Construction, calibration and validation of a macro-epidemiological surveillance indicator at a national level from wastewater analysis.

Yvon Maday

Laboratoire Jacques-Louis Lions Sorbonne Université, Paris, Roscoff, France Institut Universitaire de France

Somewhere-World — January 19, 2022

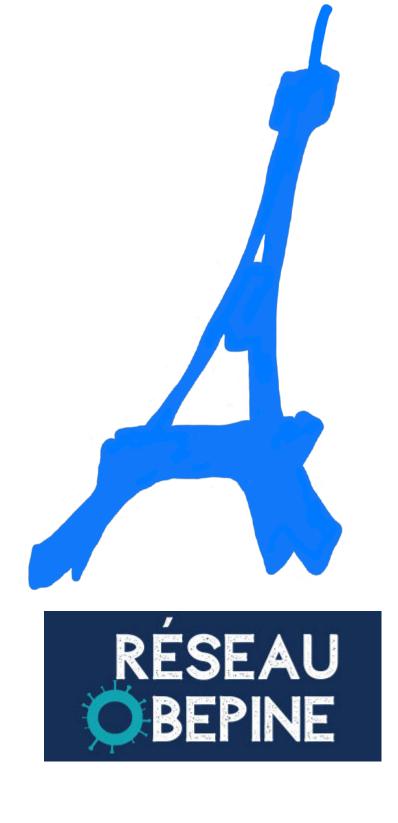
Webinar IDO











Construction, calibration and validation of a macro-epidemiological surveillance indicator at a national level from wastewater analysis.



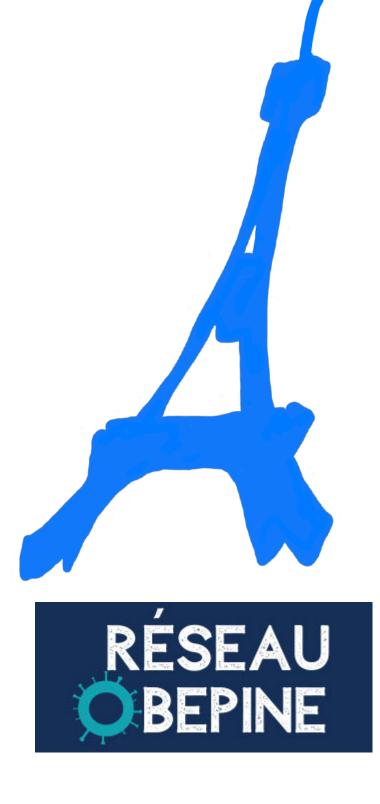












A nationwide indicator to smooth and normalize heterogeneous SARS-CoV2 RNA data in wastewater

Nicolas Cluzel^{1*}, Marie Courbariaux¹, Siyun Wang¹, Laurent Moulin², Sébastien Wurtzer², Isabelle Bertrand³, Karine Laurent¹, Patrick Monfort⁴, *Obépine* consortium^a, Christophe Gantzer³, Soizick Le Guyader⁵, Mickaël Boni⁶, Jean-Marie Mouchel^{7†}, Vincent Maréchal^{8†}, Grégory Nuel^{9,1†}, and Yvon Maday^{10*†}

















Is an interdisciplinary research group aiming at defining a macro epidemiological indicator from wastewater analysis

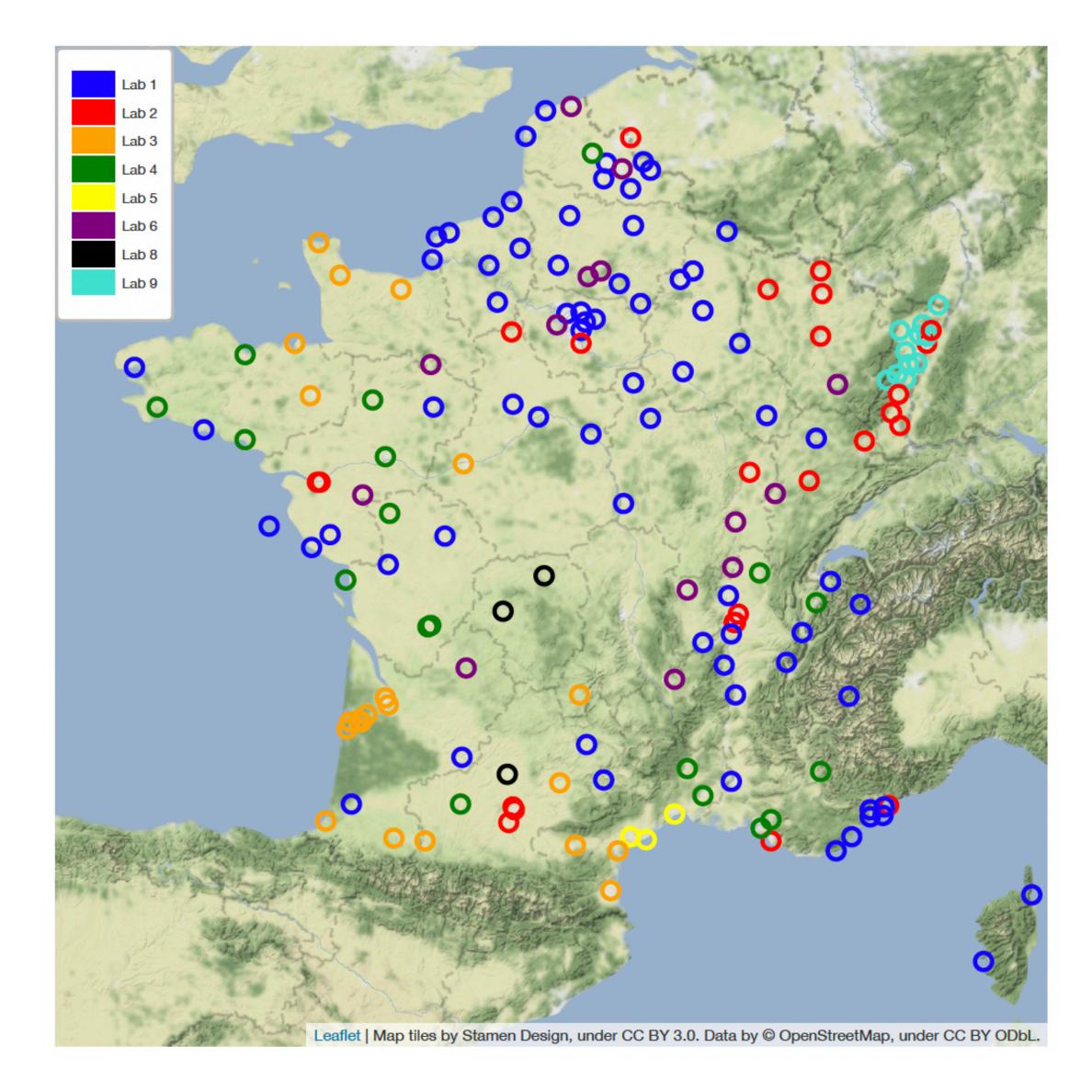


Is an interdisciplinary research group aiming at defining a macro epidemiological indicator from wastewater analysis

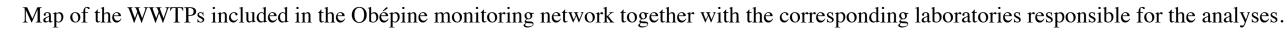
based on 6 main steps:

- qualification of the presence of the gene of the virus in wastewater at a low LOD
- quantification of the concentration of the trace of virus
- interpretation to get rid of the inherent errors
- transforming this in a reliable indicator in tendencies
- choice of the WW treatment plans
- homogenising it to propose a nation wide indicator

