013
 - Time series of cases
 - Serial interval distribution (trasmission tree) incorporates uncertainty in the distribution the time between the onset of symptoms in a primary case and the onset of symptoms in secondary cases

Correcting delays

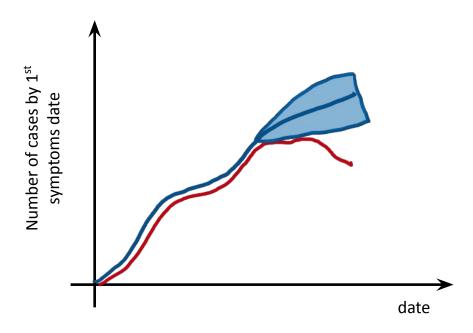
• Corrections:



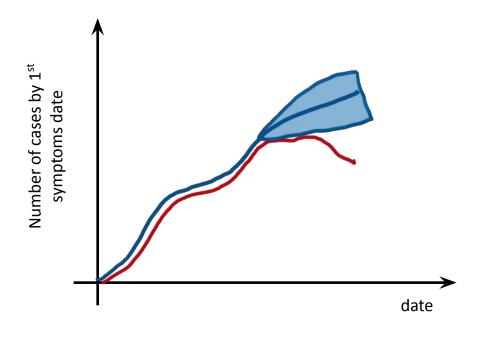
Nowcasting estimates the number of cases that have already occurred (that already have a 1st day of symptoms) but that have not been reported in the database.



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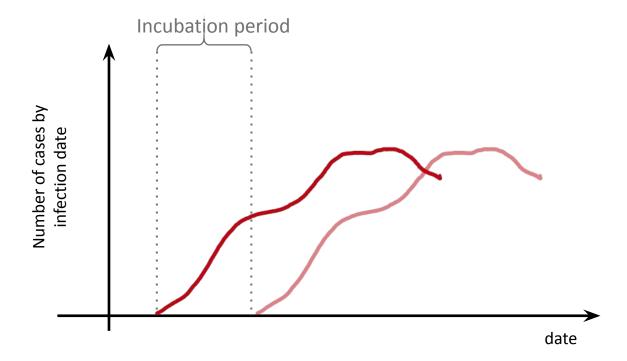


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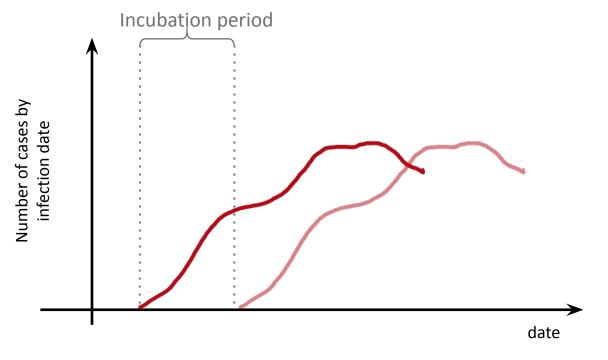


- Nowcasting by Bayesian Smoothing (McGough et al 2020)
- Posterior series

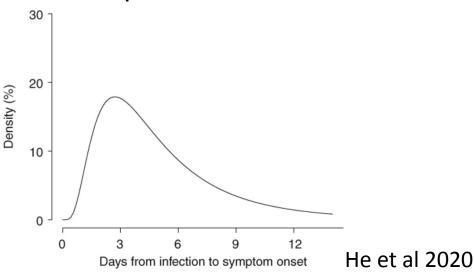
Back-projection estimates the number of cases by infection date, therefore including the incubation period (SARS-CoV2 \sim 5 days)



Back-projection deconvolution estimates the number of cases by infection date, therefore including the incubation period (SARS-CoV2 \sim 4-5 days)



