

Estimation of the reproduction number of the Covid19 pandemic

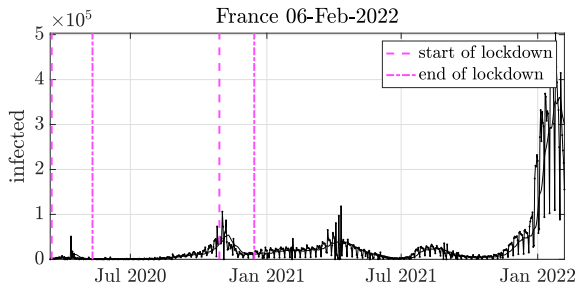
Maximum A Posteriori and credibility intervals

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Steniq, SigMA team meeting

Data: number of daily new infection counts



(partial) lockdown

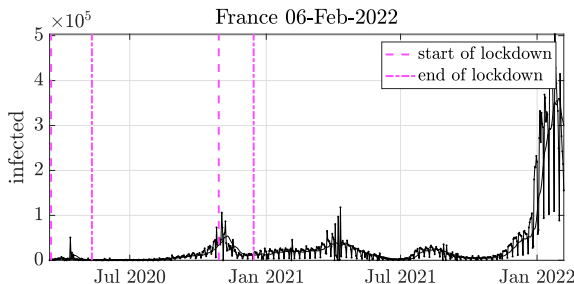
Indicator: reproduction number R_0

averaged number of people contaminated by one infected person

$R_0 > 1$: the virus propagates,

$R_0 < 1$: the epidemic slows down.

Taking, e.g., lockdown, measures requires a **real-time**, daily, estimate R_t .



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

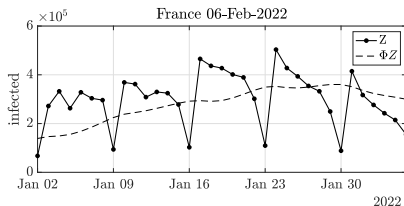
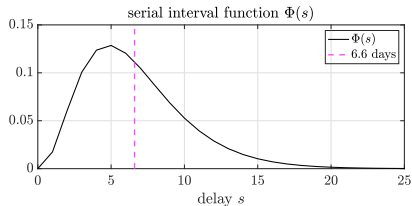
- refinement needed to get socially realistic model
- quadratic increase of the number of parameters
- Bayesian framework: heavy computational burden
- need consolidated and accurate datasets

Poisson process accounting for random contamination

$$Z_t \sim \text{Poisson}(p_t), \quad p_t = R_t \sum_{s=1}^{\tau_\Phi} \Phi(s) Z_{t-s}$$

$\Phi(s)$: **serial interval function** $\tau_\Phi = 26$ days

random delay between onset of symptoms in primary and secondary cases



> modeled by a Gamma distribution with mean and variance of 6.6 and 3.5 days

Unknown parameters: $\mathbf{R} = (R_1, \dots, R_T) \in \mathbb{R}_+^T$

Observed data: $\mathbf{Z} = (Z_1, \dots, Z_T)$

Poisson distribution of parameter $p_t = R_t \sum_{s=1}^{\tau_\Phi} \Phi(s) Z_{t-s}$

$$\mathbb{P}(Z_t | \mathbf{Z}_{t-\tau_\Phi:t-1}, R_t) = \frac{p_t^{Z_t} e^{-p_t}}{Z_t!}$$

> negative log-likelihood

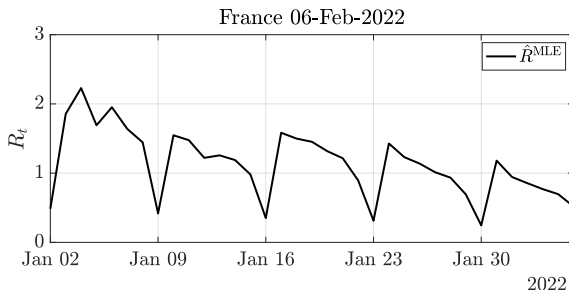
$$\begin{aligned} -\ln(\mathbb{P}(Z_t | \mathbf{Z}_{t-\tau_\Phi:t-1}, R_t)) &= p_t - Z_t \ln(p_t) + \ln(Z_t!) \\ &\underset{Z_t \gg 1}{\simeq} p_t - Z_t \ln(p_t) + Z_t \ln(Z_t) - Z_t \\ &\underset{(\text{def.})}{=} d_{\text{KL}}(Z_t | p_t) \quad \text{Kullback-Leibler divergence} \end{aligned}$$

