Epidemic monitoring:

Estimation of the reproduction number of Covid19

DATASIM

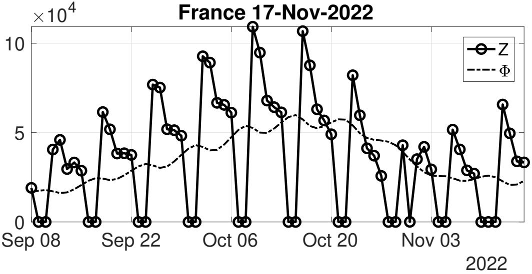
January 25th 2023

Barbara Pascal

Plots of Section III are reproduced with courtesy of N. Pustelnik and J.-C. Pesquet.

Motivation and context: pandemic surveillance

Data: counts of daily new infections



data from National Health Agencies collected by Johns Hopkins University

$\Longrightarrow$ number of cases not informative enough: need to capture the **dynamics**

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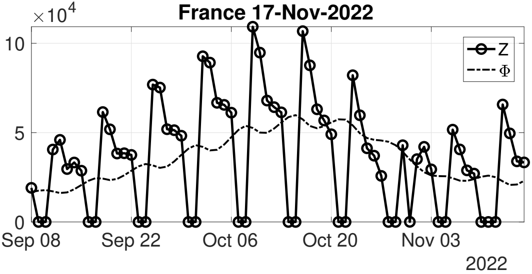

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Goal: design adapted counter measures and evaluate their effectiveness

$\longrightarrow$ efficient monitoring tools *epidemiological model,*

$\longrightarrow$ robust to low quality of the data *managing erroneous counts,*

$\longrightarrow$ **(bonus)** accompanied by reliable confidence level *credibility intervals.*

Outline

I. Epidemic modeling (Cori et al., 2013, *Am. Journal of Epidemiology*)

II. Reproduction number estimation (Pascal et al., 2022, *Trans. Sig. Process.*)

- **A)** maximum likelihood principle
- **B)** variational approaches

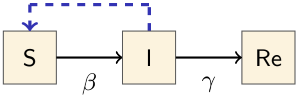
III. Nonsmooth convex optimization (Boyd et al., 2004, *Cambridge University Press*)

- **A)** basic tools and concepts
- **B)** algorithms

IV. Conclusion & Perspectives

I. Epidemic modeling: SIR model

Susceptible-Infected-Recovered (SIR), among *compartmental models*



– ODE: $\dfrac{\mathrm{dS}_t}{\mathrm{d}t} = -\beta \mathsf{S}_t \mathsf{I}_t$, $\dfrac{\mathrm{dI}_t}{\mathrm{d}t} = \beta \mathsf{S}_t \mathsf{I}_t - \gamma \mathsf{I}_t$, $\dfrac{\mathrm{dRe}_t}{\mathrm{d}t} = \gamma \mathsf{I}_t$

– Stochastic model: likelihood maximization to infer β, γ

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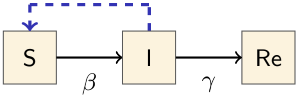

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– Stochastic model: likelihood maximization to infer β, γ

Limitations:

- refinement needed to get socially realistic model
- quadratic increase of the number of parameters
- Bayesian framework: heavy computational burden
- need consolidated and accurate datasets

✗ not adapted to real-time monitoring of Covid19 pandemic

Definition. The reproduction number associated to an epidemic is

"the averaged number of secondary cases generated by a typical infectious individual"

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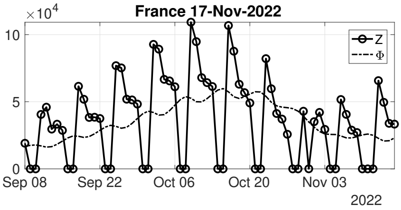
Interpretation. At day t

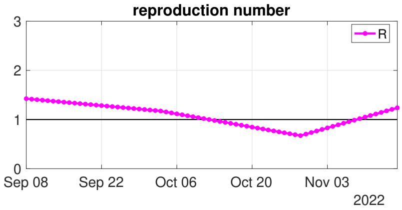
$\mathrm{R}_t > 1$ the virus propagates at exponential speed,

$\mathrm{R}_t < 1$ the epidemic shrinks with an exponential decay,

$\mathrm{R}_t = 1$ the epidemic is stable.

$\Longrightarrow$ one single indicator accounting for the overall pandemic mechanism

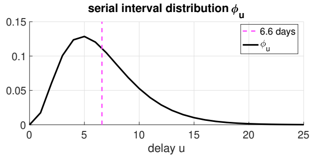




Principle: Z_t new infections at day t

$$\mathbb{E}\left[\mathrm{Z}_t\right] = \mathrm{R}_t \Phi_t, \quad \Phi_t = \sum_{u=1}^{\tau_\Phi} \phi_u \mathrm{Z}_{t-u}$$

with Φ_t global "infectiousness" in the population



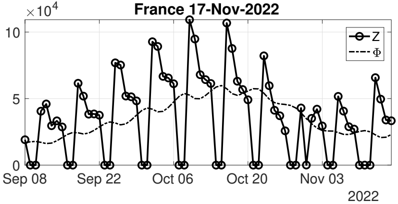
$\{\phi_u\}_{u=1}^{\tau_\Phi}$ distribution of delay between onset of symptoms in primary and secondary cases

Gamma distribution truncated at 25 days, of mean 6.6 days and standard deviation 3.5 days

Data: daily counts $\mathbf{Z} = (Z_1, \ldots, Z_T)$

Model: Poisson distribution

$$\mathbb{P}(\mathsf{Z}_t | \mathbf{Z}_{t-\tau_\Phi:t-1}, \mathsf{R}_t) = \frac{(\mathsf{R}_t \Phi_t)^{\mathsf{Z}_t} \mathrm{e}^{-\mathsf{R}_t \Phi_t}}{\mathsf{Z}_t!}$$