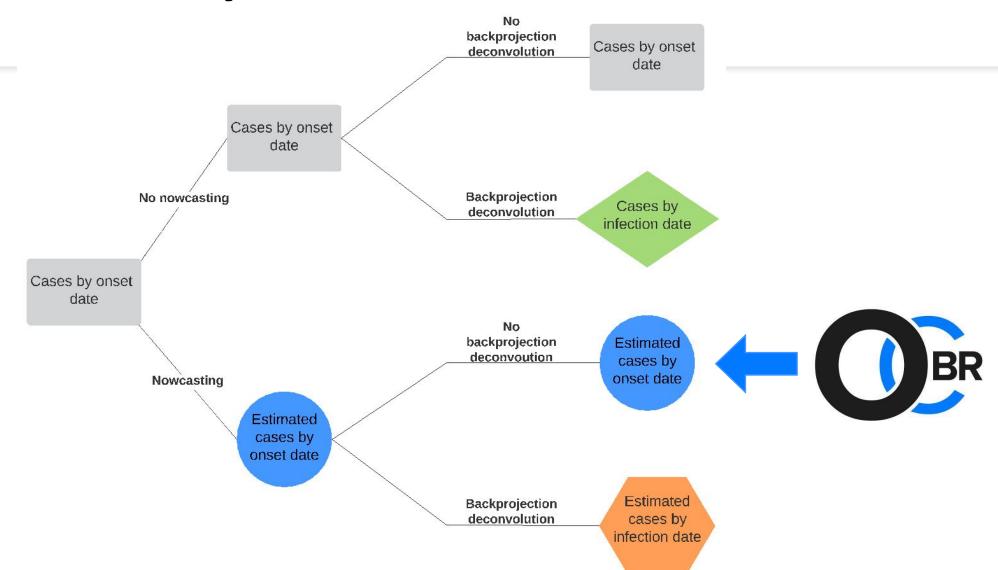
- Expectation-Maximization Smoothing procedure (Becker et al 1991)
- Bootstrap simulation



Credible intervals of the R_{eff}

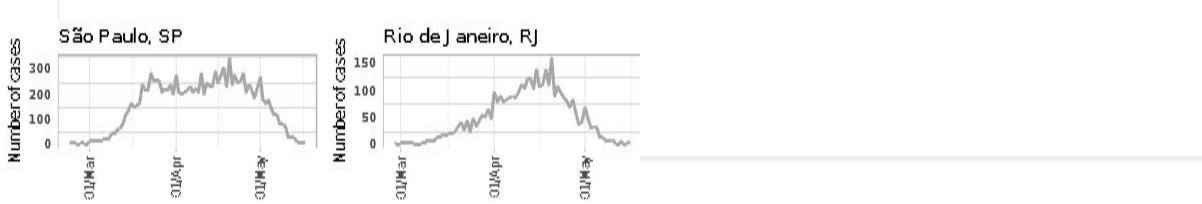
- Nowcasting CI: delays from de reporting system
- Backprojetion CI: delay from the incubation period.
- *Transmission tree CI*: from the method proposed by Cori et al 2013 based on cases that will come from the transmission tree