Currently following 200 WWTP

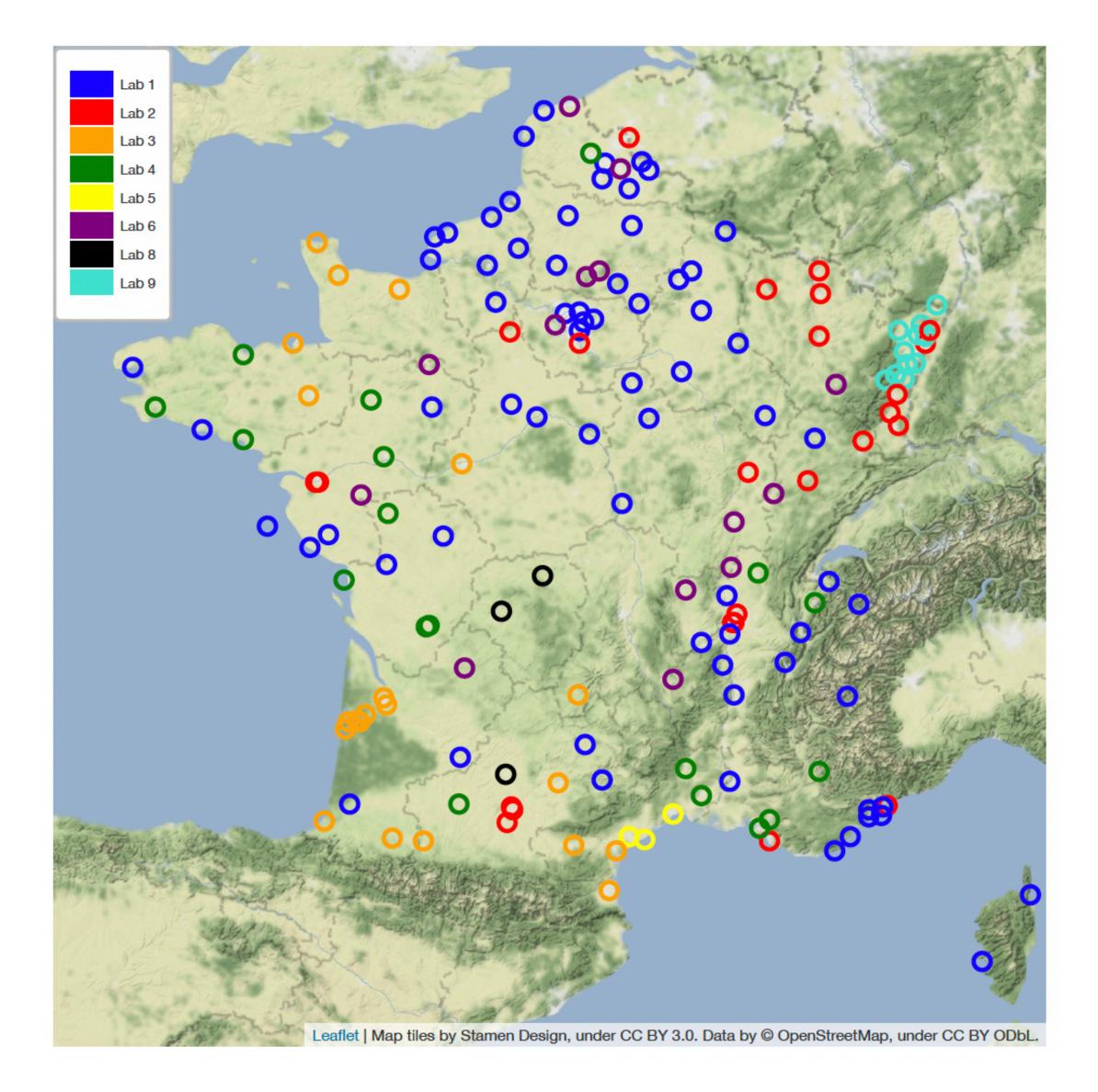


on 9 different laboratories





Currently following 200 WWTP
on integrated Sampling our 24 h



on 9 different laboratories

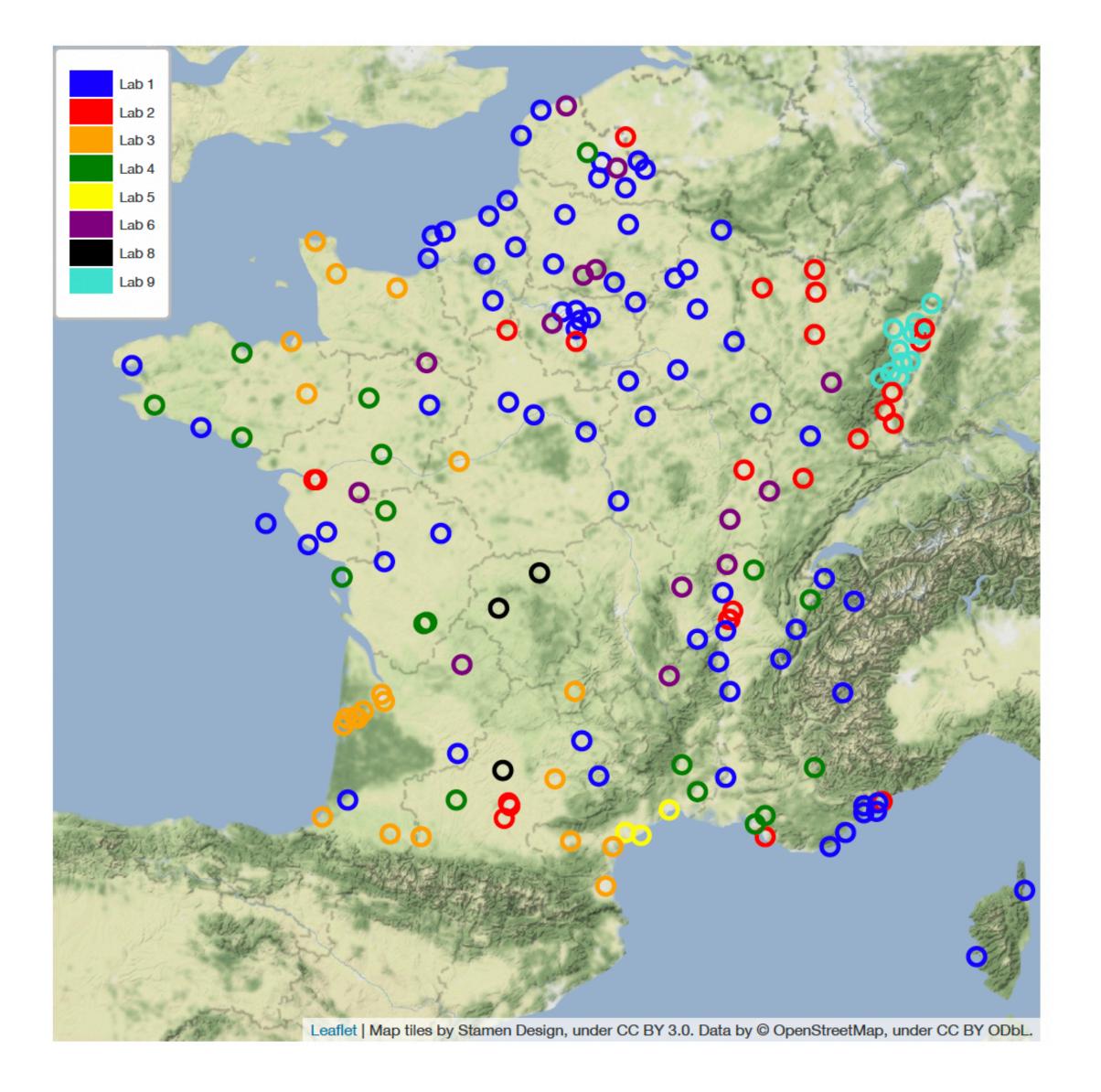
twice a week



Map of the WWTPs included in the Obépine monitoring network together with the corresponding laboratories responsible for the analyses.

Currently following 200 WWTP
on integrated sampling our 24 h

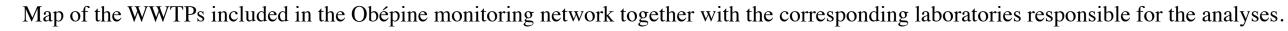
E and RdRp genes



on 9 different laboratories

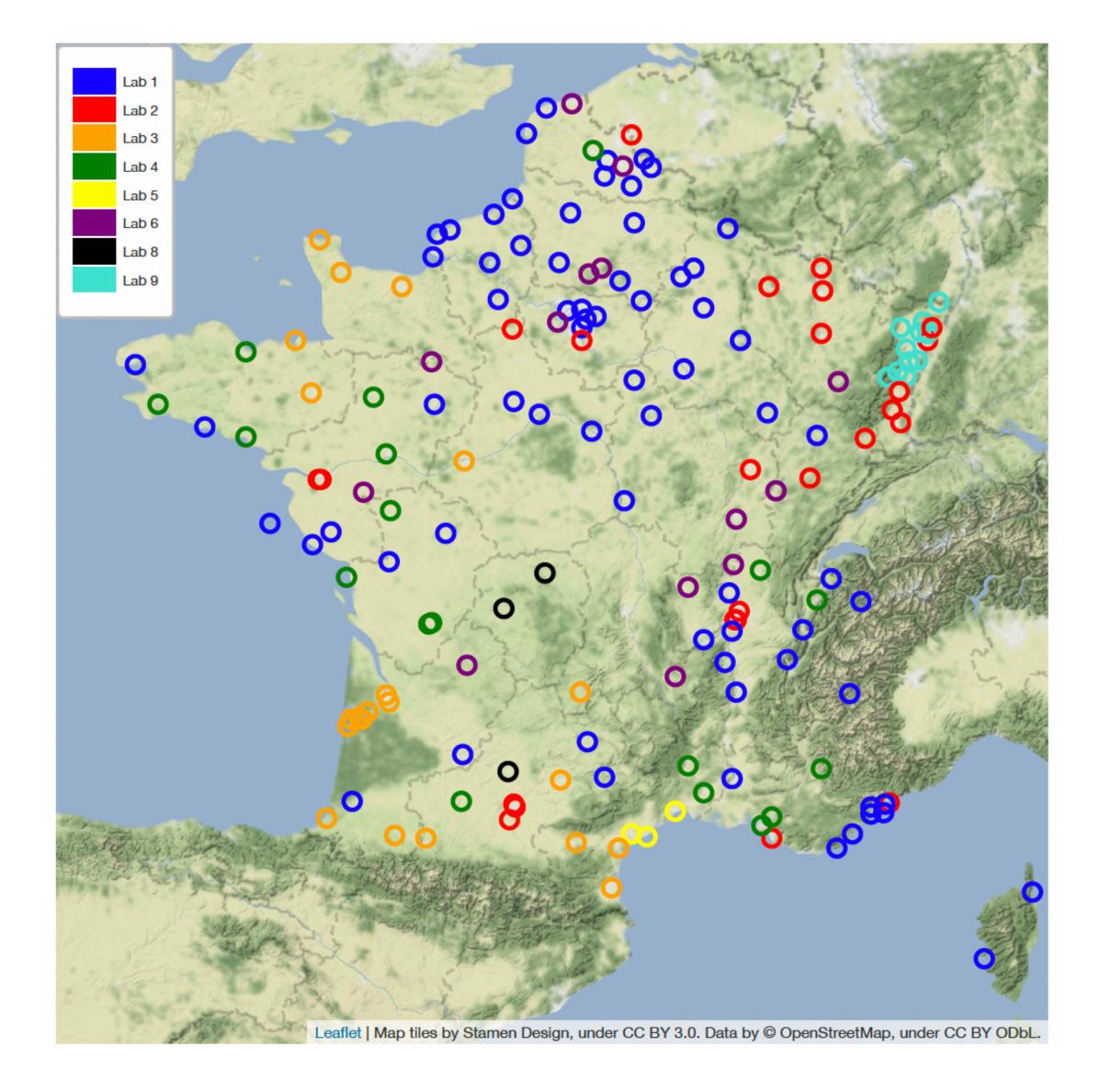
twice a week

reports sent to collectivities and MSS





Currently following 200 WWTP on integrated sampling our 24 h E and RdRp genes + characterization of variants



on 9 different laboratories

twice a week

reports sent to collectivities and MSS

Alpha, Delta, Omicron.



The concentrations are normalised

by taking into account additional data

$$C_t = (C_{0,t})_1 \times \left(\frac{V_t}{V_0}\right)_2 \times \left(\frac{Poll_0}{Poll_t}\right)_3$$

The concentrations are normalised

by taking into account additional data

$$C_t = (C_{0,t})_1 \times \left(\frac{V_t}{V_0}\right)_2 \times \left(\frac{Poll_0}{Poll_t}\right)_3$$
 different quality indexes

The concentrations are normalised

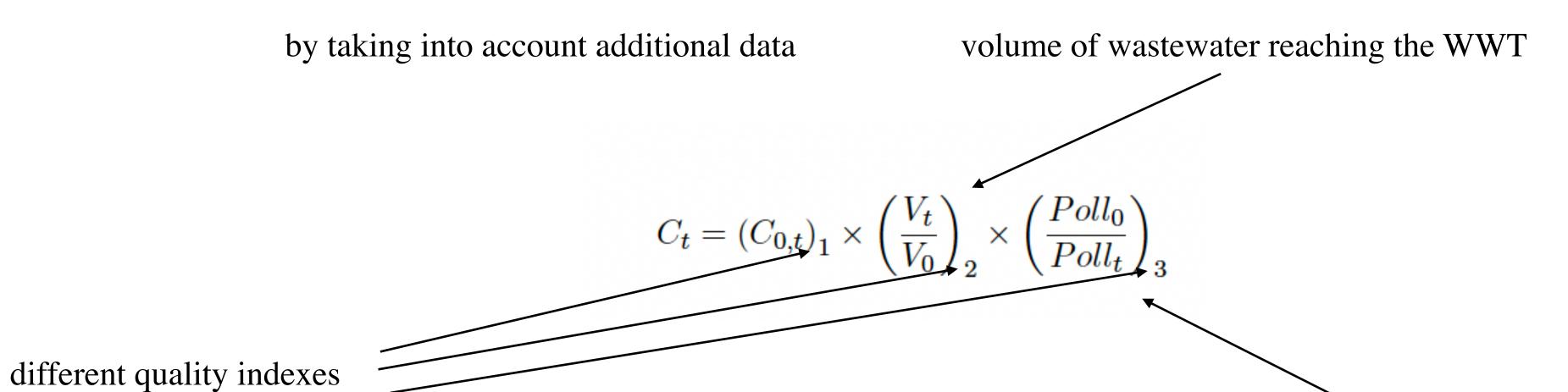
by taking into account additional data

volume of wastewater reaching the WWT

$$C_t = (C_{0,t})_1 \times \left(\frac{V_t}{V_0}\right)_2 \times \left(\frac{Poll_0}{Poll_t}\right)_3$$

different quality indexes

The concentrations are normalised



This factor is a function of ammonium, COD and conductivity concentrations, normalised by their mean dry weather value

We have developed a Kalman-smoothing algorithm adapted for the Obepine data.

This one allows to find an autoregressive process of which we observe only a noisy and censored version.

$$X_i = \eta X_{i-1} + \delta + \sigma \mathcal{N}(0, 1)$$

$$Y'_i = X_i + \varepsilon \mathcal{N}(0, 1)$$

$$Y_i = \max(Y'_i, c)$$

In our case, we are looking for an estimate of the "real" concentrations in the wastewater, X, of which we only observe noisy and censored measurements, Y.

The proposed method also allows to provide the a posteriori law of the underlying process, and thus intervals in which the "true" concentration is found in 99%, 95%, 50%, . . . of the cases.

We have developed a Kalman-smoothing algorithm adapted for the Obepine data.

This one allows to find an autoregressive process of which we observe only a noisy and censored version.

$$X_i = \eta X_{i-1} + \delta + \sigma \mathcal{N}(0,1)$$

$$Y_i' = X_i + \varepsilon \mathcal{N}(0,1) \qquad Y_t = \log(C_t)$$
). Y_t is generally only partially observed
$$Y_i = \max(Y_i',c)$$

In our case, we are looking for an estimate of the "real" concentrations in the wastewater, X, of which we only observe noisy and censored measurements, Y.

The proposed method also allows to provide the a posteriori law of the underlying process, and thus intervals in which the "true" concentration is found in 99%, 95%, 50%, . . . of the cases.

We have developed a Kalman-smoothing algorithm adapted for the Obepine data.

This one allows to find an autoregressive process of which we observe only a noisy and censored version.

true process, that we want to recover

$$X_i = \eta X_{i-1} + \delta + \sigma \mathcal{N}(0, 1)$$

$$Y'_i = X_i + \varepsilon \mathcal{N}(0, 1)$$

$$Y_i = \max(Y'_i, c)$$

In our case, we are looking for an estimate of the "real" concentrations in the wastewater, X, of which we only observe noisy and censored measurements, Y.

The proposed method also allows to provide the a posteriori law of the underlying process, and thus intervals in which the "true" concentration is found in 99%, 95%, 50%, . . . of the cases.

We have developed a Kalman-smoothing algorithm adapted for the Obepine data.