Maximum Likelihood Estimate

> maximizing the likelihood is equivalent to minimizing $-\ln \mathbb{P}$

$$\hat{\mathbf{R}}^{\text{MLE}} = \underset{\mathbf{R} \in \mathbb{R}_+^T}{\operatorname{argmin}} \sum_{t=1}^T d_{\text{KL}}(Z_t | R_t(\Phi Z)_t), \quad (\Phi Z)_t \triangleq \sum_{s=1}^{\tau_\Phi} \Phi(s) Z_{t-s}$$

Explicit solution: $\hat{R}_t^{\text{MLE}} = Z_t / (\Phi Z)_t$



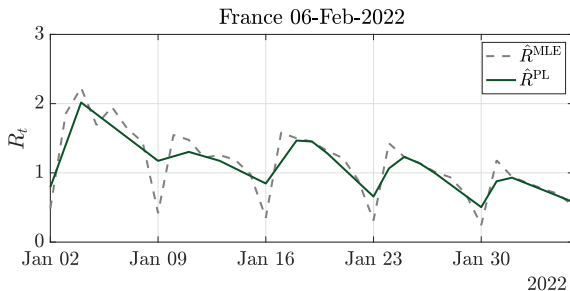
not realistic! pseudo-periodicity, irregularity, no local trend

Penalized log-likelihood

> favor piecewise linear behavior

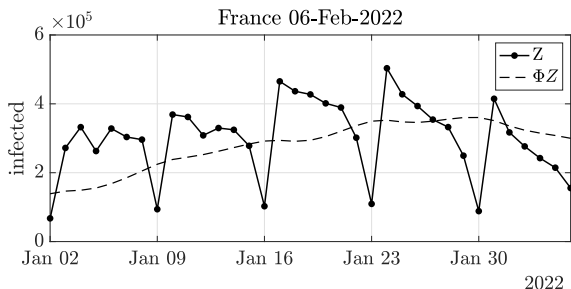
$$\hat{\mathbf{R}}^{\text{PL}} = \underset{\mathbf{R} \in \mathbb{R}_+^T}{\operatorname{argmin}} \sum_{t=1}^T d_{\text{KL}}(Z_t | R_t(\Phi Z)_t) + \lambda_{\text{time}} \|\mathbf{D}_2 \mathbf{R}\|_1$$

$(\mathbf{D}_2 \mathbf{R}) = R_{t+1} - 2R_t + R_{t-1}$ second order discrete derivative



better, but still pseudo-oscillations

Managing low quality of the data



New infection counts $\mathbf{Z} = (Z_1, \dots, Z_T)$ are corrupted by

- missing samples,
- non meaningful negative counts,
- retrospected cumulated counts spread over few days,
- pseudo-seasonality effects, with less counts on non working days, ...

> parametric modeling out of reach

Nonstationary Poisson process with *outliers*

$$Z_t \sim \text{Pois}(R_t(\Phi Z)_t + O_t)$$

O_t : significant values, concentrated on specific days (Sundays, day-offs, ...)

Unknown parameters: $(\mathbf{R}, \mathbf{O}) = (R_1, \dots, R_T, O_1, \dots, O_T) \in \mathbb{R}_+^T \times \mathbb{R}^T$

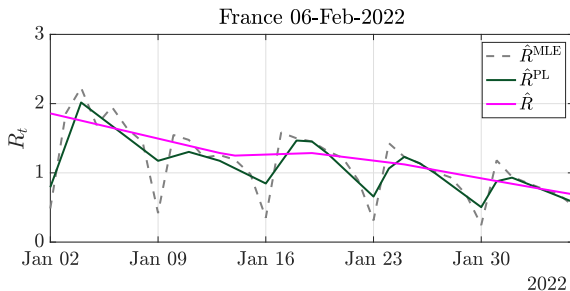
Extended penalized log-likelihood

$$(\hat{\mathbf{R}}, \hat{\mathbf{O}}) = \underset{(\mathbf{R}, \mathbf{O}) \in \mathbb{R}_+^T \times \mathbb{R}^T}{\operatorname{argmin}} \sum_{t=1}^T d_{\text{KL}}(Z_t | R_t(\Phi Z)_t + O_t) + \lambda_{\text{time}} \|\mathbf{D}_2 \mathbf{R}\|_1 + \lambda_{\text{O}} \|\mathbf{O}\|_1$$