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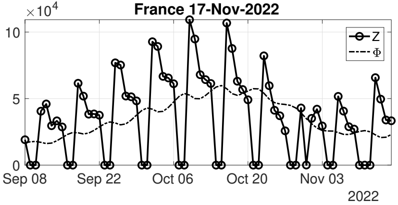

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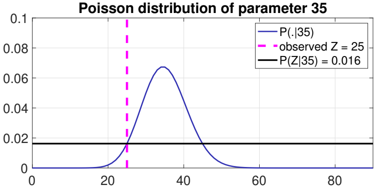
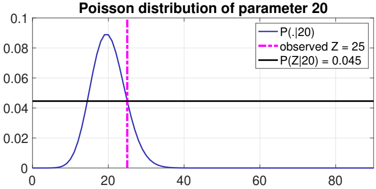
Maximum Likelihood Principle: If one observes a given Z_t, how to infer R_t?





observation $Z_t = 25 \quad \Rightarrow \quad \widehat{R}_t^{\mathrm{MLE}} \Phi_t =$

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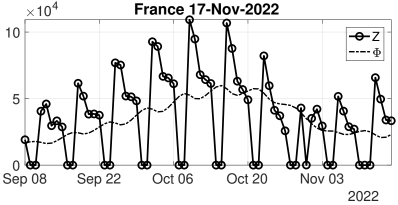

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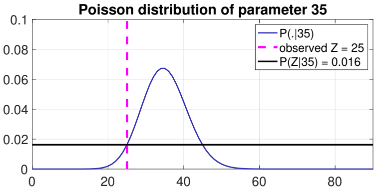

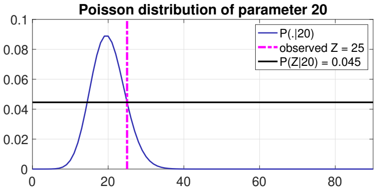

$$\text{observation } Z_t = 25 \quad \Rightarrow \quad \widehat{R}_t^{\text{MLE}} \Phi_t = 25$$

Maximum Likelihood Estimator. $\widehat{\mathrm{R}}_t^{\mathrm{MLE}} := \underset{\mathrm{R}_t}{\operatorname{argmax}}\ \mathbb{P}(\mathrm{Z}_t|\mathbf{Z}_{t-\tau_\Phi:t-1}, \mathrm{R}_t)$

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\ln\left(\mathbb{P}(\mathrm{Z}_t|\mathbf{Z}_{t-\tau_\Phi:t-1}, \mathrm{R}_t)\right) &= \mathrm{Z}_t \ln(\mathrm{R}_t\Phi_t) - \mathrm{R}_t\Phi_t - \ln(\mathrm{Z}_t!) \\
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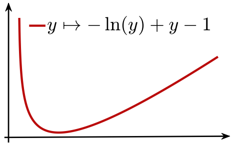
Definition. (Kullback-Leibler divergence)

$$\mathrm{d}_{\mathrm{KL}}(\mathrm{Z}|\mathrm{p}) = \begin{cases} \mathrm{Z}\ln(\mathrm{Z}/\mathrm{p}) + \mathrm{p} - \mathrm{Z} & \text{if } \mathrm{Z} > 0\ \&\ \mathrm{p} > 0 \\ \mathrm{p} & \text{if } \mathrm{Z} = 0\ \&\ \mathrm{p} \geq 0 \\ \infty & \text{otherwise.} \end{cases}$$



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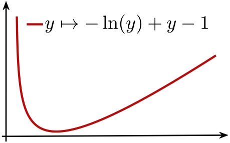

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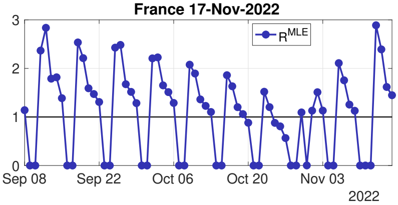

- ♣ huge variability along time/ no local trend
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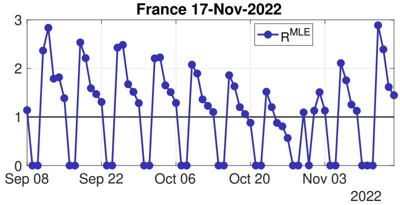

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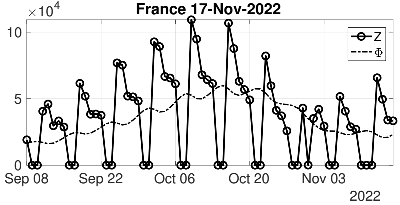
Explanation.



New infection counts **Z** are corrupted by

- missing samples,
- non meaningful negative counts,
- retrospected cumulated counts,
- pseudo-seasonality effects.