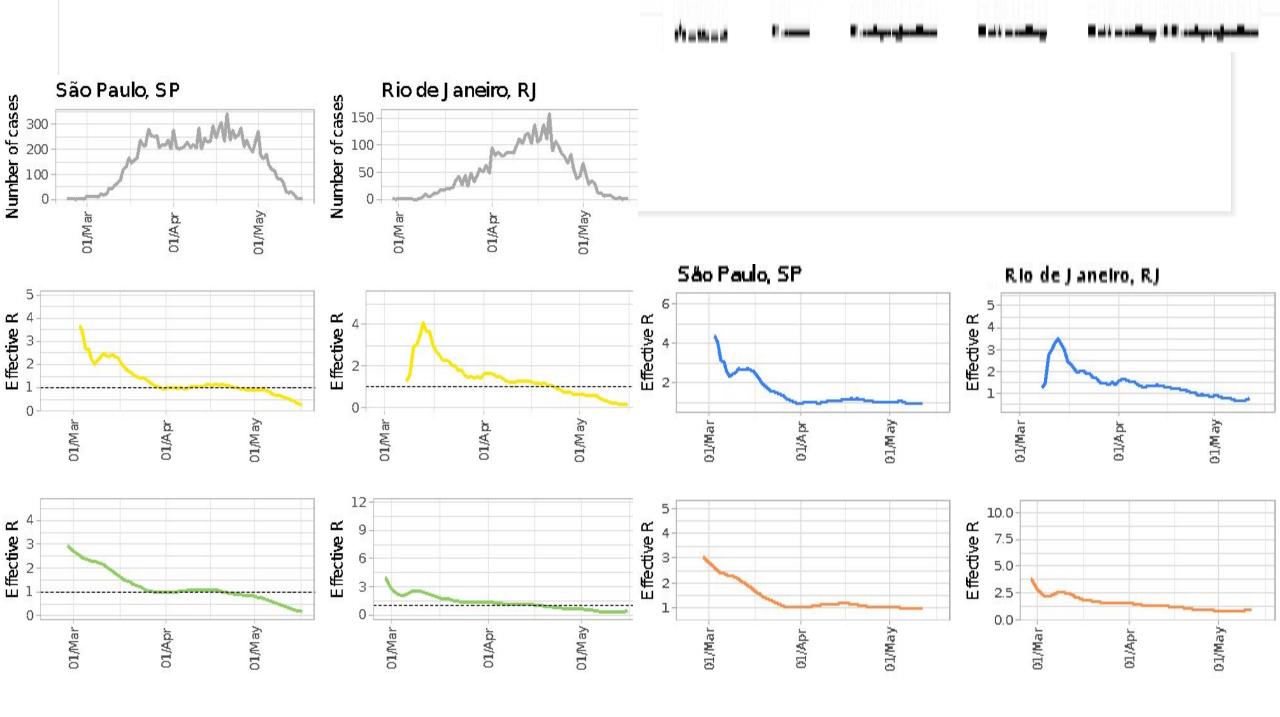
Different delay corrections

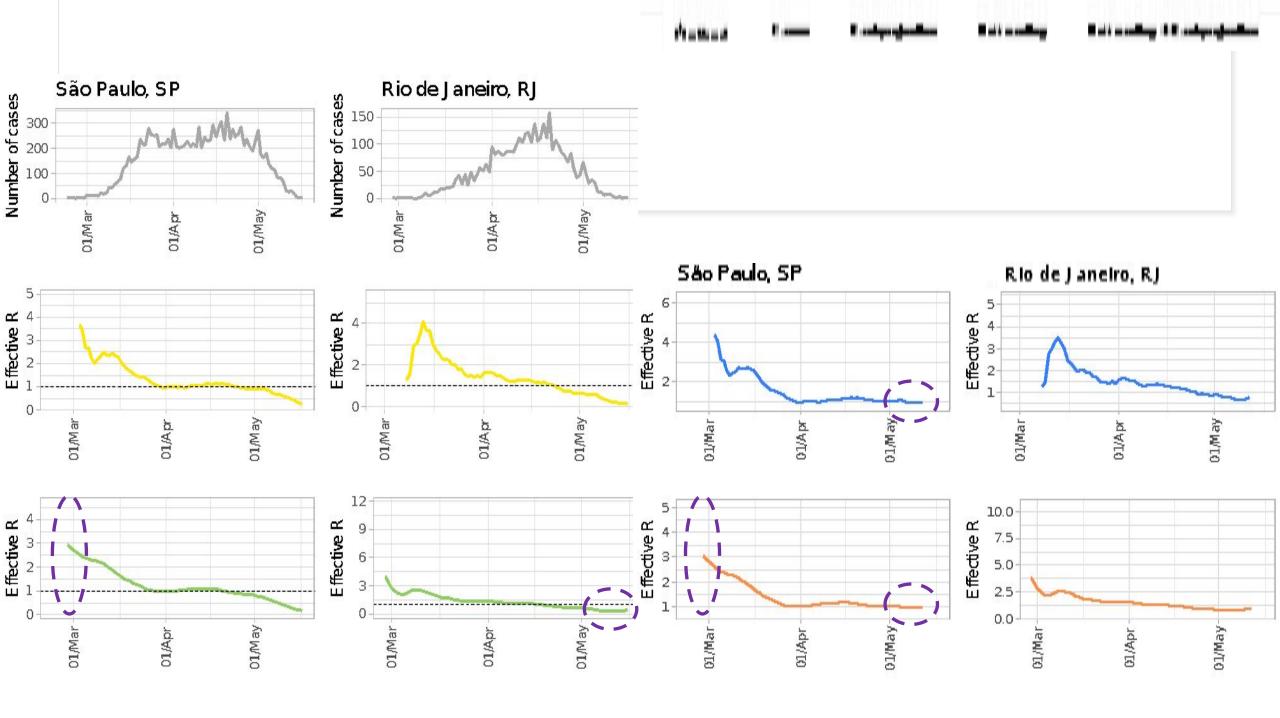


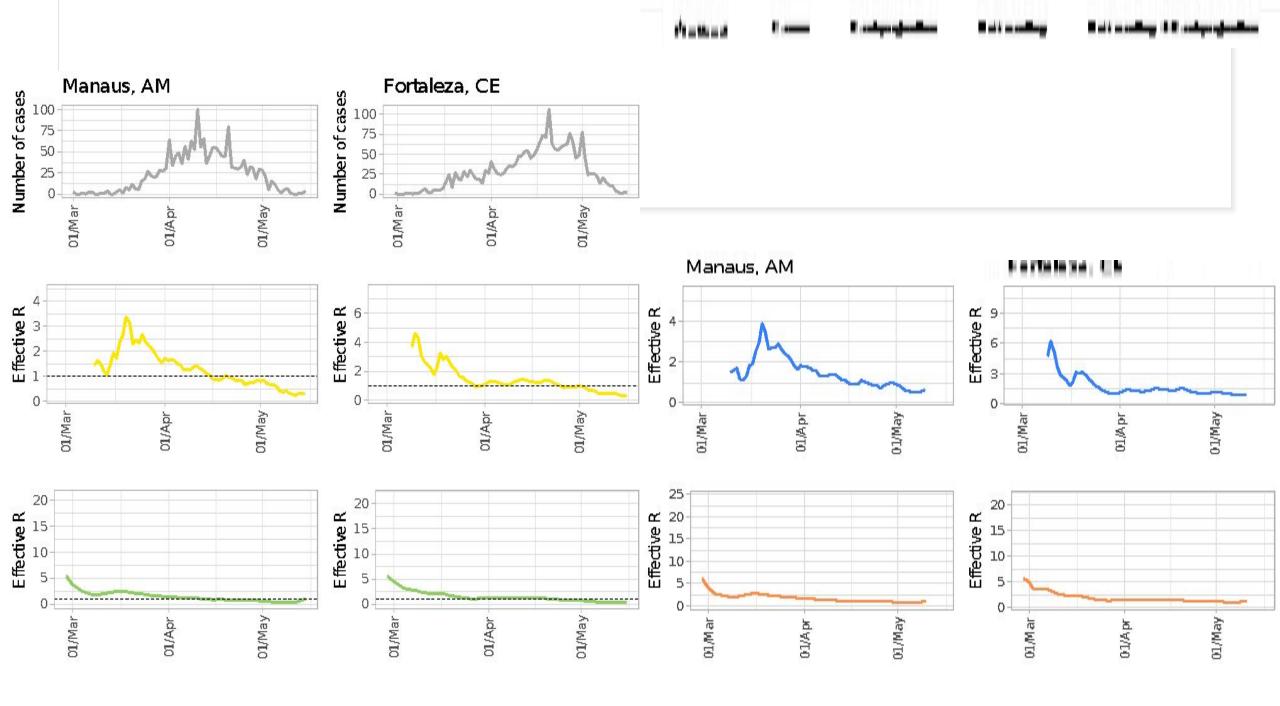
Results

- SIVEP-Gripe (hospitalized cases)
- Positive cases diagnosed by RT-PCR
- Base from May 18,2020 (with a consolidated base of June 30,2020)
- State capitals: São Paulo (SP), Rio de Janeiro (RJ), Manaus (AM), Fortaleza (CE)