This one allows to find an autoregressive process of which we observe only a noisy and censored version.

true process, that we want to recover

$$X_i = \eta X_{i-1} + \delta + \sigma \mathcal{N}(0, 1)$$

$$Y'_i = X_i + \varepsilon \mathcal{N}(0, 1)$$

$$Y_i = \max(Y'_i, c)$$

In our case, we are looking for an estimate of the "real" concentrations in the wastewater, X, of which we only observe noisy and censored measurements, Y.

The proposed method also allows to provide the a posteriori law of the underlying process, and thus intervals in which the "true" concentration is found in 99%, 95%, 50%, . . . of the cases.

takes into account the noise (errors) in the measures

We have developed a Kalman-smoothing algorithm adapted for the Obepine data.

This one allows to find an autoregressive process of which we observe only a noisy and censored version.

true process, that we want to recover

$$Y_i' = \eta X_{i-1} + \delta + \sigma \mathcal{N}(0, 1)$$

$$Y_i' = X_i + \varepsilon \mathcal{N}(0, 1)$$

$$Y_i = \max(Y_i', c)$$

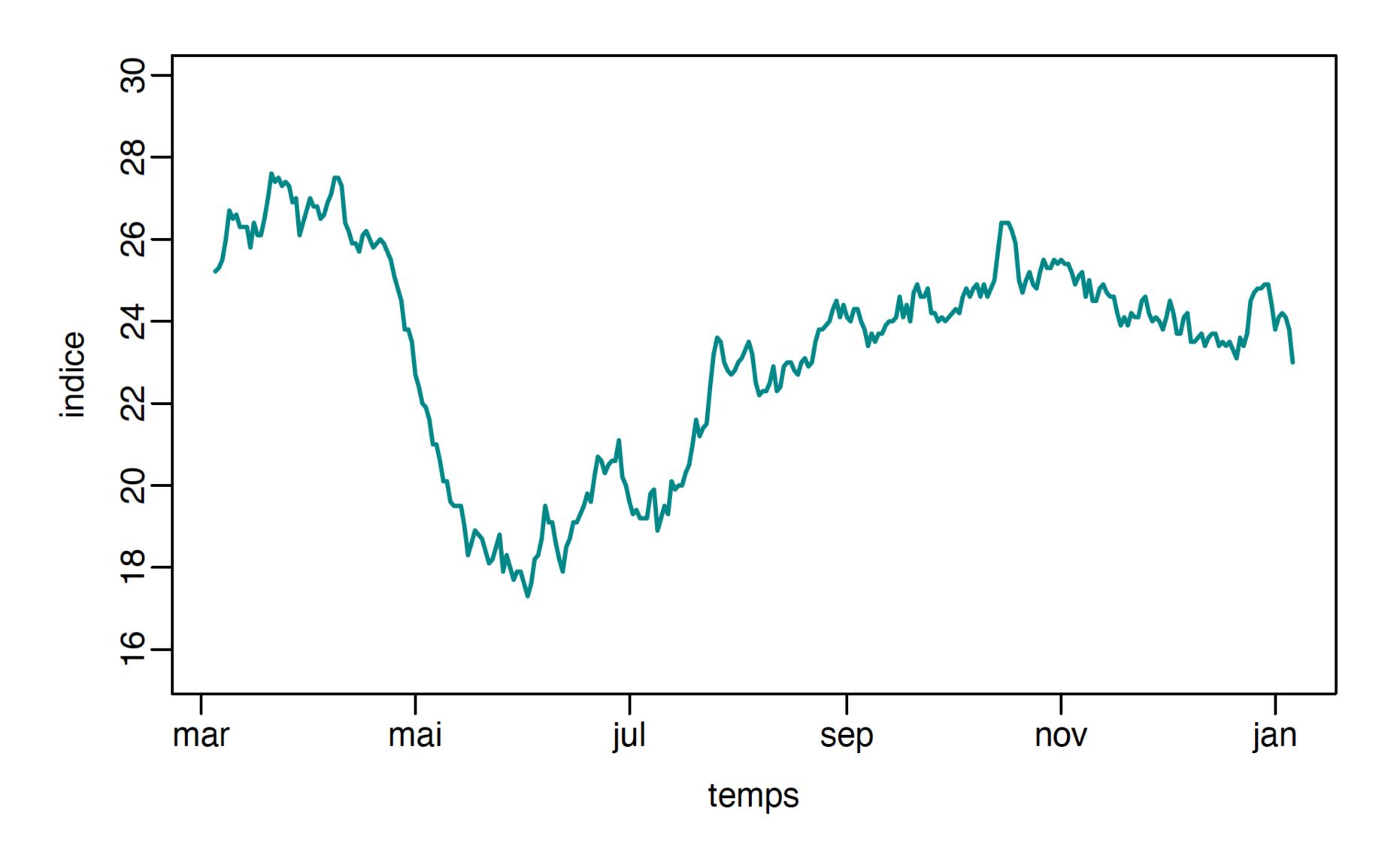
In our case, we are looking for an estimate of the "real" concentrations in the wastewater, X, of which we only observe noisy and censored measurements, Y.

The proposed method also allows to provide the a posteriori law of the underlying process, and thus intervals in which the "true" concentration is found in 99%, 95%, 50%, . . . of the cases.

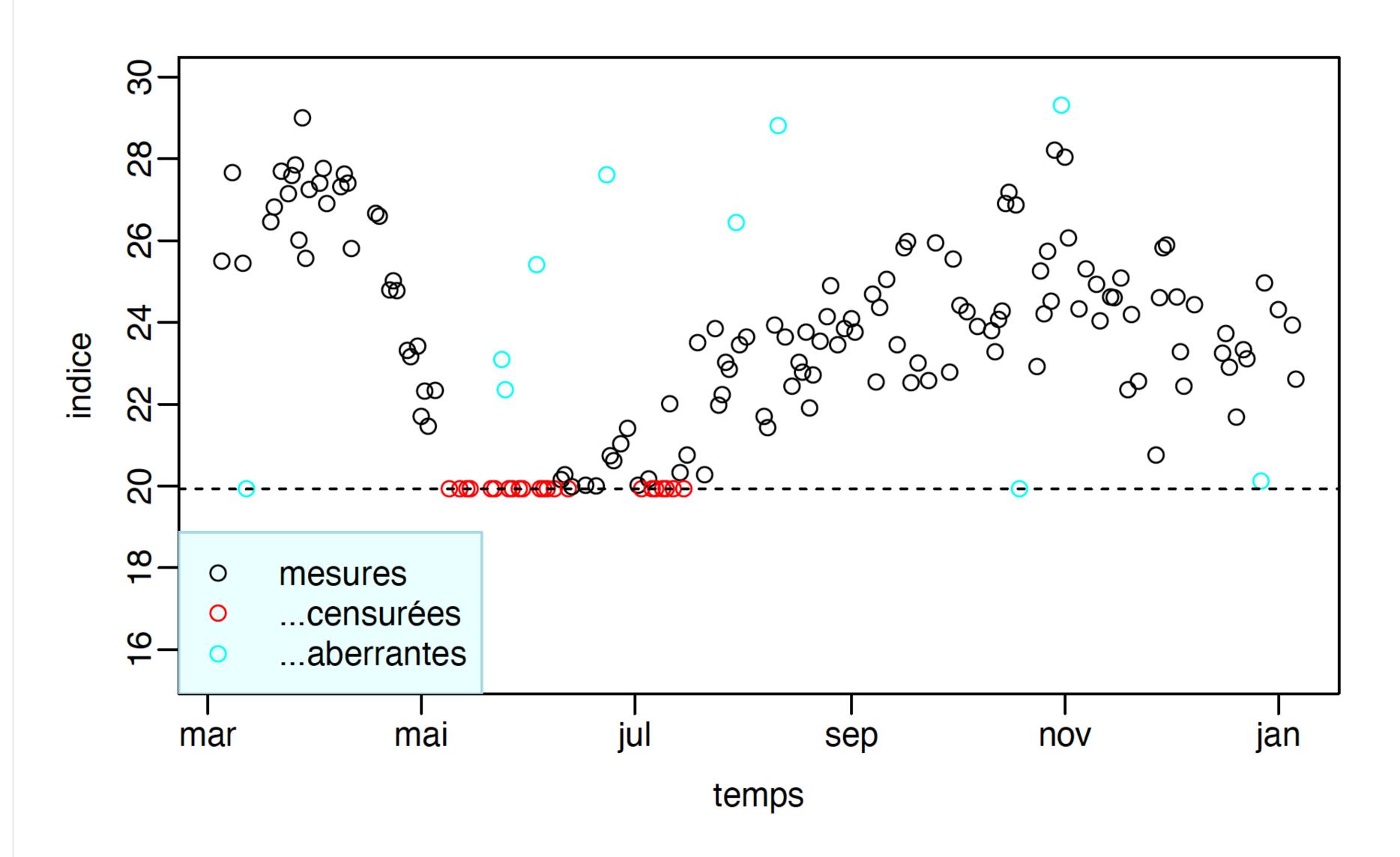
takes into account the noise (errors) in the measures