State-of-the-art in epidemiology. Smoothing over a temporal window

$$\widehat{\mathrm{R}}^{\mathrm{MLE}}_{t,s}, \text{ with } s = 7 \text{ days}$$

(Cori et al., 2013, *Am. Journal of Epidemiology*)

$\Longrightarrow$ not able to detect rapid surge, nor fast decrease following sanitary restrictions

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Penalized likelihood. Regularization through nonlinear filtering

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with $\mathcal{P}(\mathbf{R})$ favoring some temporal regularity (Abry et al., 2020, *PlosOne*)

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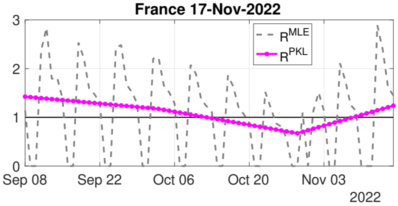
$$\mathcal{P}(\mathbf{R}) = \|\mathbf{D}_2\mathbf{R}\|_1$$

$$(\mathbf{D}_2\mathbf{R})_t = R_{t+1} - 2R_t + R_{t-1}$$

2nd order derivative & ℓ_1-norm

$\Longrightarrow$ piecewise linearity



captures global **trend**, more **regular** than MLE, detect **ruptures**

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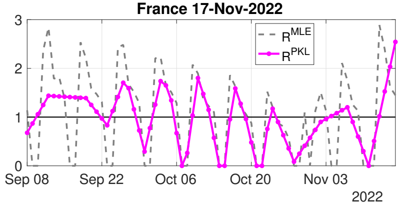
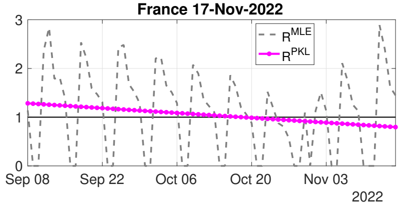
Balance between data-fidelity and temporal regularity.



small λ_{R}



large λ_{R}

Data. Daily reported counts $\mathbf{Z} = (Z_1, \ldots, Z_T)$

Model. Poisson distribution $\mathbb{P}(Z_t | \mathbf{Z}_{t-\tau_\Phi:t-1}, R_t) = \dfrac{(R_t \Phi_t)^{Z_t} e^{-(R_t \Phi_t)}}{Z_t!}$

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properties of the objective function:

- sum of convex functions composed with linear operators $\implies$ globally convex;
- feasible domain: $\{\text{if } Z_t > 0,\ R_t \Phi_t > 0, \text{ else } R_t \Phi_t \geq 0\}$;
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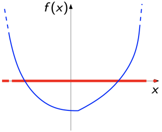
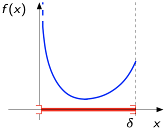
Theorem (Pascal et al., 2022, *Trans. Sig. Process.*)

+ The minimization problem has at least one solution $\widehat{\mathbf{R}}^{\mathrm{PKL}}$.
+ The estimated time-varying Poisson intensity $\widehat{p}_t^{\mathrm{PKL}} = \widehat{R}_t^{\mathrm{PKL}}\Phi_t$ is unique.

Definition. Let $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$, the domain of f is $\operatorname{dom} f = \{\boldsymbol{x} \in \mathbb{R}^T \mid f(x) < \infty\}$



$\operatorname{dom} f = \mathbb{R}$



$\operatorname{dom} f =]0, \delta]$

If $\operatorname{dom} f \neq \emptyset$, f is said to be proper.

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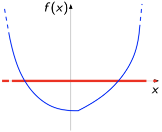

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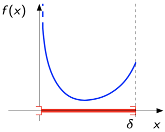

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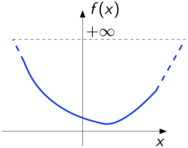
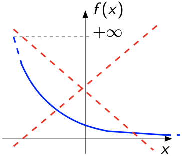
Definition. Let $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$. If

$$\lim_{\|\boldsymbol{x}\|_2 \to \infty} f(\boldsymbol{x}) = \infty$$

then f is said to be coercive.





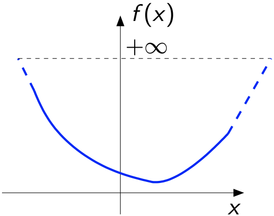
Theorem. If $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$ is proper, continuous on $\operatorname{dom} f$, coercive then

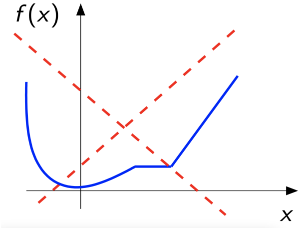
$$\operatorname{Argmin} f = \{\boldsymbol{x} \in \operatorname{dom} f \mid f(\boldsymbol{x}) = \inf f\}$$

is nonempty. If f is convex, then $\operatorname{Argmin} f$ is convex.

Theorem. If $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$ is proper, C^1 on $\operatorname{dom} f$, coercive, and convex

$$\widehat{\boldsymbol{x}} \in \operatorname{Argmin} f \quad \Leftrightarrow \quad \nabla f(\widehat{\boldsymbol{x}}) = 0$$



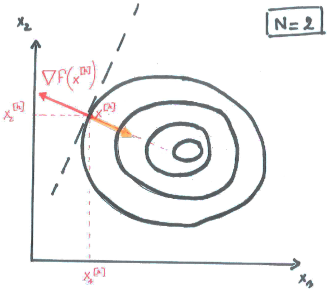


Gradient descent algorithm.

$f : \mathbb{R}^T \to \mathbb{R}$, continuously differentiable

for $k = 1, 2 \dots$ **do**

$$\boldsymbol{x}^{[k+1]} = \boldsymbol{x}^{[k]} - \gamma \nabla f(\boldsymbol{x}^{[k]})$$



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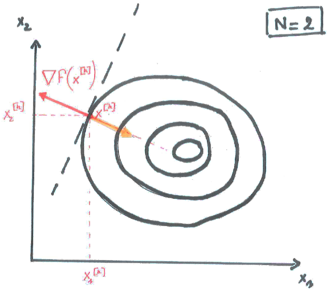

Definition. Let $f : \mathbb{R}^T \to \mathbb{R}$, continuously differentiable, and $\beta > 0$. If

$$\forall \boldsymbol{u}, \boldsymbol{v} \in \mathbb{R}^T, \quad \|\nabla f(\boldsymbol{u}) - \nabla f(\boldsymbol{v})\|_2 \leq \beta \|\boldsymbol{u} - \boldsymbol{v}\|_2$$

f is said to be β-smooth, i.e., f has a β-Lipschitz gradient.