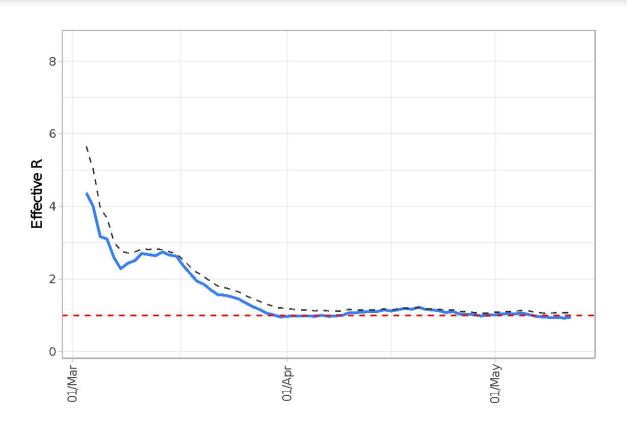








Results

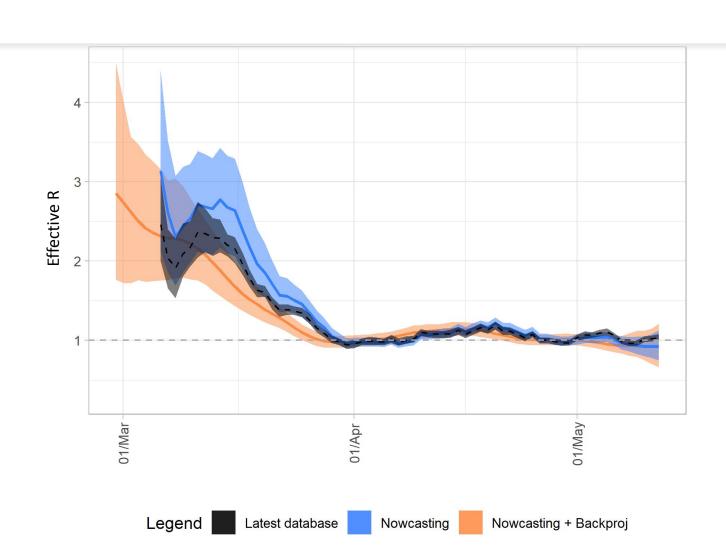


Nowcasting

Consolidated

Method

Results



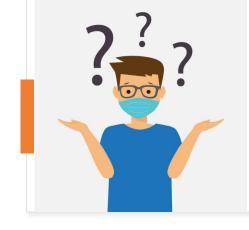
Discussion

- Estimate the R₊ based on the notification date is not accurate
- Important delays that can be corrected: reporting system and incubation period
- Backprojection can affect the end of the R_t time series, decreasing the R_t values due to the lack of data feeding the end of the time series
- The nowcasting increases the CI of the R₊

Discussion

 Decisions made by policy makers and health authorities must consider the CI and not only the mean value. The entire range of the CI must be lower than 1 to be considered controlled.

- When using the back-projection correction, one should avoid using the most recent estimates of R₊ (final temporal series).
- Other metrics should also be taken into account for public health decisions. values of R₊ around 1 associated with a high number of new cases per day can easily fuel a new rise



Questions?

Reff estimate by Cori et al 2013 method

- We use a modified estimate_R function from EpiEstim R package
- We use a sliding window of 6 days
- Serial interval by Nishiura et al 2020 for MCMC estimation for the transmission tree