takes into account the censoring

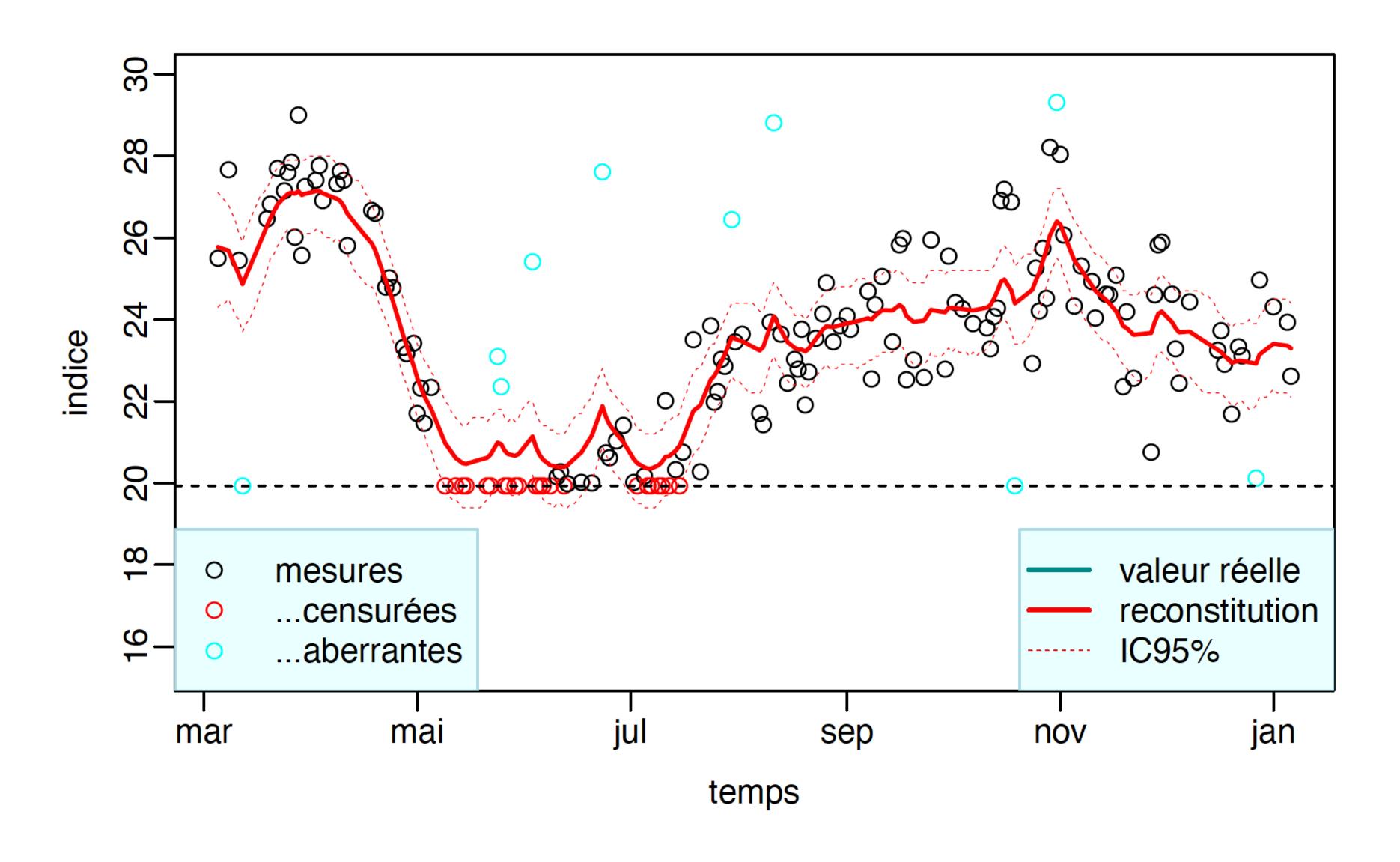
True measures

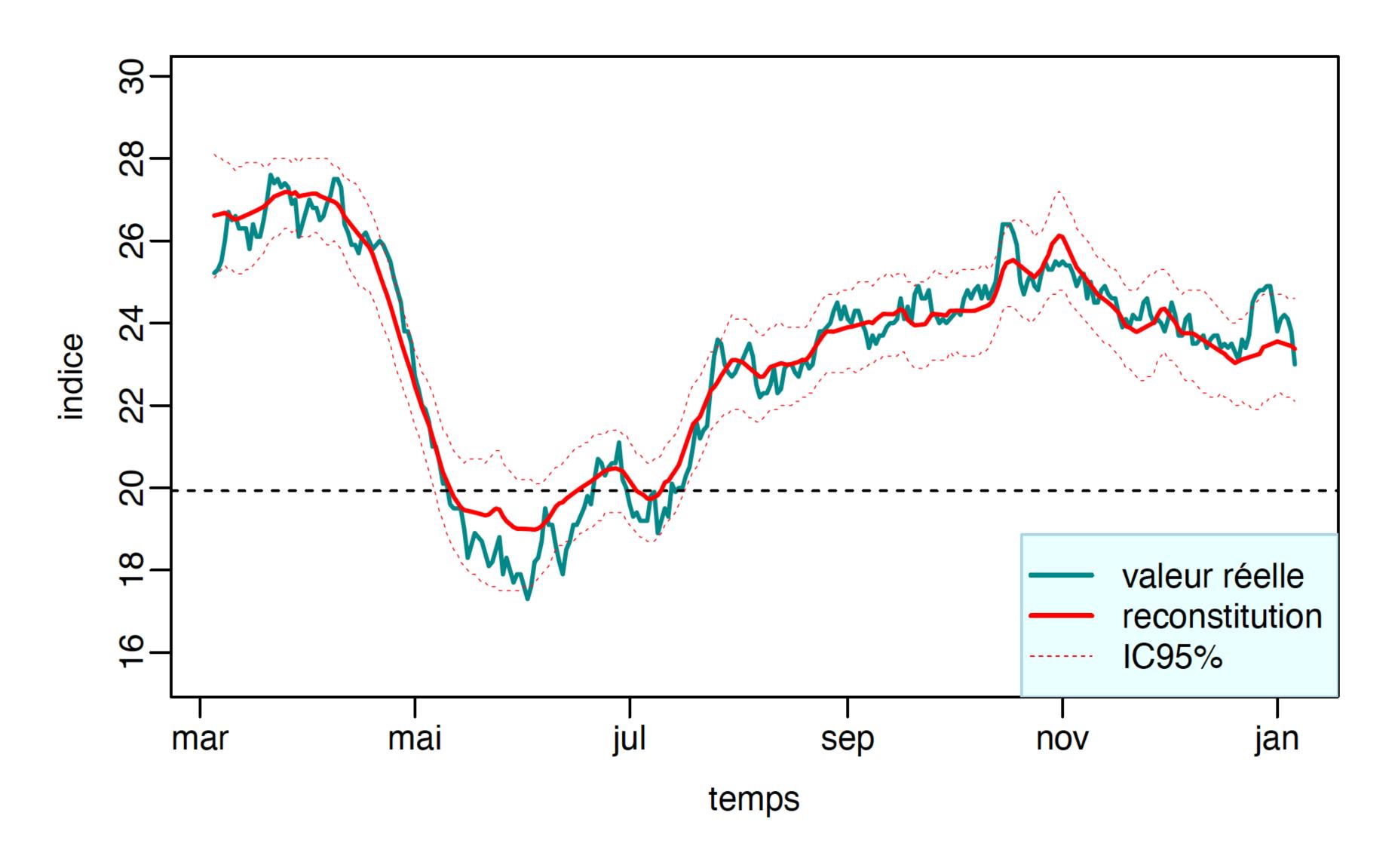


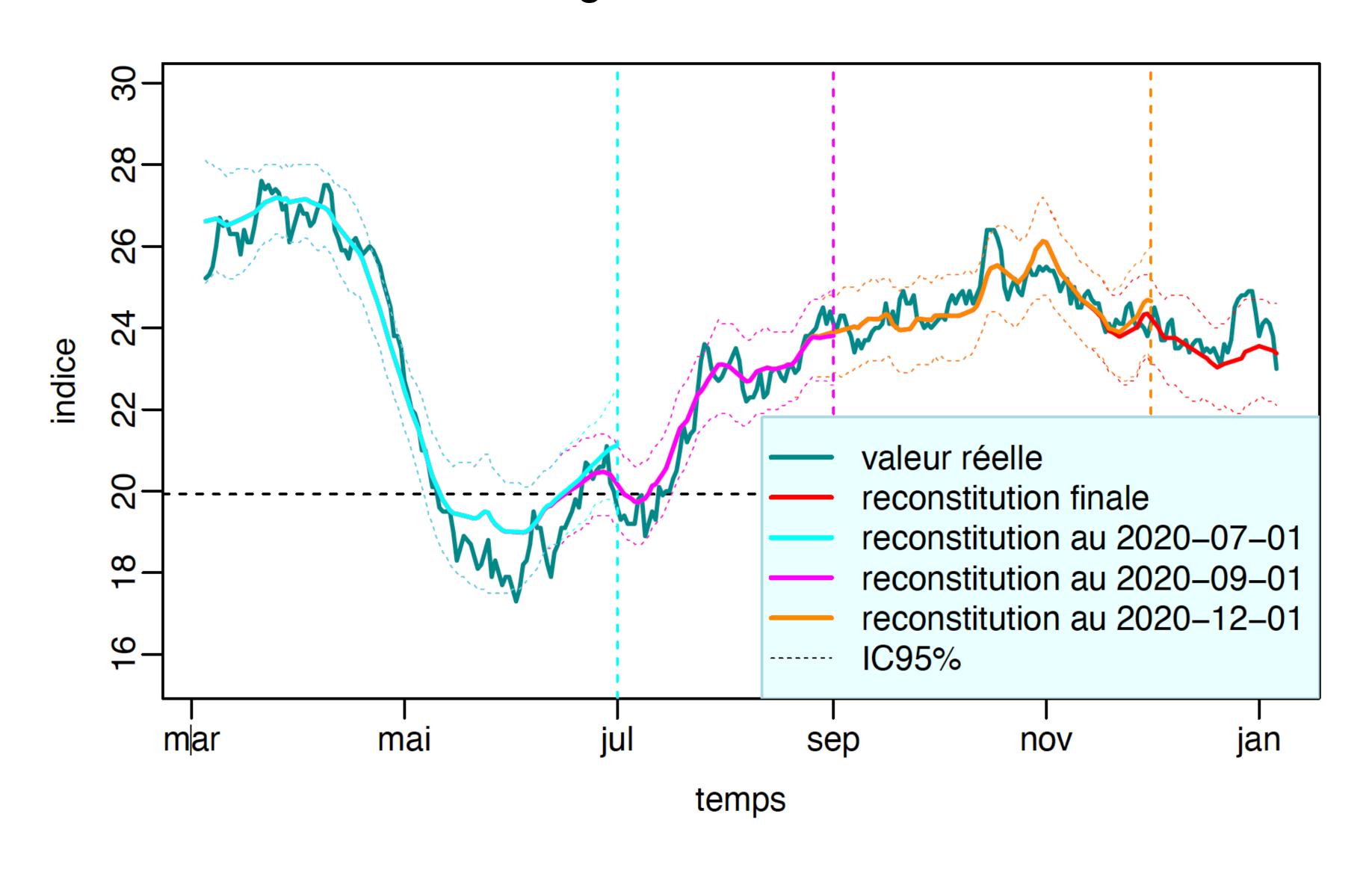
raw data



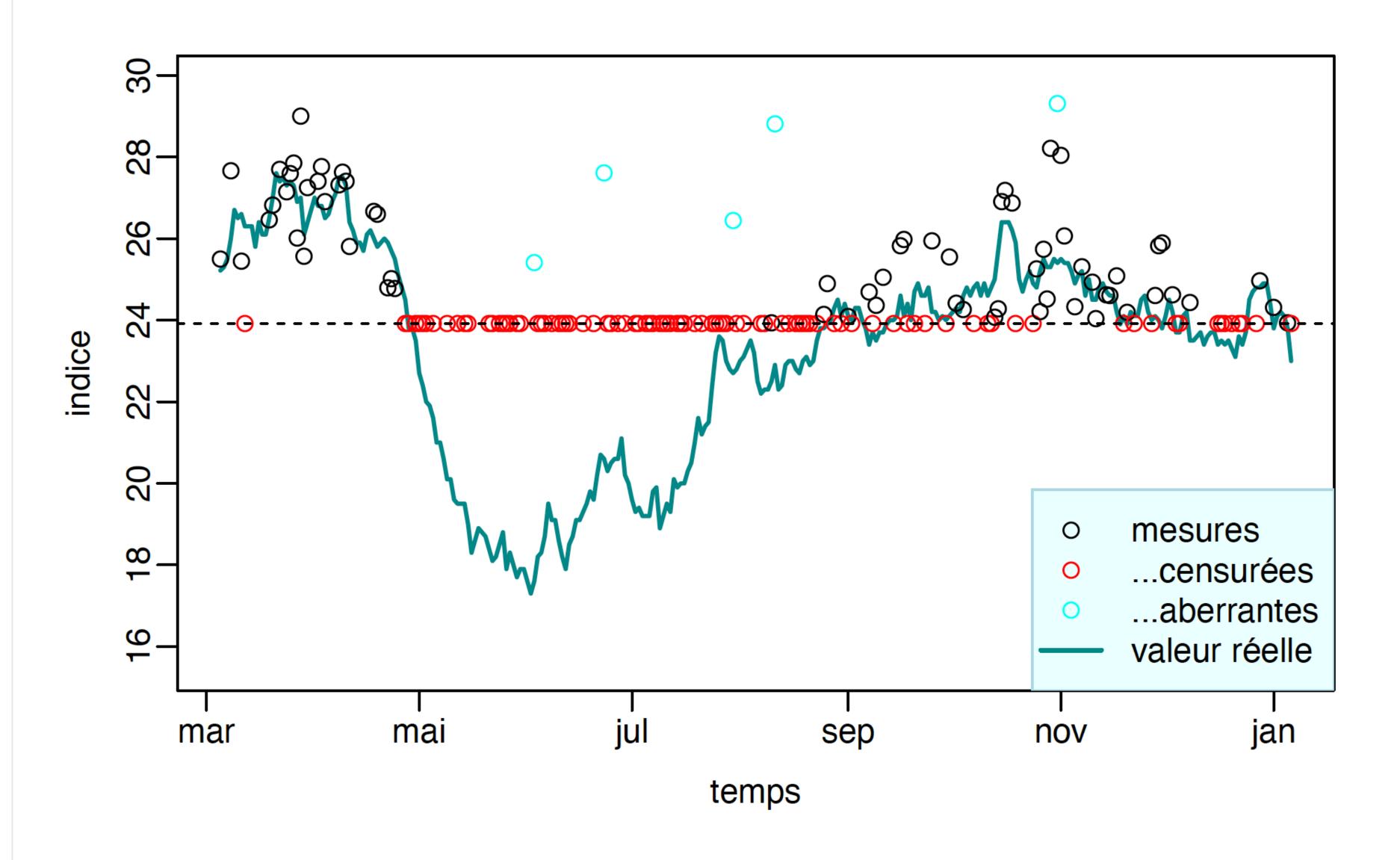
Raw treatment

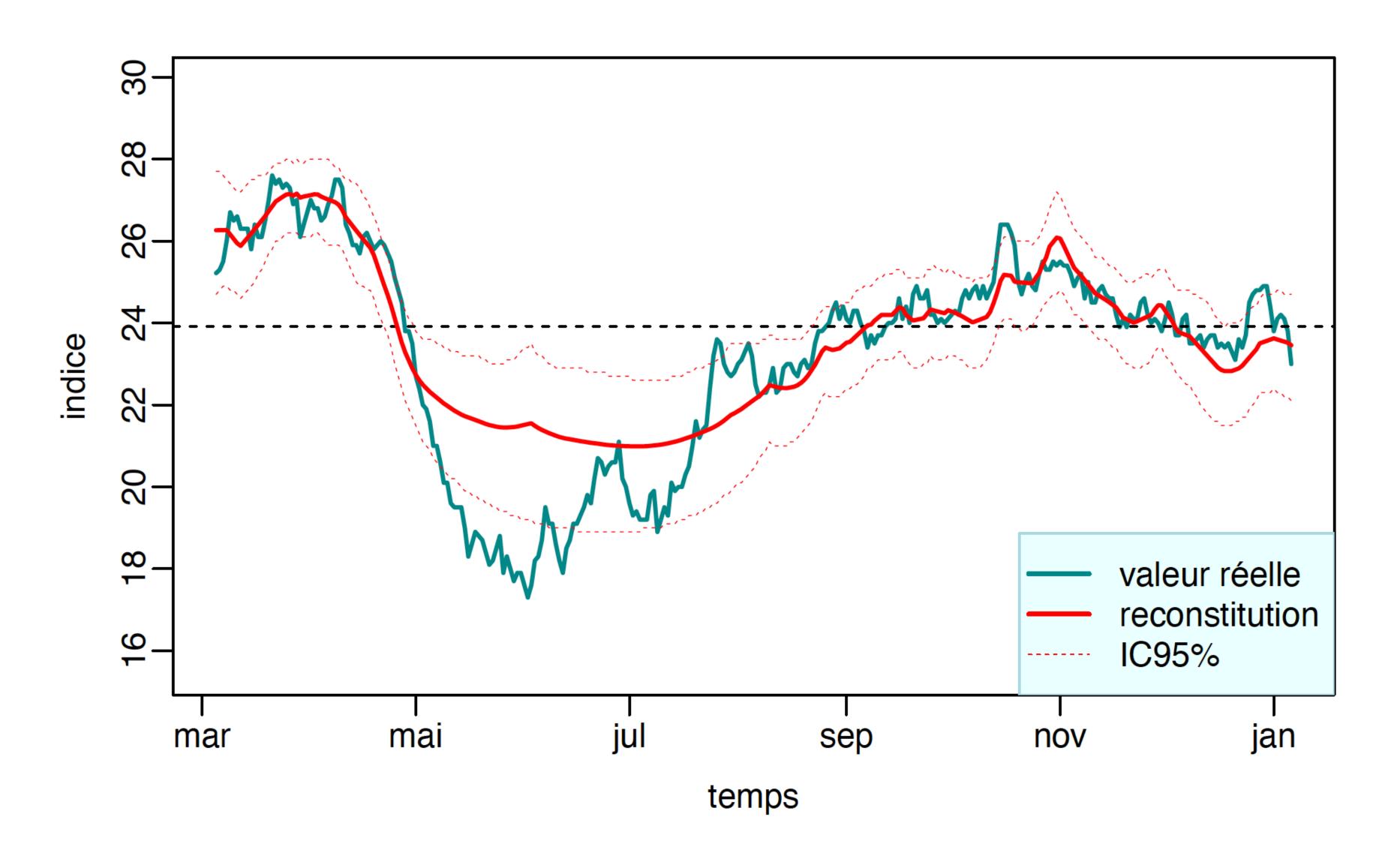