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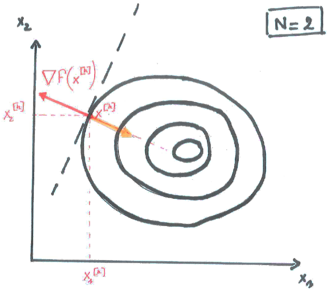

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Theorem. If $f : \mathbb{R}^T \to \mathbb{R}$ is convex, coercive, C^1, and β-smooth, with $\beta > 0$, then

$$\exists \widehat{\boldsymbol{x}} \in \mathbb{R}^T, \quad \lim_{k \to \infty} \boldsymbol{x}^{[k]} = \widehat{\boldsymbol{x}} \quad \text{with} \quad \nabla f(\widehat{\boldsymbol{x}}) = 0.$$

Definition. Let $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$, proper, the <u>subdifferential</u> of f at $\boldsymbol{x}$ is

$$\partial f(\boldsymbol{x}) = \{\boldsymbol{u} \in \mathbb{R}^T \mid \forall \boldsymbol{y} \in \mathbb{R}^T, \quad f(\boldsymbol{y}) \geq f(\boldsymbol{x}) + \langle \boldsymbol{y} - \boldsymbol{x}, \boldsymbol{u} \rangle\}$$

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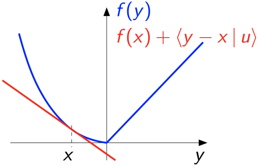

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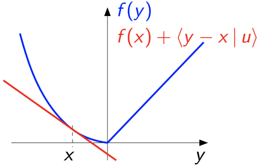

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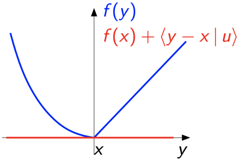
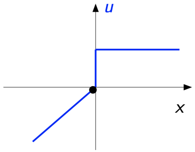


$\partial f(\mathbf{0}) = [0, 1]$



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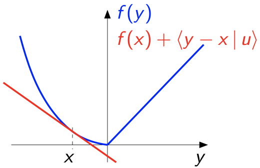

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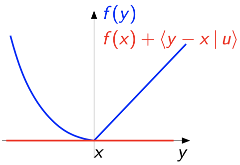

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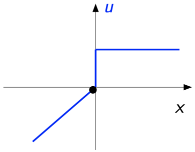

Theorem. (Fermat's rule) Let $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$ a proper function

$$\widehat{\boldsymbol{x}} \in \operatorname{Argmin} f \quad \Leftrightarrow \quad 0 \in \partial f(\widehat{\boldsymbol{x}}).$$

Subgradient descent algorithm.

$f : \mathbb{R}^T \to \mathbb{R}$, convex, continuous

for $k = 1, 2 \dots$ **do**

$$\boldsymbol{x}^{[k+1]} = \boldsymbol{x}^{[k]} - \gamma_k \boldsymbol{u}^{[k]}, \quad \boldsymbol{u}^{[k]} \in \partial f(\boldsymbol{x}^{[k]})$$

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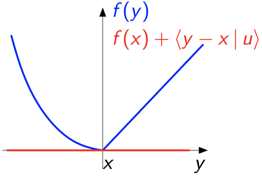

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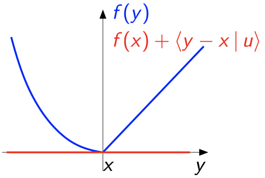

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- need a vanishing sequence $(\gamma_k)_{k\in\mathbb{N}}$: $\gamma_k \underset{k\to\infty}{\to} 0$;
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Explanation. $\partial f : \mathbb{R}^T \to 2^{\mathbb{R}^T}$ set-valued

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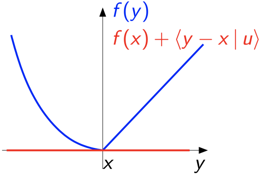

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Solution. Turn to an implicit scheme

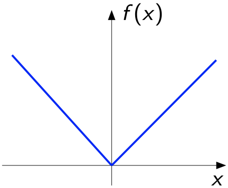
$$\boldsymbol{x}^{[k+1]} = \boldsymbol{x}^{[k]} - \gamma_k \boldsymbol{u}^{[k]}, \quad \boldsymbol{u}^{[k]} \in \partial f(\boldsymbol{x}^{[k+1]}) \Rightarrow \text{how to compute } \boldsymbol{x}^{[k+1]}?$$

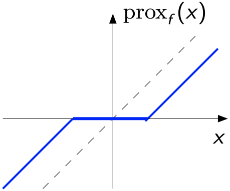
Definition. Let $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$, proper, convex, continuous, $\gamma > 0$

$$\mathrm{prox}_{\gamma f}(\boldsymbol{x}) := \underset{\boldsymbol{y} \in \mathbb{R}^T}{\mathrm{argmin}} \, \frac{1}{2} \|\boldsymbol{x} - \boldsymbol{y}\|_2^2 + \gamma f(\boldsymbol{y})$$

is the proximity operator of γf at point $\boldsymbol{x}$.

Example.





