

StatGroup19 empirical models for the Italian epidemic indicators

P. Alaimo Di Loro M. Mingione

F. Divino A. Farcomeni G. Jona Lasinio G. Lovison A. Maruotti

ECDC Forecast Hub Seminar

October 27, 2021



1 Introduction

- Modeling COVID-19 evolution
- The data

2 Incidence indicators

- Richards driven GLM model
- Fast estimation with independent components
- Bayesian space-time extension

3 Prevalence indicators

- An ensemble approach for the nowcasting of ICUs

4 Discussion

Compartmental models - **SIR** (Diekmann et al., 2012)

- **Features**

- **Epidemiologically** correct models
- **Mechanistically** describe the underlying **dynamic**
- Can model the pandemic **all-round**

Compartmental models - SIR (Diekmann et al., 2012)

- **Features**

- **Epidemiologically** correct models
- **Mechanistically** describe the underlying **dynamic**
- Can model the pandemic **all-round**

- **Pitfalls** (Ioannidis et al., 2020)

- Based on **chaotic systems**
- Rely on **hypothetical constants** and **require accurate data**
- **Bias** and **misspecification** lead to completely different results

Compartmental models - SIR (Diekmann et al., 2012)

● Features

- **Epidemiologically** correct models
- **Mechanistically** describe the underlying **dynamic**
- Can model the pandemic **all-round**

● Pitfalls (Ioannidis et al., 2020)

- Based on **chaotic systems**
- Rely on **hypothetical constants** and **require accurate data**
- **Bias** and **misspecification** lead to completely different results

At the beginning of the Italian pandemic
most **forecasts** proved to be completely **wrong!**

Errors and heterogeneity

- Different transmission and data collection system
- Measurement errors and errors in data entry
- Delays in reporting
- Temporal misalignment between indicators
- Tracking was highly symptoms driven



Phenomenological models (Chowell et al., 2016)

- **Features**

- **Macroscopic** models
- Draw inference from **global statistical properties** of the data
- **Statistical artifacts** modeling the observables' mean

Phenomenological models (Chowell et al., 2016)

- **Features**

- **Macroscopic** models
- Draw inference from **global statistical properties** of the data
- **Statistical artifacts** modeling the observables' mean

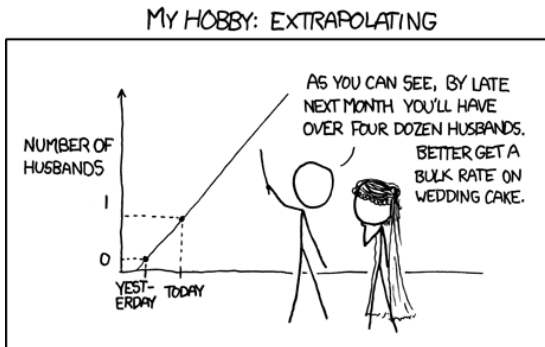
- **Pitfalls**

- Entirely ignore the **nature of the data**
- Assume **Gaussianity**
- Use flexible (non-semiparametric) models to **extrapolate**

Common conceptual mistakes

Pick a model, validate its performances, but **be aware of its limitations!**

- Never ignore **confidence intervals**
- Good fit does not imply good **extrapolation**
- Predictions at **unreasonable time horizons** are **unreliable**



- **Data sources:** GitHub repository of *Italian Protezione Civile*
<https://github.com/pcm-dpc/COVID-19>
- **Detail:** collected and updated daily by the regional Health Systems
- **Type (main):**
 - **Prevalence:** current positives, intensive care occupancy (stock)

$$Y_t = Y_{t-1} + I_t - O_t$$

- **Incidence:** cumulative positives, cumulative deceased (flow)

$$Y_t^C = Y_{t-1}^C + \mathbf{Y}_t$$

- **Data sources:** Gi tHub repository of *Italian Protezione Civile*
<https://github.com/pcm-dpc/COVID-19>
- **Detail:** collected and updated daily by the regional Health Systems
- **Type (main):**
 - **Prevalence:** current positives, intensive care occupancy (stock)

$$Y_t = Y_{t-1} + I_t - O_t$$

- **Incidence:** cumulative positives, cumulative deceased (flow)

$$Y_t^C = Y_{t-1}^C + \mathbf{Y}_t$$

Logistic curves

- *S-shaped* curves
- Widely used to model various **growth** phenomena (biological, population, etc.)
- **Exponential** growth followed by a sudden **level off**

Logistic curves

- *S-shaped* curves
- Widely used to model various **growth** phenomena (biological, population, etc.)
- **Exponential** growth followed by a sudden **level off**

In the **epidemic** context

- Finite elements solutions/approximations to epidemiological **ode**
- Describe the **macroscopic** behavior of an **infection trajectory**
- Fit **globally** on the data respecting their **epidemic** nature

Cumulative counts follow a Logistic Growth

- Model the mean as a **modified Richards' curve**

$$E[Y_t^c] = \lambda_\gamma(t) = b \left(t + \frac{r}{[1 + 10^{h(p-t)}]^s} \right), \quad \gamma = [b, r, h, p, s]^>$$