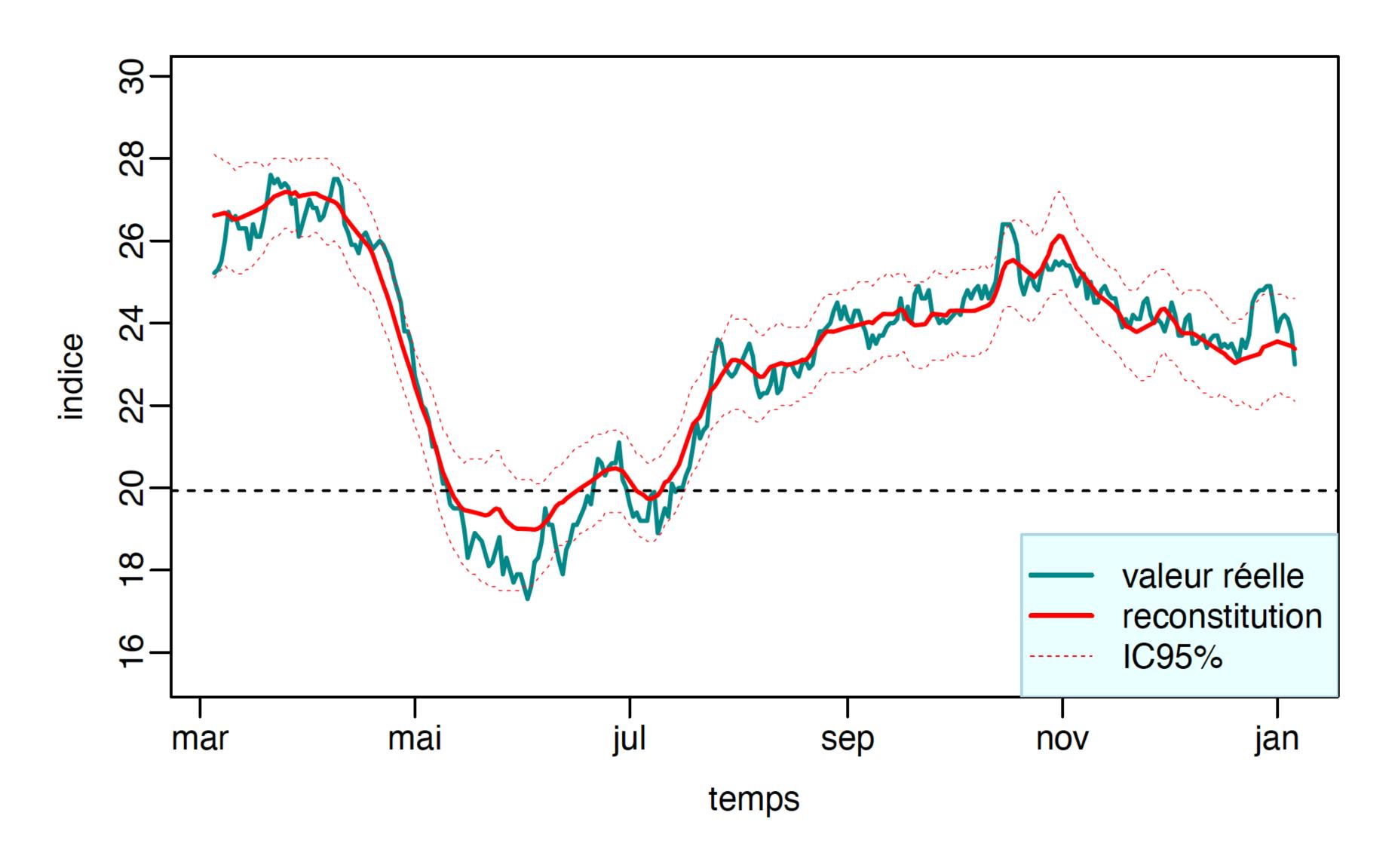


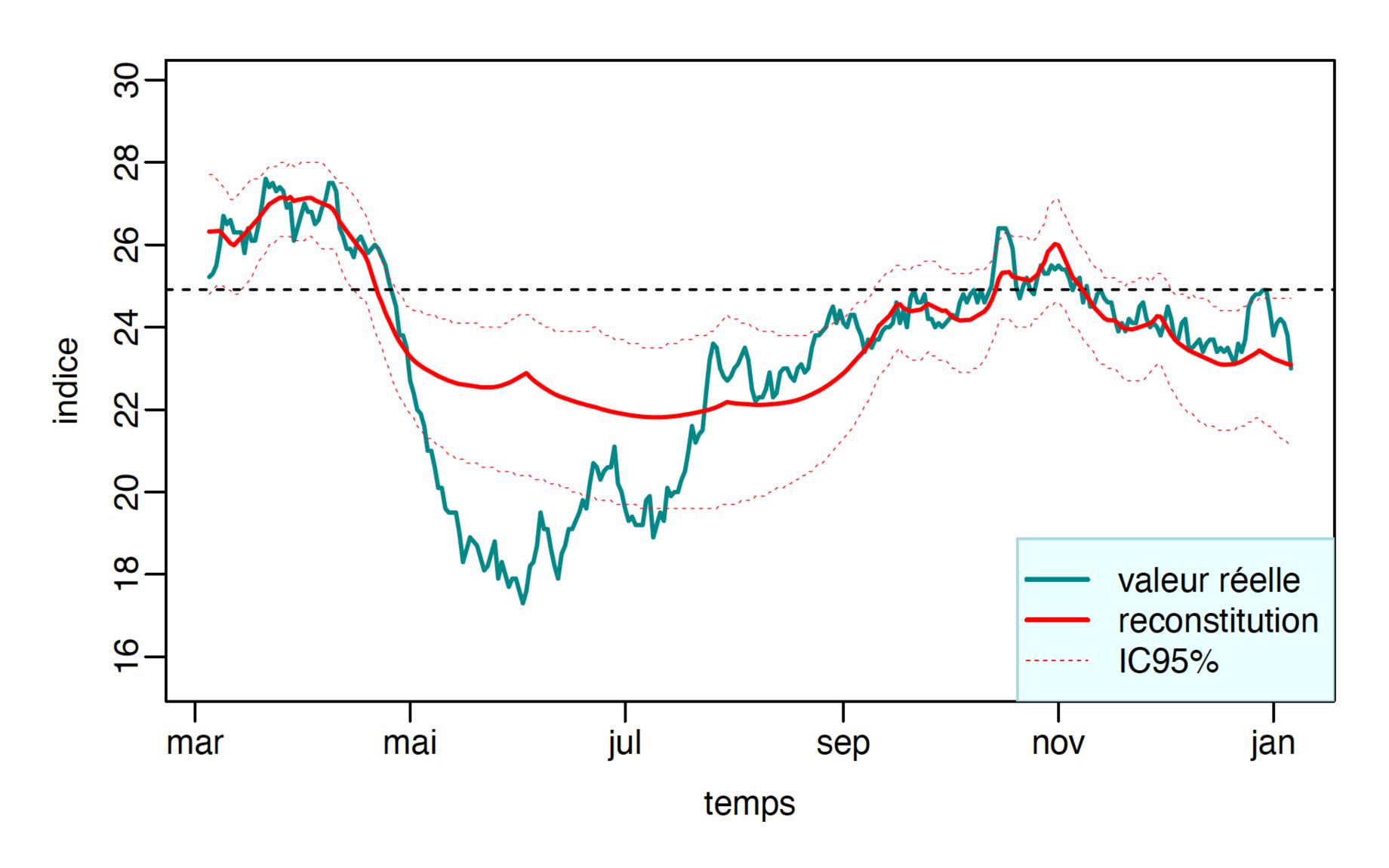


raw data









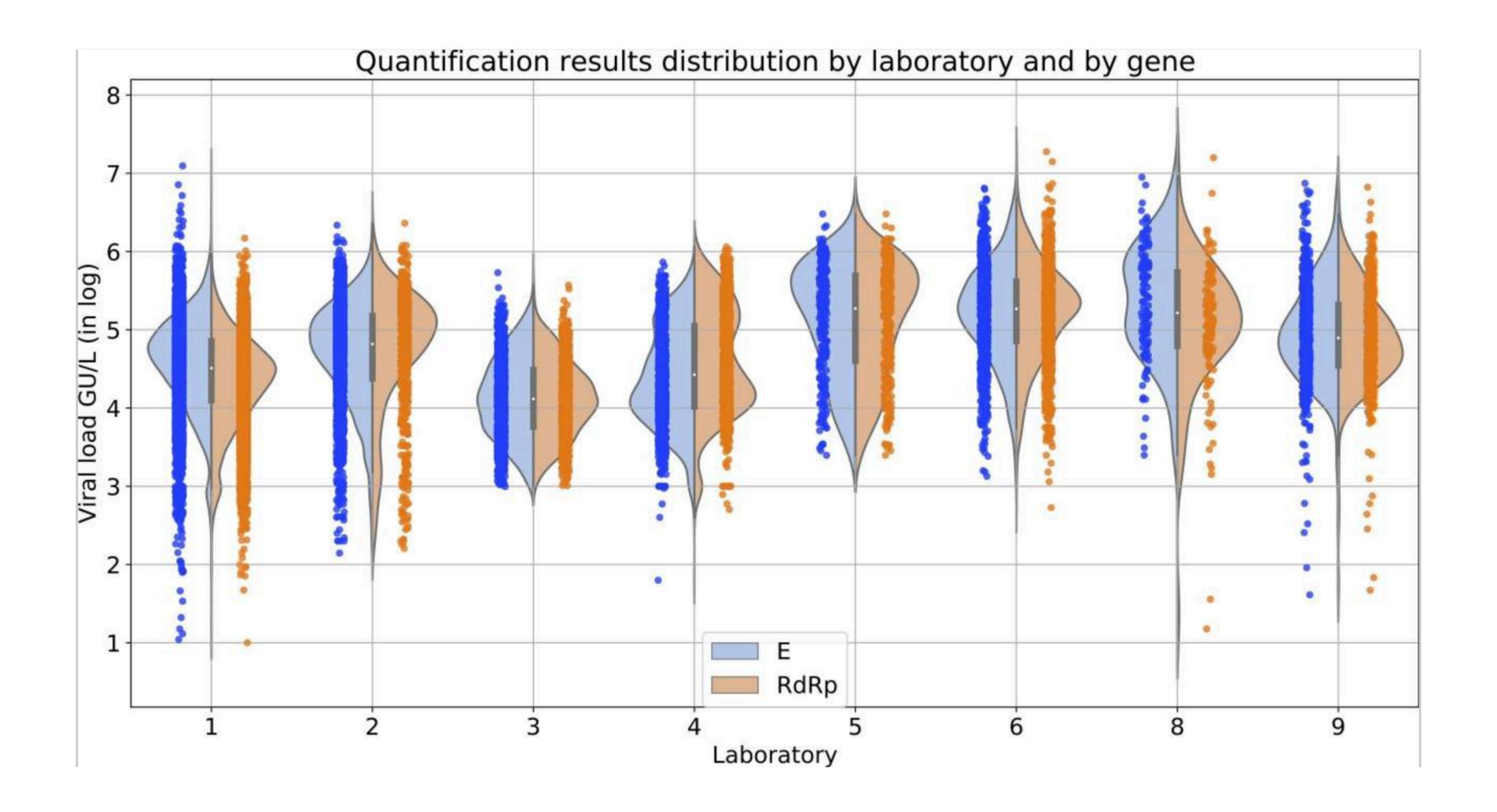
Normalization among the different laboratories

Organization of inter lab qualification and quantification

Normalization among the different laboratories