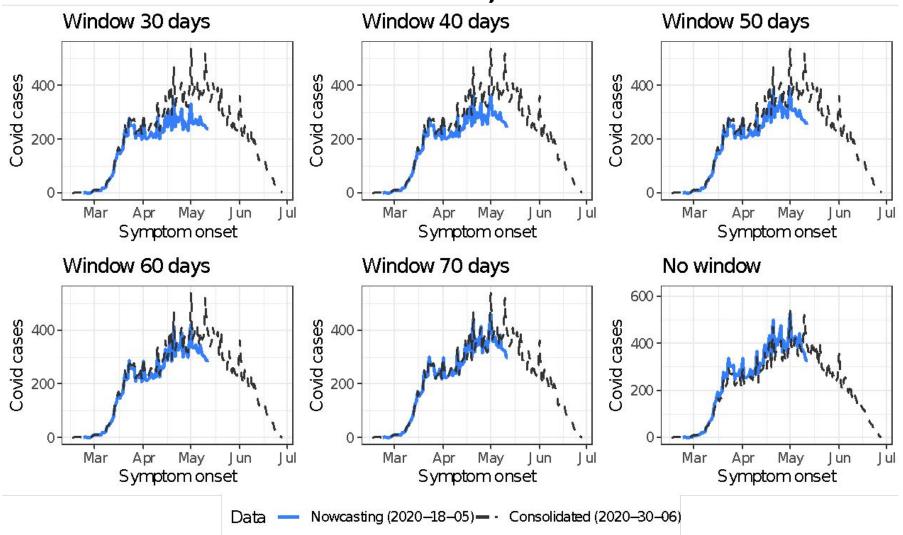
- We use a modified function from the NobBs R package (McGough et al 2020)
- Trim of 5 days in the time series end
- Time window of delay:

São Paulo (SP): 70 days

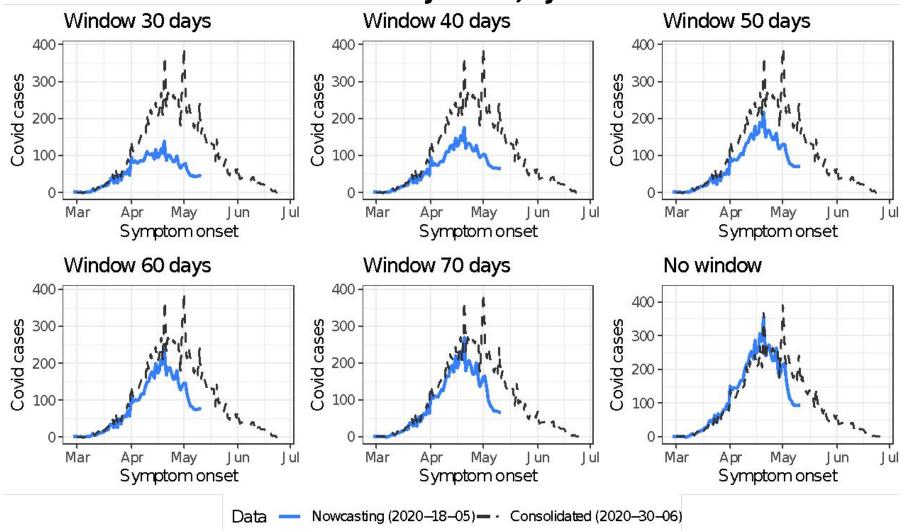
Manaus (AM): 60 days.

Rio de Janeiro (RJ) and Fortaleza (CE) no window.