> favors piecewise linear reproduction number and sparse outliers

Nonsmooth convex optimization for the estimation of \mathbf{R}

$$\left(\hat{\mathbf{R}}, \hat{\mathbf{O}}\right) = \underset{(\mathbf{R}, \mathbf{O}) \in \mathbb{R}_+^T \times \mathbb{R}^T}{\operatorname{argmin}} \quad \mathrm{D}_{\mathrm{KL}}(\mathbf{Z} | \mathbf{R} \cdot \Phi \mathbf{Z} + \mathbf{O}) + \lambda_{\text{time}} \|\mathbf{D}_2 \mathbf{R}\|_1 + \lambda_{\mathbf{O}} \|\mathbf{O}\|_1$$

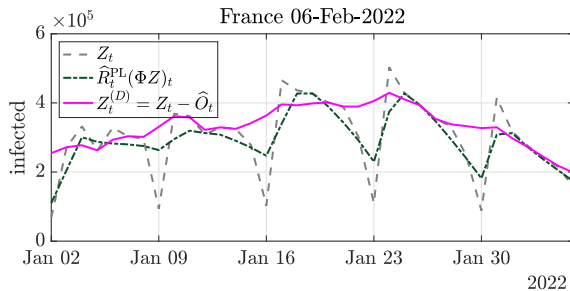


> no more pseudo-seasonality, local trends well captured, smooth behavior

Denoised infection counts

As a byproduct

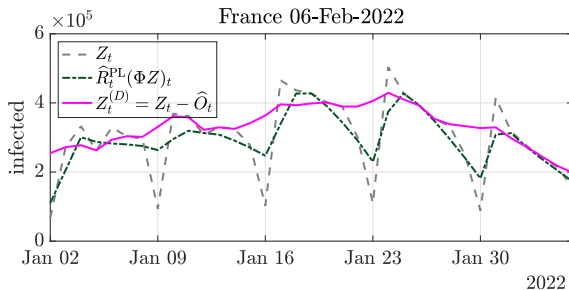
$$\hat{\mathbf{z}}^{(D)} = \mathbf{z} - \hat{\mathbf{o}}$$



Denoised infection counts

As a byproduct

$$\hat{\mathbf{Z}}^{(D)} = \mathbf{Z} - \hat{\mathbf{O}}$$



> level of confidence in the reproduction number \hat{R}_t and corrected count $\hat{Z}_t^{(D)}$?

Idea: interpret the minimization problem

$$\underset{\mathbf{R}, \mathbf{O}}{\text{minimize}} \quad D_{\text{KL}}(\mathbf{Z} | \mathbf{R} \cdot \Phi \mathbf{Z} + \mathbf{O}) + \lambda_{\text{time}} \|\mathbf{D}_2 \mathbf{R}\|_1 + \lambda_{\text{O}} \|\mathbf{O}\|_1$$

as a Maximum A Posteriori estimate of the parameter $\theta = (\mathbf{R}, \mathbf{O})$.

> θ , \mathbf{Z} realizations of random vectors whose distributions are to be specified

Purpose: reformulation of the estimation in a Bayesian framework

$$\hat{\theta} = \underset{\theta}{\operatorname{argmax}} \pi(\theta | \mathbf{Z})$$

π the density of the a posteriori distribution.

Quantiles of the distribution $\pi \implies$ **credibility intervals**

A posteriori distribution

$$\pi(\boldsymbol{\theta}|\mathbf{Z}) \sim \prod_{t=1}^T \underbrace{\mathbb{P}(Z_t|\mathbf{Z}_{1:t-1}, R_t, O_t)}_{\text{likelihood}} \underbrace{\mathbb{P}(R_t, O_t|\mathbf{R}_{1:t-1}, \mathbf{O}_{1:t-1})}_{\text{prior}}$$

Likelihood: standard Kullback-Leibler based for Poisson model

$$\mathbb{P}(Z_t|\mathbf{Z}_{1:t-1}, R_t, O_t) \propto \exp(-d_{\text{KL}}(Z_t|R_t(\Phi Z)_t + O_t))$$

Prior: **R** and **O** supposed *mutually independent*

- Laplace auto-regressive AR(2) for R_t , $\forall t > 2$

$$\mathbb{P}(R_t|\mathbf{R}_{t-2:t-1}) = \frac{\lambda_{\text{time}}}{2} \exp(-\lambda_{\text{time}}|R_t - 2R_{t-1} + R_{t-2}|)$$