Organization of inter lab qualification and quantification

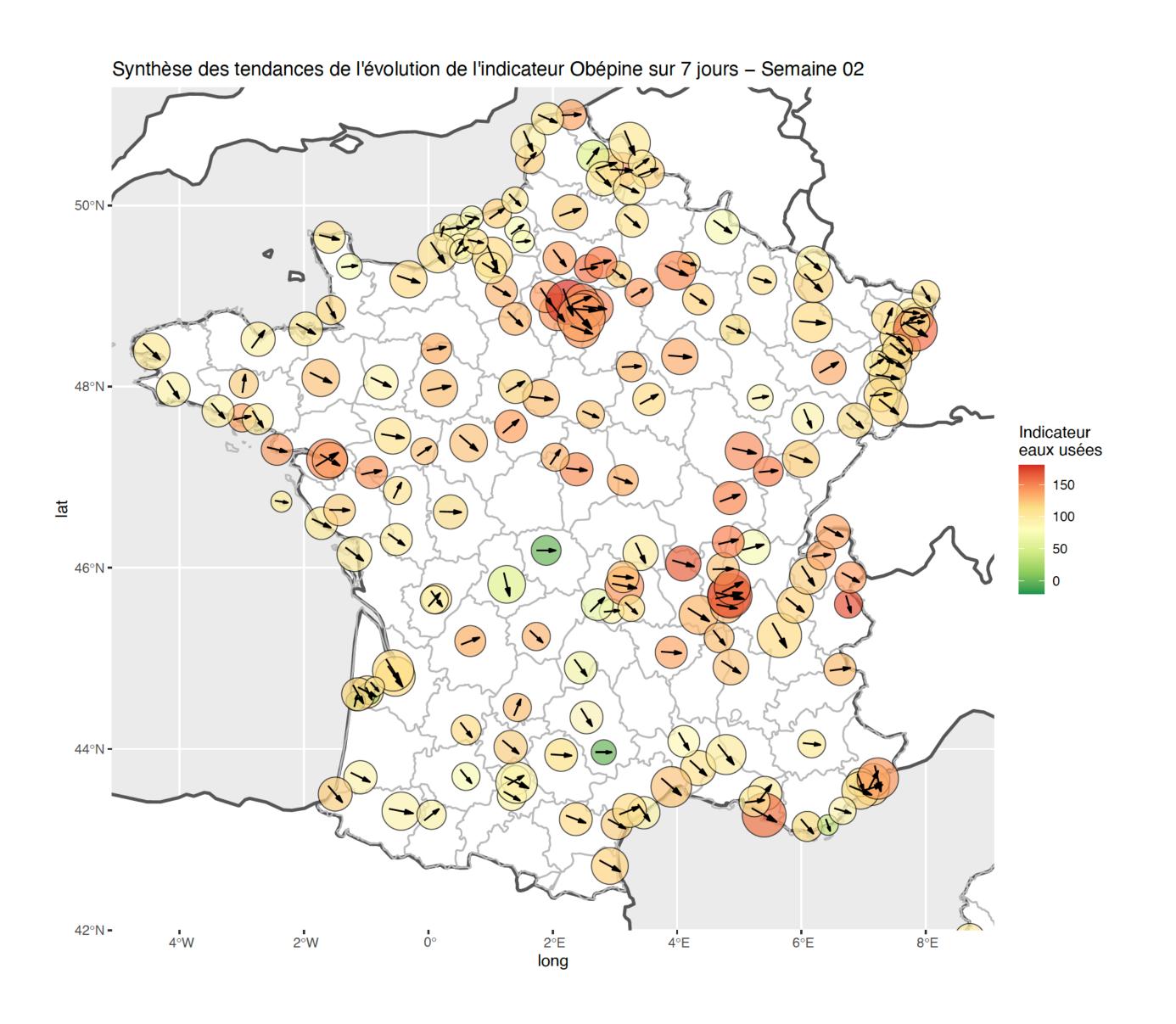
about 25 have participated: 10 were selected

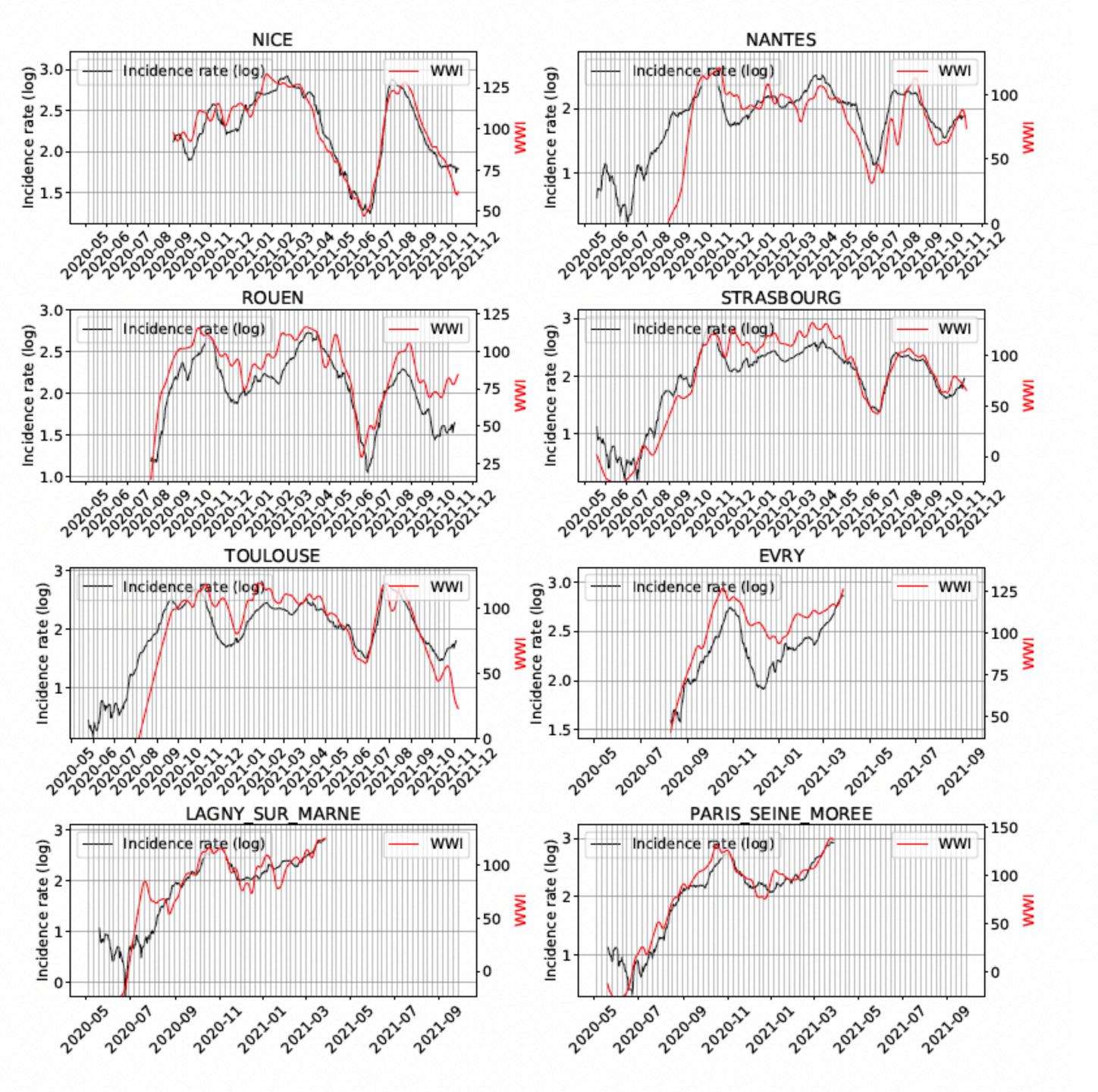


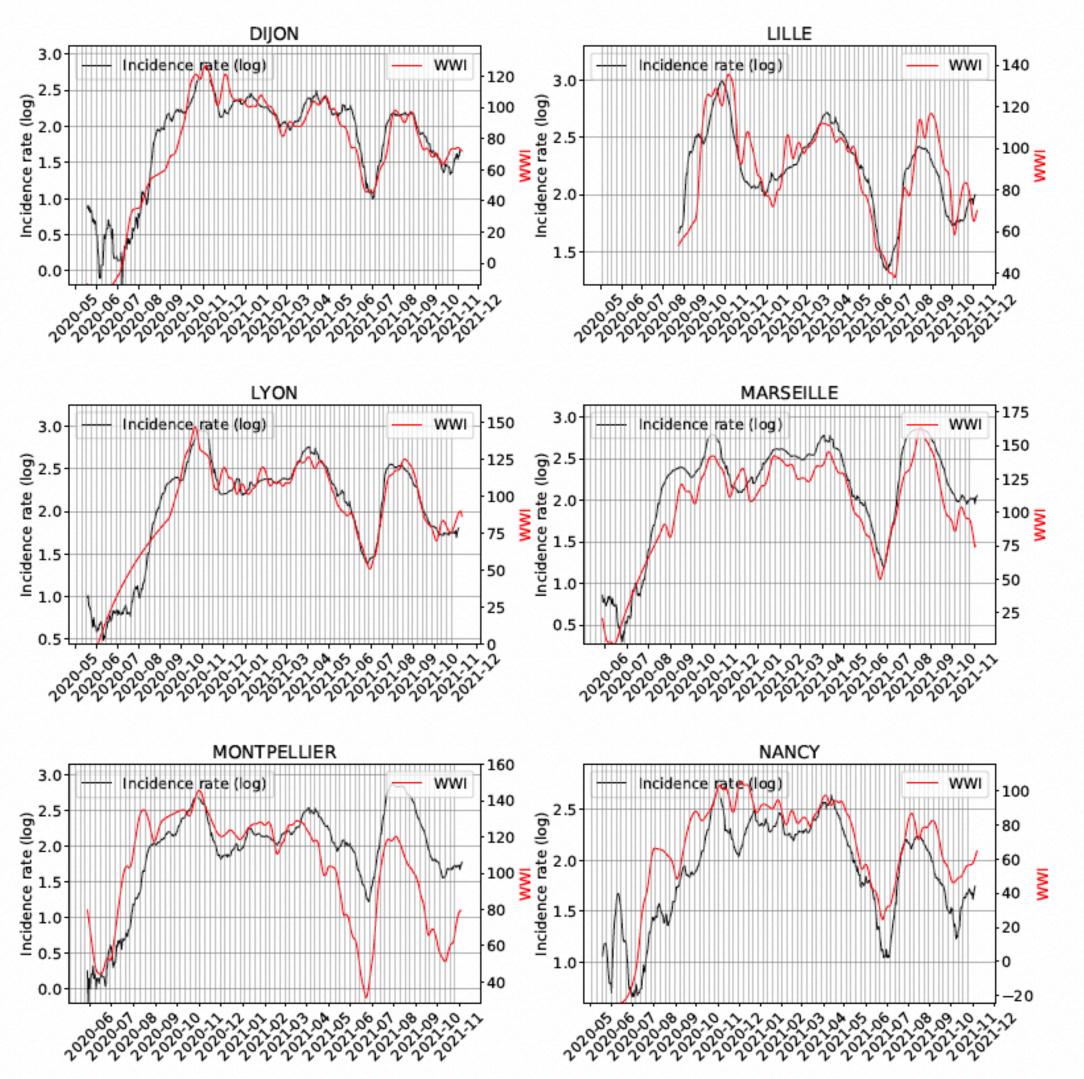
Normalization among the different laboratories