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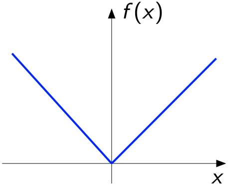

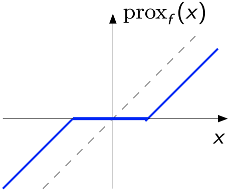

Theorem. Let $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$ a proper, convex, continuous function

$$\boldsymbol{p} = \mathrm{prox}_{\gamma f}(\boldsymbol{x}) \quad \Leftrightarrow \quad \boldsymbol{x} \in \boldsymbol{p} + \partial f(\boldsymbol{p})$$

Implicit scheme.

$$\boldsymbol{x}^{[k+1]} = \boldsymbol{x}^{[k]} - \gamma_k \boldsymbol{u}^{[k]}, \quad \boldsymbol{u}^{[k]} \in \partial f(\boldsymbol{x}^{[k+1]}) \Rightarrow \text{how to compute } \boldsymbol{x}^{[k+1]}?$$

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Solution. Apply the theorem in the $\Leftarrow$ sense with $\boldsymbol{x} = \boldsymbol{x}^{[k]}$ and $\boldsymbol{p} = \boldsymbol{x}^{[k+1]}$

$$\boldsymbol{x}^{[k]} = \boldsymbol{x}^{[k+1]} + \gamma_k \boldsymbol{u}^{[k]}, \quad \boldsymbol{u}^{[k]} \in \partial f(\boldsymbol{x}^{[k+1]})$$

Proximal point algorithm. $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$, proper, convex, continuous

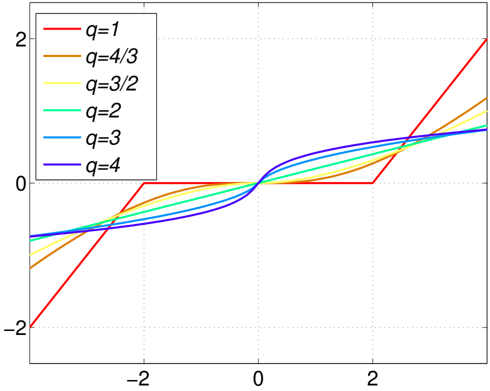
for $k = 1, 2 \ldots$ **do**

$$\boldsymbol{x}^{[k+1]} = \operatorname{prox}_{\gamma f}(\boldsymbol{x}^{[k]})$$

Theorem. For any $\gamma > 0$, $\left(\boldsymbol{x}^{[k]}\right)_{k \in \mathbb{N}}$ converges toward some $\widehat{\boldsymbol{x}} \in \operatorname{Argmin} f$.

Power q function with $q \geq 1$. Let $\eta > 0$, $q \in [1, +\infty[$

$$f : \mathbb{R} \to \mathbb{R} \cup \{\infty\},\ x \mapsto \eta |x|^q$$



many more explicit proximal operators at `http://proximity-operator.net/`

Property. If $f : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$ is separable, i.e.,

$$\forall \boldsymbol{x} \in \mathbf{R}^T, \quad f(\boldsymbol{x}) = \sum_{t=1}^{T} f_t(x_t), \quad \text{with } f_t \text{ proper, convex, continuous}$$

then the proximal operator can be computed component-wise and

$$\boldsymbol{p} = \operatorname{prox}_{\gamma f}(\boldsymbol{x}) \quad \Leftrightarrow \quad \forall t = 1, \ldots, T, \quad p_t = \operatorname{prox}_{\gamma f_t}(x_t).$$

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Problematic. $f, g : \mathbb{R}^T \to \mathbb{R} \cup \{\infty\}$ convex, proper, continuous

$$\underset{\boldsymbol{x} \in \mathbb{R}^T}{\text{minimize}} \; f(\boldsymbol{x}) + g(\boldsymbol{x}).$$

$\Rightarrow$ compute $\operatorname{prox}_{f+g}$: in general intractable!

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Hypotheses. f is continuously differentiable and β-smooth, with $\beta > 0$.
g is proximable, i.e., $\text{prox}_{\gamma g}$ has an explicit formula.

Forward-backward algorithm. or "Proximal-gradient"

for $k = 1, 2 \ldots$ **do**

$$\boldsymbol{x}^{[k+1]} = \text{prox}_{\gamma g}(\boldsymbol{x}^{[k]} - \gamma \nabla f(\boldsymbol{x}^{[k]}))$$

explicit-implicit scheme: $\boldsymbol{x}^{[k+1]} = \boldsymbol{x}^{[k]} - \gamma \nabla f(\boldsymbol{x}^{[k]}) - \gamma \boldsymbol{u}^{[k]}, \; \boldsymbol{u}^{[k]} \in \partial g(\boldsymbol{x}^{[k+1]})$

Theorem. If $\gamma \in]0, 2/\beta[$, $\left(\boldsymbol{x}^{[k]}\right)_{k \in \mathbb{N}}$ converges toward some $\widehat{\boldsymbol{x}} \in \text{Argmin}\, f + g$.

$$\underset{\mathbf{R}\in\mathbb{R}^T}{\text{minimize}} \ \sum_{t=1}^{T} \mathrm{d}_{\mathrm{KL}}\left(\mathrm{Z}_t \,|\, \mathrm{R}_t \Phi_t\right) + \lambda_{\mathrm{R}} \|\mathbf{D}_2 \mathbf{R}\|_1$$

- each term of the functional is convex;
- ℓ_1-norm and indicative functions $\Longrightarrow$ nonsmooth; ✗ gradient descent
- gradient of $\mathrm{p}_t \mapsto \mathrm{d}_{\mathrm{KL}}\left(\mathrm{Z}_t \,|\, \mathrm{p}_t\right)$ is not Lipschitzian; ✗ forward-backward
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$$\Longleftrightarrow \underset{\mathbf{R}\in\mathbb{R}^T}{\text{minimize}} \quad f(\mathbf{R}|\mathbf{Z}) + h(\mathbf{D}_2\mathbf{R}), \quad \mathbf{D}_2 \text{ linear; } f, h \text{ proximable}$$