- independent Laplace for O_t , $\forall t > 0$

$$\mathbb{P}(O_t) = \frac{\lambda_O}{2} \exp(-\lambda_O|O_t|)$$

Markov chain Monte Carlo $\theta = (\mathbf{R}, \mathbf{O})$

Purpose: sample from the distribution

$$\pi(\theta|\mathbf{Z}) \sim \exp \left(- \sum_{t=1}^T d_{\text{KL}}(Z_t | R_t(\Phi Z)_t + O_t) \right. \\ \left. - \lambda_{\text{time}} \sum_{t=3}^T |R_t - 2R_{t-1} + R_{t-2}| - \lambda_O \sum_{t=1}^T |O_t| \right)$$

To be compared with

$$\underset{\mathbf{R}, \mathbf{O}}{\text{minimize}} \quad D_{\text{KL}}(\mathbf{Z} | \mathbf{R} \cdot \Phi \mathbf{Z} + \mathbf{O}) + \lambda_{\text{time}} \|\mathbf{D}_2 \mathbf{R}\|_1 + \lambda_O \|\mathbf{O}\|_1$$

MCMC principle: $\pi(\theta|\mathbf{Z})$ is **intractable**, thus

generate a sequence $\{\theta^n, n \geq 0\}$ with π as invariant measure

\implies for n sufficiently large, $\theta^n \sim \pi(\theta|\mathbf{Z})$ are representative samples

Metropolis Hastings with Gaussian proposals

Drift term: $\xi^{n+1} \sim \mathcal{N}(\mathbf{0}_{2T}, \mathbf{C})$, and $\mathbf{C} \in \mathbb{R}^{2T \times 2T}$ covariance matrix

$$\theta^{n+1/2} = \mu(\theta^n) + \xi^{n+1}$$

Langevin: $\mu = \mathbf{I} - \gamma \nabla(-\ln \pi)$ drives the chain toward *high probability* regions

for the Covid19 application $\ln \pi$ **not differentiable**

> we proposed a *proximal* Langevin strategy, with $\nabla \rightarrow \partial f$ *sub-differential*

$$\mu(\theta) = \text{prox}_{\gamma \|\cdot\|_1}(\theta - \gamma \nabla D_{\text{KL}}(\theta))$$

Acceptance-rejection Metropolis mechanism: q Gaussian kernel

> set $\theta^{n+1} = \theta^{n+1/2}$ with probability

$$\min \left(1, \frac{\pi(\theta^{n+1/2})}{\pi(\theta^n)} \frac{q(\theta^n, \theta^{n+1/2})}{q(\theta^{n+1/2}, \theta^n)} \right)$$

5% credibility intervals for Covid19 reproduction number

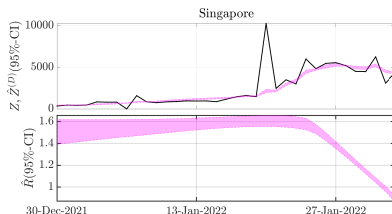
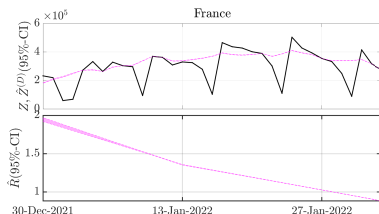
Data from *Johns Hopkins University*

<https://coronavirus.jhu.edu/>

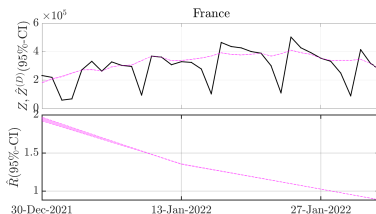
- from National Health Authorities
- 200+ countries
- since the outbreak of the pandemic
- updated on a daily basis

Credibility intervals for R and $Z^{(D)}$

- > $\lambda_{\text{time}} = 3.5 \times \text{std}(\mathbf{Z})$, λ_O for **all** countries/time periods
- > 10 million points in the Markov chain $\{\theta^n, n \geq 0\}$



Take home message



- real-time credibility interval estimate of R_t
 > really informative for Health Authorities
- leverage connection between *variational* and *Bayesian* formulations
- using nonsmooth optimization tools smarter than random walk

 > Estimate of R_t from convex optimization:

<https://perso.ens-lyon.fr/patrice.abry/>

 > Credibility interval estimates of R_t :

<https://perso.math.univ-toulouse.fr/gfort/>