Organization of inter lab qualification and quantification

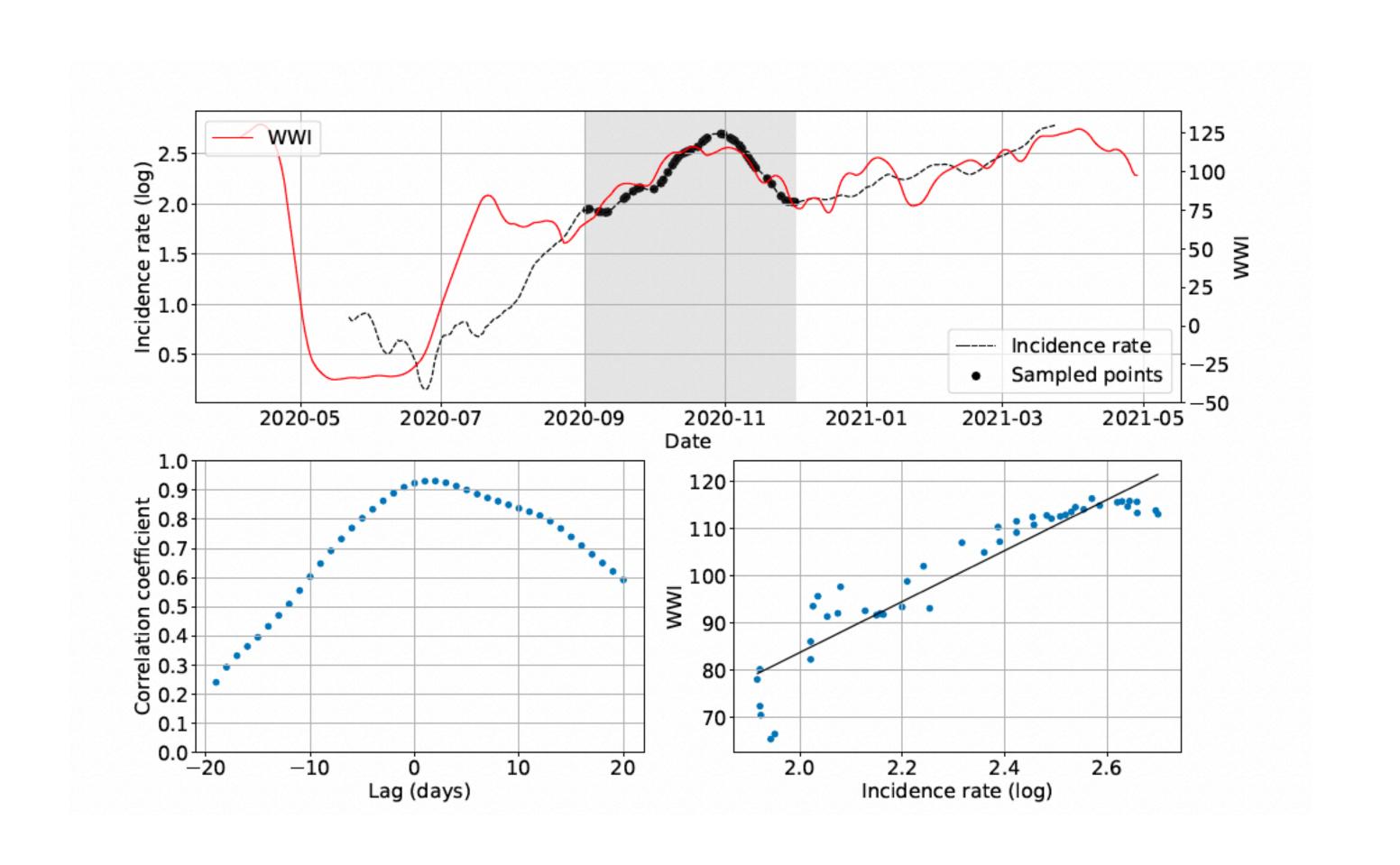
allows to present a comparable treatment over the network



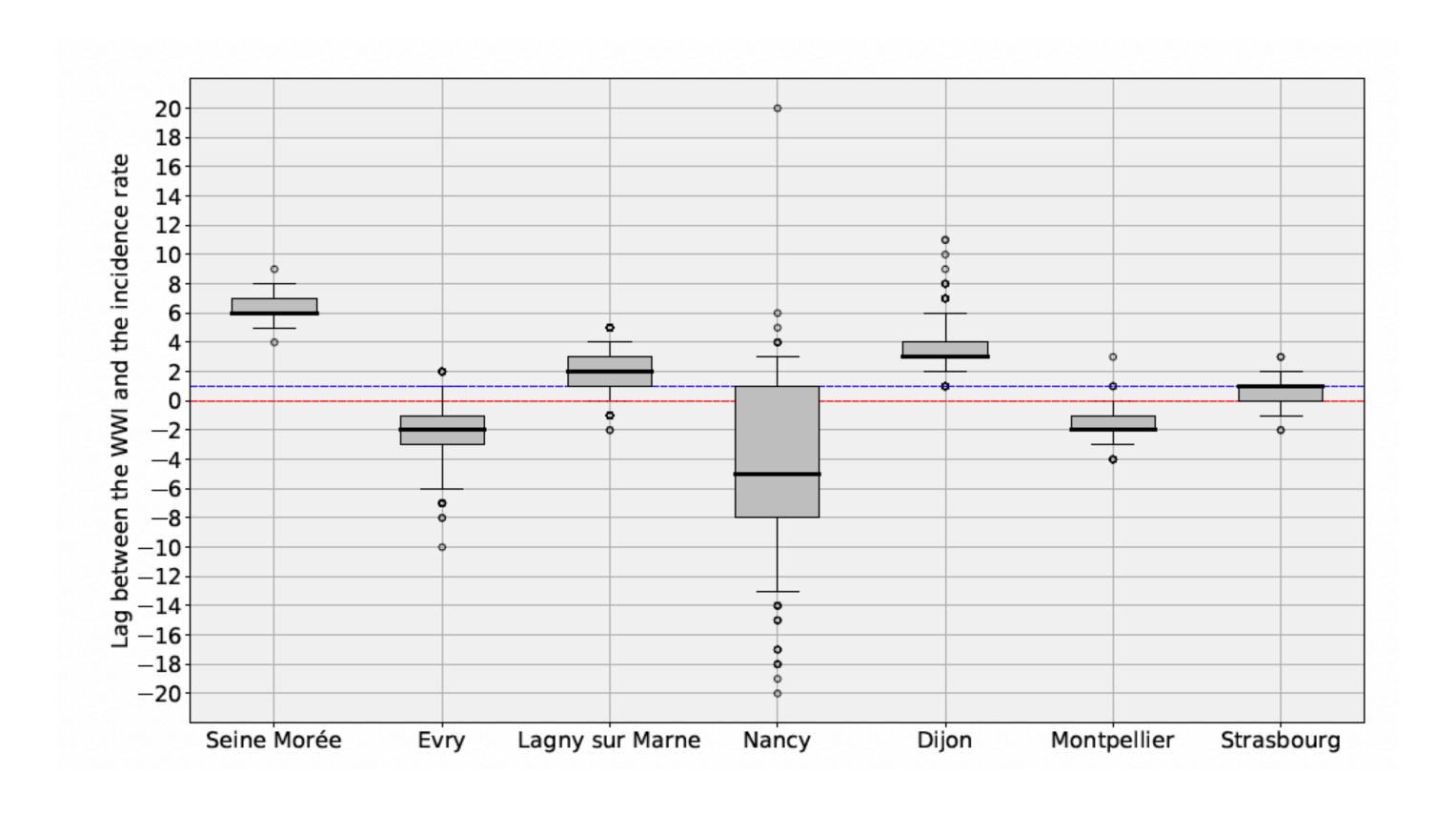




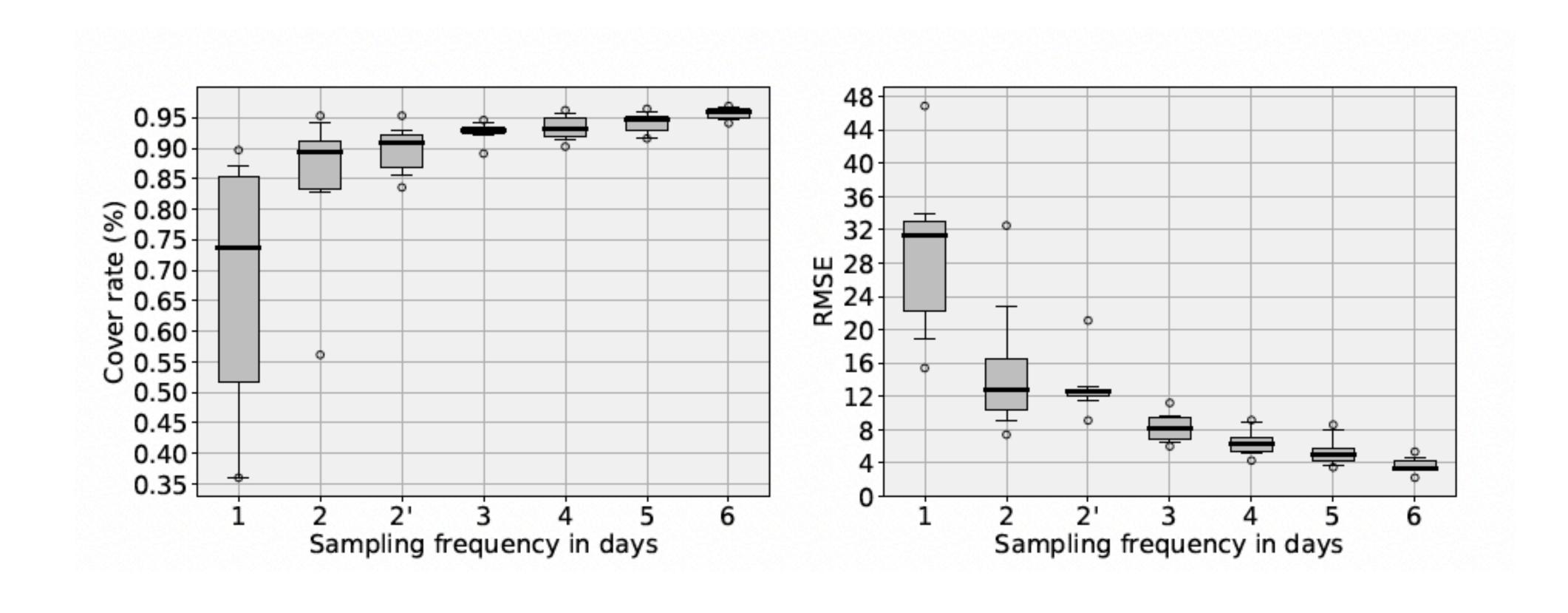
Correlation and lag between the WWI and the incidence rate