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Primal-dual algorithm (Chambolle et al., 2011, *Int. Conf. Comput. Vis.*)

for $k = 1, 2\ldots$ **do**

$$\mathbf{Q}^{[k+1]} = \mathbf{Q}^{[k]} + \sigma\mathbf{D}_2\overline{\mathbf{R}}^{[k]} - \sigma\mathrm{prox}_{\sigma^{-1}h}(\sigma^{-1}\mathbf{Q}^{[k]} + \mathbf{D}_2\overline{\mathbf{R}}^{[k]}) \quad \text{dual}$$

$$\mathbf{R}^{[k+1]} = \mathrm{prox}_{\tau f(\cdot|\mathbf{Z})}(\mathbf{R}^{[k+1]} - \tau\mathbf{D}_2^*\mathbf{Q}^{[k+1]}) \quad \text{primal}$$

$$\overline{\mathbf{R}}^{[k+1]} = 2\mathbf{R}^{[k+1]} - \mathbf{R}^{[k]} \quad \text{auxiliary}$$

Theorem. If $\tau\sigma\|\mathbf{D}_2\|_{\mathrm{op}}^2 < 1$, $\left(\mathbf{R}^{[k]}\right)_{k\in\mathbb{N}}$ converges toward $\widehat{\mathbf{R}}^{\mathrm{PKL}}$.

New infection counts per county: $\mathbf{Z} = \left\{ \mathrm{Z}_t^{(d)},\, d \in [1, D],\, t \in [1,\, T] \right\}$

$\Rightarrow$ multivariate time-varying reproduction number $\mathrm{R}_t^{(d)}$

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Multivariate extended penalized Kullback-Leibler

$$\widehat{\mathbf{R}} = \underset{\mathbf{R} \in \mathbb{R}^{D \times T}}{\operatorname{argmin}} \sum_{d=1}^{D} \sum_{t=1}^{T} \mathsf{d}_{\mathsf{KL}} \left(\mathsf{Z}_t^{(d)} \,\middle|\, \mathsf{R}_t^{(d)} \Phi_t^{(d)} \right) + \lambda_{\mathsf{R}} \|\mathbf{D}_2 \mathbf{R}\|_1 + \lambda_{\text{space}} \|\mathbf{G}\mathbf{R}\|_1$$

$\Longrightarrow$ $\|\mathbf{G}\mathbf{R}\|_1$ favors **piecewise constancy** in space

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$\Longrightarrow$ $\|\mathbf{GR}\|_1$ favors **piecewise constancy** in space

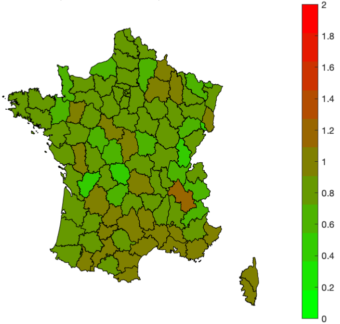
Graph Total Variation

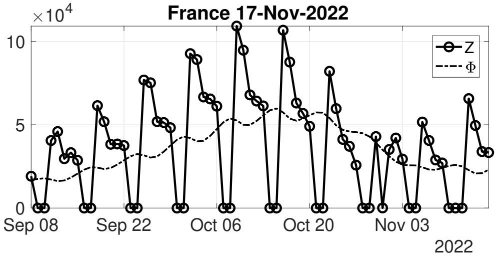
$$\|\mathbf{GR}\|_1 = \sum_{t=1}^{T} \sum_{d_1 \sim d_2} \left| \mathrm{R}_t^{(d_1)} - \mathrm{R}_t^{(d_2)} \right|$$

sum over neighboring counties

here: $d_1 \sim d_2 \Leftrightarrow$ share terrestrial border

France (Hospital based data) - R - 15-Nov-2022

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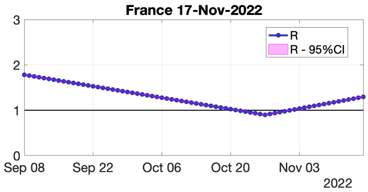
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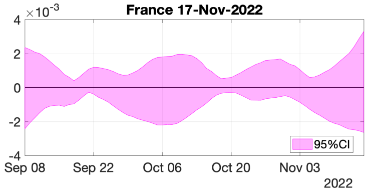
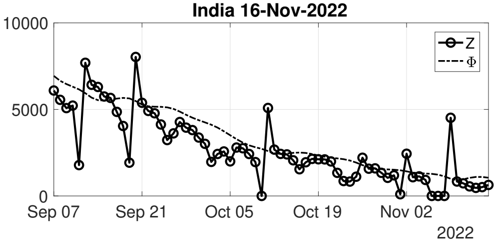
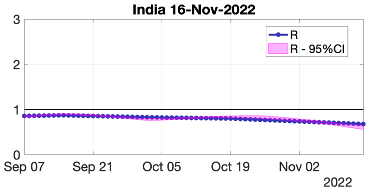
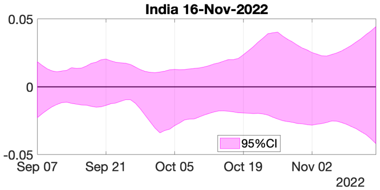

France 17-Nov-2022
×10^-3
4
2
0
-2
-4
95%CI
Sep 08
Sep 22
Oct 06
Oct 20
Nov 03
2022

Worldwide Covid19 monitoring







Why not United Kingdom?