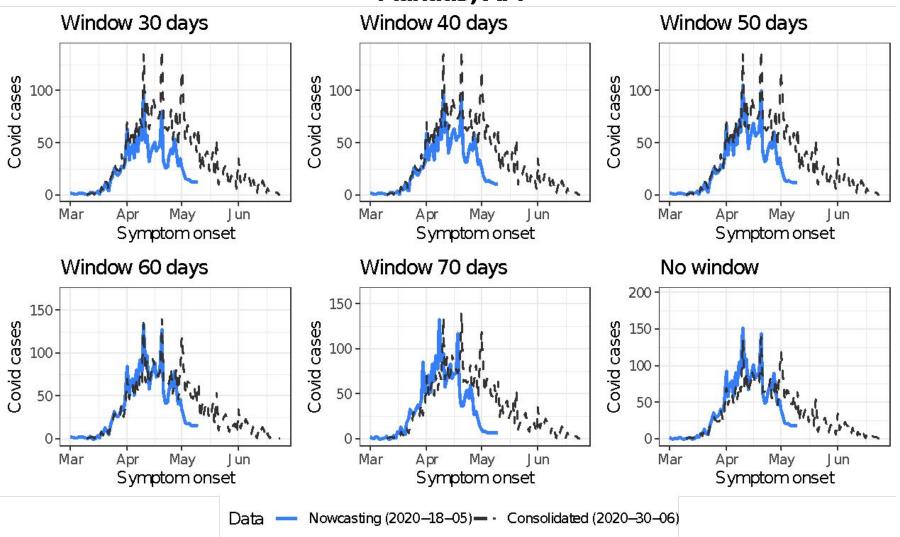
São Paulo, SP



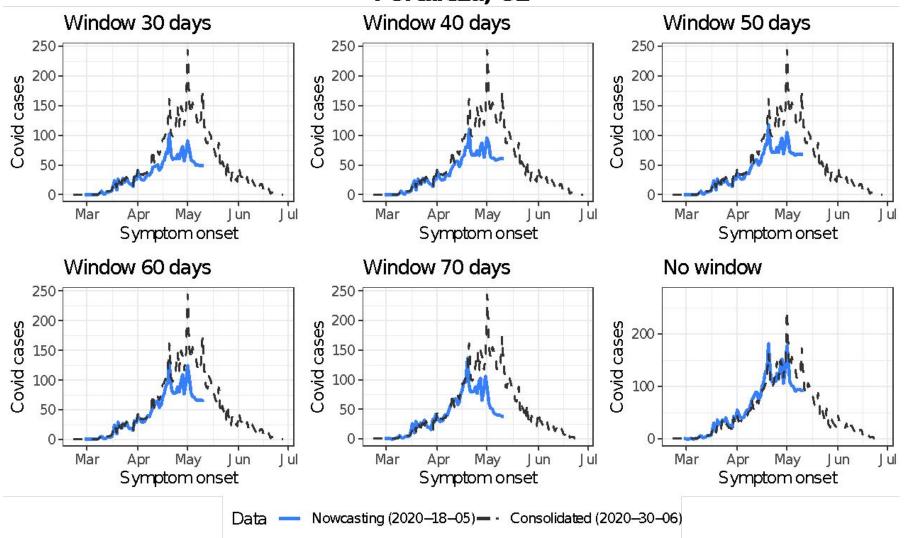
Rio de Janeiro, RJ



Manaus, AM



Fortaleza, CE



• The incidence data is corrected by discounting delays from infection to the first day of symptoms using an incubation distribution defined by a lognormal (mean=5.2 days and 95% CI = [4.1, 7.0] days (He et al 2020).

• In contrast to the nowcasting method, which includes cases considering the delay of the notification system, {\fla the back-projection method just estimate the shift in the time series backwards with no addition of cases

• The Expectation-Maximization-Smoothing procedure of Becker et al (1991) gives a sample estimation of the mean number of individuals infected at time \boldsymbol{t} and is denoted by $\boldsymbol{\lambda}_{\scriptscriptstyle t}$.

 The sample estimation must be accompanied by a confidence interval which can be estimated by a bootstrap method (see Yip 2008).

• We choose to return a sample draw from a Poisson distribution with mean $\lambda_{\!\scriptscriptstyle \perp}$ after we have back-projected our data.

• This procedure is repeated for 1,000 posterior nowcasted series. The sample error for the λ_t computation is incorporated when we consider a large number of back-projected samples of a nowcasted series.

 The complete routine is implemented in C++ and is already integrated with R via the Rcpp package

More information and analysis in our website

https://covid19br.github.io/



Observatório COVID-19 BR

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Contact:

Instagram, Facebook, Twitter: @obscovid19br

Correções do dia de ocorrência dos eventos

• Uma outra correção possível é deslocar a série para trás de acordo com um distribuição do número de dias entre o dia da infecção e o 1o sintoma. O método é geralmente chamado de *Backprojection deconvolution*.