Correlation and lag between the WWI and the incidence rate



Impact of the sampling frequency



Different types of filtering processes

Different types of filtering processes

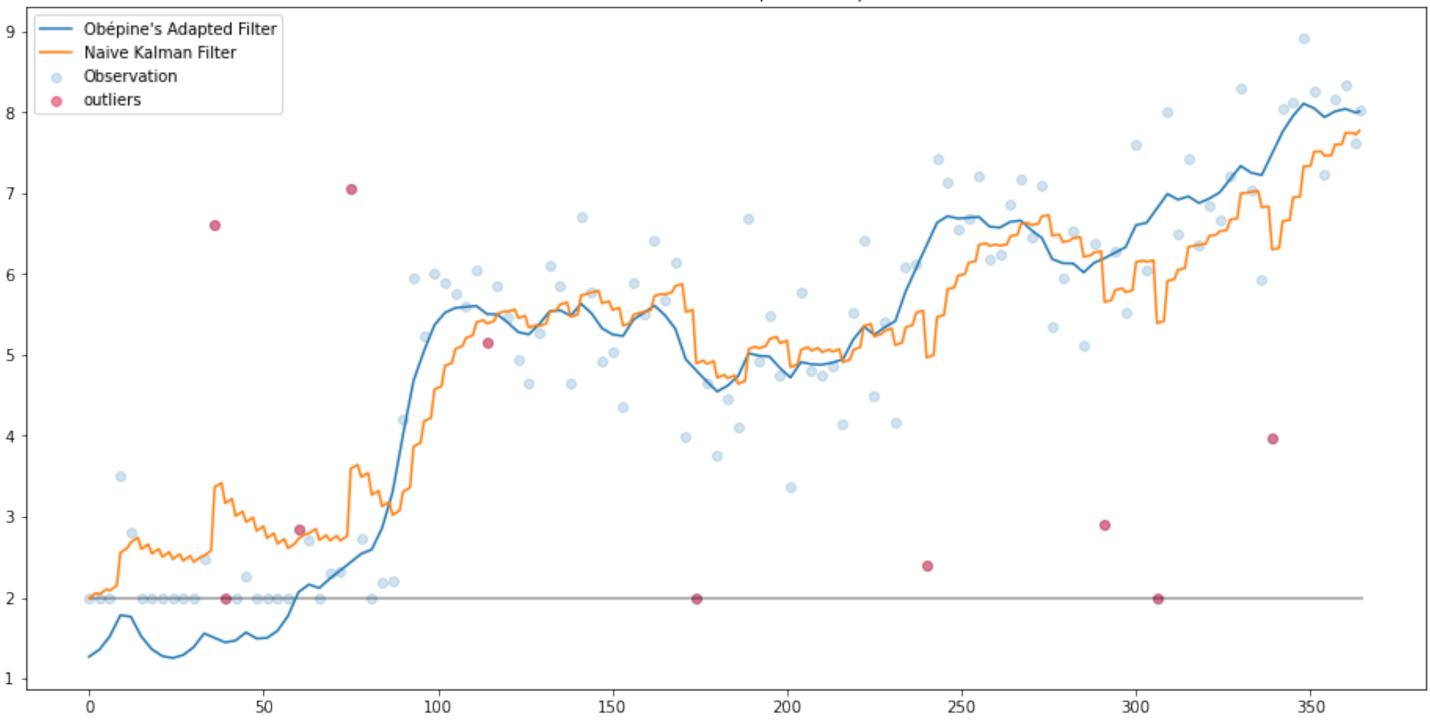
Data filtering methods for SARS-CoV-2 wastewater surveillance

Rezgar Arabzadeh ¹, Daniel Martin Grünbacher ¹, Heribert Insam ², Norbert Kreuzinger ³, Rudolf Markt ², Wolfgang Rauch ¹

Different types of filtering processes

Method	Reference	Sample
TUK	Mallows (1979)	Fiskeaux & Ling (1982)
KAF	Tusell (2011)	Pan et al. (2016)
FFT	Cochran <i>et al.</i> (1967)	Yang et al. (2004)
SPL ^{a,e}	Reinsch (1967)	Eubank (1988)
KER ^{a,e}	Härdle & Vieu (1992)	Speckman (1988))
SMA ^a	Hyndman (2011)	He et al. (2020)
RRM ^a	Friedman & Stuetzle (1982)	Polasek (1984)
SUP ^{a,e}	Friedman (1984)	Friedman & Silverman (1989)
$POL^{a,e}$	Atkeson <i>et al.</i> (1997)	Rajagopalan & Lall (1998)
SGF^b	Press & Teukolsky (1990)	Bromba & Ziegler (1981)
ARI ^{b,e}	Akaike (1969)	Lohani et al. (2012)
$ADP^{c,e}$	Barak (1995)	Jakubowska & Kubiak (2004)
$GAM^{d,e}$	Hastie (2017)	Murphy et al. (2019)





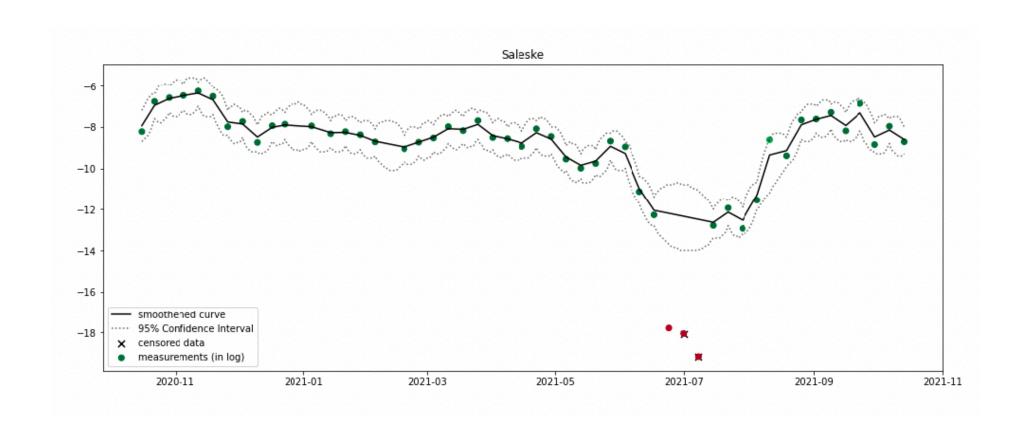


NACIONALNI INŠTITUT ZA BIOLOGIJO NATIONAL INSTITUTE OF BIOLOGY





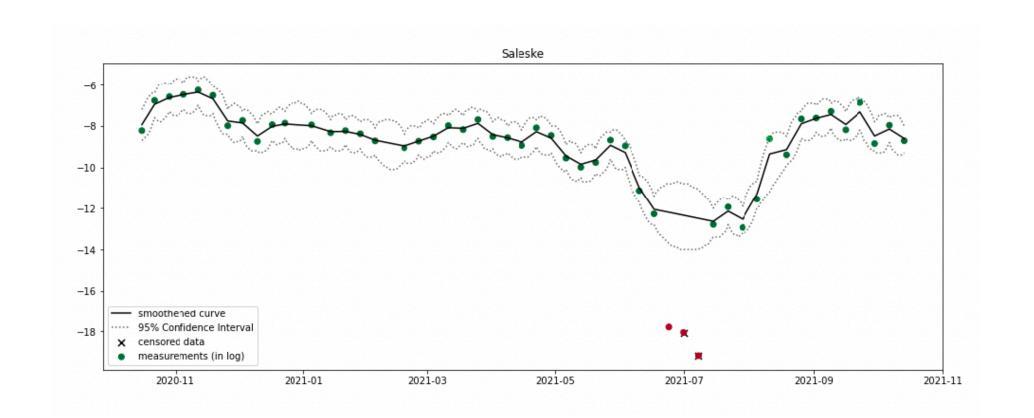


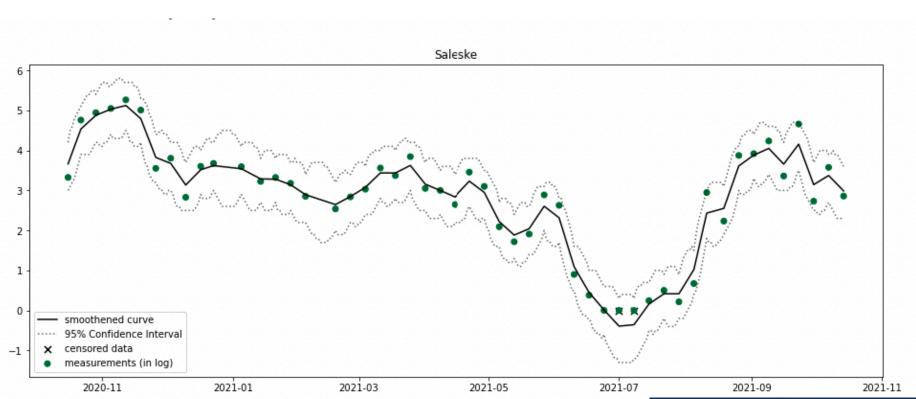








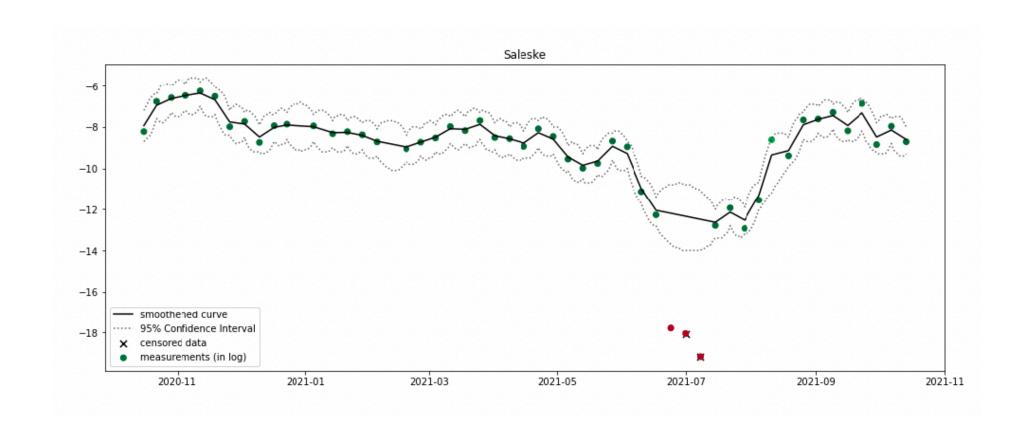


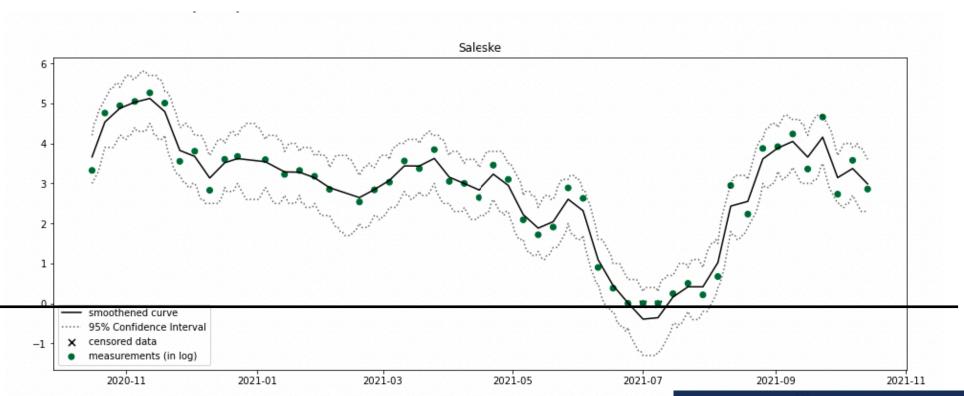








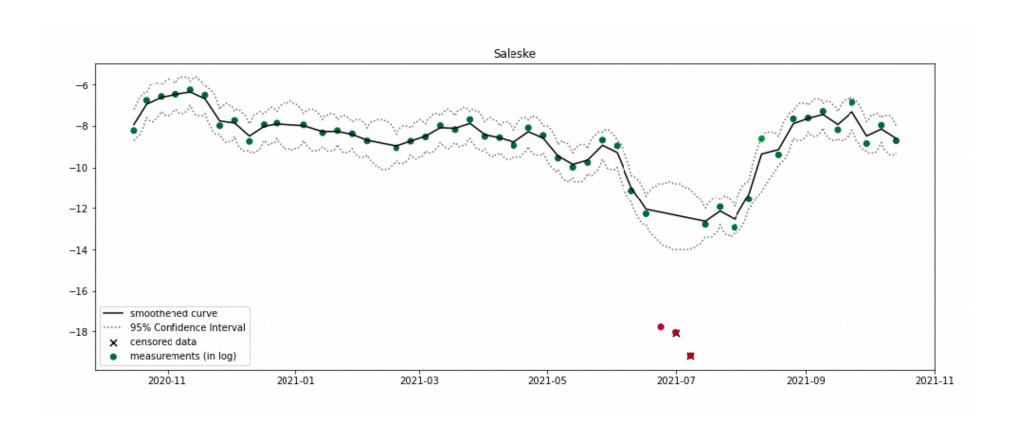












Kranj, Celje, Ljubljana, Maribor, Koper