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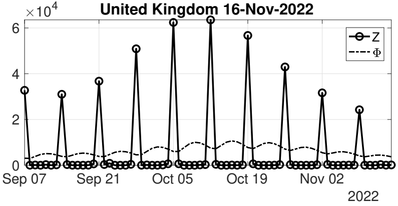
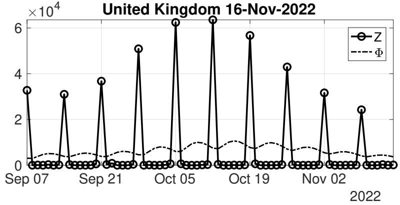


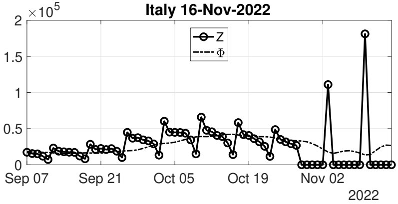
rate of erroneous counts: 6/7!

Why not United Kingdom?



rate of erroneous counts: 6/7!

And Italy?



seems to adopt the same reporting rate ...

$\Longrightarrow$ call for new tools, robust to very scarce data

Pointwise estimate of parameter $\boldsymbol{\theta} = \mathbf{R}$ from observations $\mathbf{Z}$

$$\underset{\mathbf{R} \in \mathbb{R}^T}{\text{minimize}} \quad f(\boldsymbol{\theta}|\mathbf{Z}) + h(\mathbf{A}\boldsymbol{\theta}) \quad \text{(Pascal et al., 2022, \textit{Trans. Sig. Process.})}$$

Bayesian framework for credibility interval estimation

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Q: what is the value of R today? **R:** solve the minimization problem and output $\widehat{\mathrm{R}}_T$.

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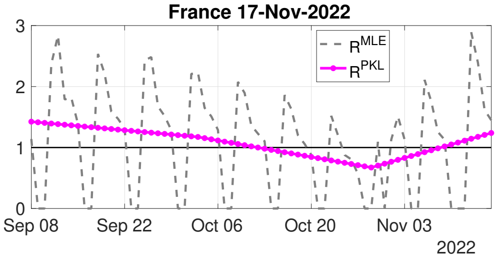

$\widehat{\mathrm{R}}_T = 1.2955$

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Bayesian reformulation: interpret $\widehat{\mathbf{R}}^{\mathrm{PL}}$ as the Maximum A Posteriori of

$$\pi(\boldsymbol{\theta}) \propto \exp(-f(\boldsymbol{\theta}|\mathbf{Z}) - h(\mathbf{A}\boldsymbol{\theta}))$$

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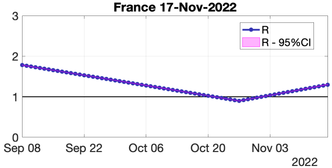
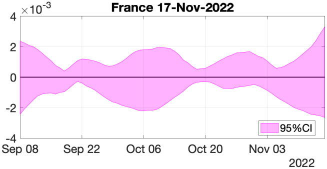
$\Longrightarrow$ instead of focusing on $\widehat{\mathrm{R}}_t$, the **pointwise** MAP, probe π to get

$\mathrm{R}_t \in [\underline{\mathrm{R}}_t, \overline{\mathrm{R}}_t]$ with 95% probability, i.e., **credibility interval** estimates





$$\widehat{\mathrm{R}}_T \in [1.2987, 1.3047]$$

Markov Chain Monte Carlo sampling

Purpose: sampling the random variable $\boldsymbol{\theta} = \mathbf{R} \in \mathbb{R}^T$ according to the posterior[†]

$$\pi(\boldsymbol{\theta}) \propto \exp\left(-f(\boldsymbol{\theta}) - g(\boldsymbol{\theta})\right) \mathbb{1}_{\mathcal{D}}(\boldsymbol{\theta})$$

[†] π is defined up to a normalizing constant

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Principle: 1) generate a random sequence $\{\boldsymbol{\theta}^n, n \in \mathbb{N}\}$ such that

- $\boldsymbol{\theta}^{n+1}$ only depends on $\boldsymbol{\theta}^n$,
- at convergence, i.e., as $n \to \infty$, $\boldsymbol{\theta}^n \sim \pi$,

2) compute Bayesian estimators, e.g., credibility intervals, on samples $\{\boldsymbol{\theta}^n, n \geq N\}$

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State-of-the-art: *Hastings-Metropolis random walk*

(i) propose a random move according to

$$\boldsymbol{\theta}^{n+\frac{1}{2}} = \boldsymbol{\theta}^n + \sqrt{2\gamma}\Gamma\xi^{n+1}, \quad \xi^{n+1} \sim \mathcal{N}_T(0, \mathrm{I})$$

with γ positive step size, $\Gamma \in \mathbb{R}^{T \times T}$

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(ii) accept: $\boldsymbol{\theta}^{n+1} = \boldsymbol{\theta}^{n+\frac{1}{2}}$, with probability $1 \wedge \dfrac{\pi(\boldsymbol{\theta}^{n+\frac{1}{2}})}{\pi(\boldsymbol{\theta}^n)}$, or reject: $\boldsymbol{\theta}^{n+1} = \boldsymbol{\theta}^n$

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Metropolis Adjusted Langevin Algorithm (MALA)

Langevin dynamics: $\boldsymbol{\theta}^{n+\frac{1}{2}} = \mu(\boldsymbol{\theta}^n) + \sqrt{2\gamma}\xi^{n+1}$, (Kent, 1978, *Adv Appl Probab*)

$$\mu(\boldsymbol{\theta}) \text{ adapted to } \pi(\boldsymbol{\theta}) = \exp(-f(\boldsymbol{\theta}) - g(\boldsymbol{\theta}))\mathbb{1}_{\mathcal{D}}(\boldsymbol{\theta})$$

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Case 1: $g = 0$ and $-\ln \pi = f$ is smooth (Roberts & Tweedie, 1996, *Bernoulli*)

$$\mu(\boldsymbol{\theta}) = \boldsymbol{\theta} - \gamma \Gamma\Gamma^\top \nabla f(\boldsymbol{\theta}) = \boldsymbol{\theta} + \gamma \Gamma\Gamma^\top \nabla \ln \pi(\boldsymbol{\theta})$$

$\Longrightarrow$ move towards areas of higher probability

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Case 2: $-\ln\pi = f + g$ is nonsmooth

$$\mu(\boldsymbol{\theta}) = \mathrm{prox}_{\gamma g}^{\Gamma\Gamma^\top}(\boldsymbol{\theta} - \gamma\Gamma\Gamma^\top\nabla f(\boldsymbol{\theta}))$$

combining *Langevin* and *proximal*[†] approaches

[†] $\mathrm{prox}_{\gamma g}^{\Gamma\Gamma^\top}(y) = \underset{x\in\mathbb{R}^d}{\mathrm{argmin}}\left(\frac{1}{2}\|x-y\|^2_{\Gamma\Gamma^\top} + \gamma g(x)\right)$: preconditioned proximity operator of g

Proximal-Gradient dual sampler PGdual

Posterior density of $\boldsymbol{\theta} = \mathbf{R}$: $\quad \pi(\boldsymbol{\theta}) \propto \exp\left(-f(\boldsymbol{\theta}) - g(\boldsymbol{\theta})\right) \mathbb{1}_{\mathcal{D}}(\boldsymbol{\theta})$

- smooth negative log-likelihood

$$\text{if } \boldsymbol{\theta} \in \mathcal{D}, \quad f(\boldsymbol{\theta}) = -\sum_{t=1}^{T} \left(\mathsf{Z}_t \ln \mathsf{p}_t(\boldsymbol{\theta}) - \mathsf{p}_t(\boldsymbol{\theta})\right), \quad \mathsf{p}_t(\boldsymbol{\theta}) = \mathsf{R}_t(\Phi \mathsf{Z})_t$$

- nonsmooth convex lower-semicontinuous negative a priori log-distribution

$$g(\boldsymbol{\theta}) = \lambda_{\mathsf{R}} \|\mathbf{D}_2 \mathbf{R}\|_1 = h(\mathbf{A}\boldsymbol{\theta})$$

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Case 3: $-\ln \pi = f + h(\mathbf{A}\cdot)$ (Fort et al., 2022, *preprint*)

closed-form expression of $\text{prox}_{\gamma h}$ but not of $\text{prox}_{\gamma h(\mathbf{A}\cdot)}$

1) extend $\mathbf{A}$ into **invertible** $\overline{\mathbf{A}}$, and h in $\bar{h}$ such that $\bar{h}(\overline{\mathbf{A}}\boldsymbol{\theta}) = h(\mathbf{A}\boldsymbol{\theta})$

2) reason on the **dual** variable $\tilde{\boldsymbol{\theta}} = \overline{\mathbf{A}}\boldsymbol{\theta}$

Markov Chain Monte Carlo sampling scheme

Proximal-gradient drift. $\mu(\boldsymbol{\theta}) = \overline{\mathbf{A}}^{-1} \mathrm{prox}_{\gamma \bar{h}} \left(\overline{\mathbf{A}} \boldsymbol{\theta} - \gamma \overline{\mathbf{A}}^{-\top} \nabla f(\boldsymbol{\theta}) \right)$

Data: $\overline{\mathbf{D}} = \overline{\mathbf{D}}_2$ (`Invert`) or $\overline{\mathbf{D}} = \overline{\mathbf{D}}_o$ (`Ortho`)
$\gamma_{\mathrm{R}} > 0$, $N_{\max} \in \mathbb{N}_\star$, $\boldsymbol{\theta}^0 = \mathbf{R}^0 \in \mathcal{D}$
Result: A $\mathcal{D}$-valued sequence $\{\boldsymbol{\theta}^n = \mathbf{R}^n, n \in 0, \dots, N_{\max}\}$
for $n = 0, \dots, N_{\max} - 1$ **do**
Sample $\xi_{\mathrm{R}}^{n+1} \sim \mathcal{N}_T(0, \mathrm{I})$;
Set $\mathbf{R}^{n+\frac{1}{2}} = \mu(\boldsymbol{\theta}^n) + \sqrt{2\gamma_{\mathrm{R}}} \overline{\mathbf{D}}^{-1} \overline{\mathbf{D}}^{-\top} \xi_{\mathrm{R}}^{n+1}$;
$\boldsymbol{\theta}^{n+\frac{1}{2}} = \mathbf{R}^{n+\frac{1}{2}}$;
Set $\boldsymbol{\theta}^{n+1} = \boldsymbol{\theta}^{n+\frac{1}{2}}$ with probability

$$1 \wedge \frac{\pi(\boldsymbol{\theta}^{n+\frac{1}{2}})}{\pi(\boldsymbol{\theta}^n)} \frac{q_{\mathrm{R}}(\boldsymbol{\theta}^{n+\frac{1}{2}}, \boldsymbol{\theta}^n)}{q_{\mathrm{R}}(\boldsymbol{\theta}^n, \boldsymbol{\theta}^{n+\frac{1}{2}})},$$

q_{R} : Gaussian kernel stemming from nonsymmetric proposal

and $\boldsymbol{\theta}^{n+1} = \boldsymbol{\theta}^n$ otherwise.

Algorithm 1: Proximal-Gradient dual: `PGdual Invert` and `PGdual Ortho`

Few illustrations and resources

- Daily estimation of the reproduction number of Covid19
  https://perso.ens-lyon.fr/patrice.abry/Covid_France_trendOutlier.png
- Estimation of R_t with credibility intervals
  https://perso.math.univ-toulouse.fr/gfort/project/opsimore-2/
- A broad audience article The Conversation France