Cumulative counts follow a Logistic Growth

- Model the mean as a **modified Richards' curve**

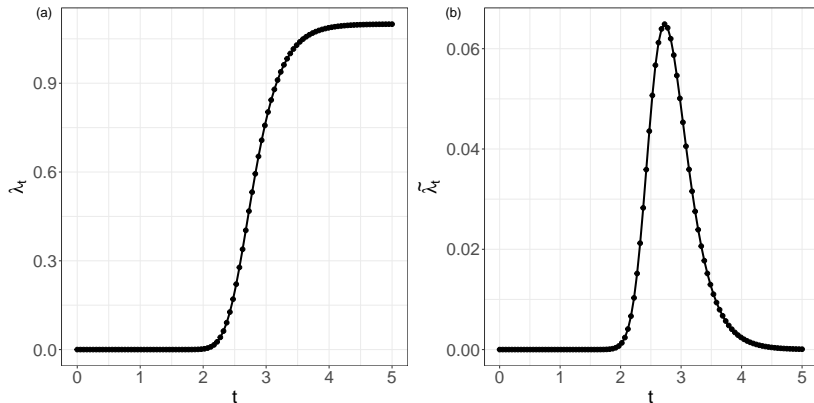
$$E[Y_t^c] = \lambda_\gamma(t) = b \cdot t + \frac{r}{[1 + 10^{h(p-t)}]^s}, \quad \gamma = [b, r, h, p, s]^>$$



- Innovations follow **first differences**

$$E[\mathbf{Y}_t] = E[Y_t^c] - E[Y_{t-1}^c] = \lambda_\gamma(t) - \lambda_\gamma(t-1)$$
$$\frac{d}{dt} \lambda_\gamma(t) = \tilde{\lambda}_\gamma(\mathbf{t}) = \mathbf{b} + \mathbf{f}_\lambda(\mathbf{t})$$

Richards' driven GLM



Example of a Richards' curve (a) and its first differences (b).

Modeling key-points of Alaimo Di Loro et al. (2021)

- Consider the effect of **covariates** through a link function

$$\eta_{\beta}(\mathbf{X}) = \beta\mathbf{X} \quad) \quad g_{\beta}(\mathbf{X}) = \exp f_{\eta_{\beta}}(\mathbf{X})g$$

- Additive:**

$$\mu_{\theta}(t, \mathbf{X}) = b_{\beta}(\mathbf{X}) + r \tilde{\lambda}_{\gamma}(t), \quad b_{\beta}(\mathbf{X}) = g_{\beta}(\mathbf{X})$$

- Multiplicative:**

$$\mu_{\theta}(t, \mathbf{X}) = b + r_{\beta}(\mathbf{X}) \tilde{\lambda}_{\gamma}(t), \quad r_{\beta}(\mathbf{X}) = g_{\beta}(\mathbf{X})$$

- Behold to the **discrete** nature of counts
Negative Binomial

$$Y_{tj} | \theta, \nu \sim \text{NegBin}(\mu_{\theta}(t), \nu)$$

- Behold to the **discrete** nature of counts

Negative Binomial

$$Y_t | \theta, \nu \sim \text{NegBin}(\mu_\theta(t), \nu)$$

- Given $\mu_\theta(\cdot)$ the Y_t 's is **stochastically independent** from Y_{t-1}^c :

$$Y_t \perp Y_{t-1}^c \quad \forall t < T$$

Cumulative counts **likelihood**:

$$f_{Y^c}(y_1^c, \dots, y_T^c | y_0; \theta) = \prod_{t=1}^T f_{Y_t^c}(y_t^c | y_{t-1}^c; \theta) = \prod_{t=1}^T f_{Y_t}(y_t | \theta)$$



Optimal parameters $\hat{\theta}$:

- **Log-Likelihood maximization**
- **Multi-start** routine based on **Fisher-scoring**
- Gradient $r(\cdot)$ and Hessian $H(\cdot)$ **computed analytically**

Optimal parameters $\hat{\theta}$:

- **Log-Likelihood maximization**
- **Multi-start** routine based on **Fisher-scoring**
- Gradient $r(\cdot)$ and Hessian $H(\cdot)$ **computed analytically**

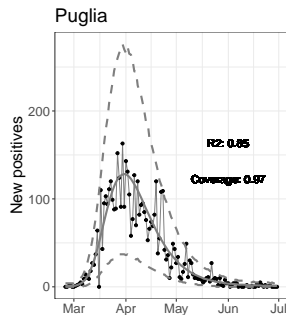
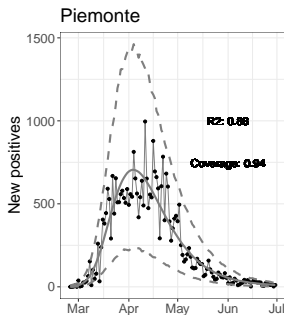
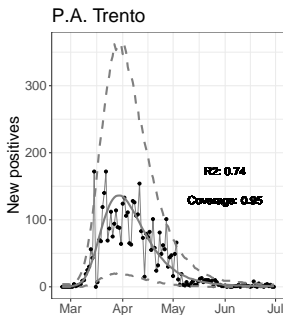
Uncertainty:

- Parameters' uncertainty quantified by the **Hessian**
- **Huber Sandwich** correction

$$V[\hat{\theta}] = \left(H(\hat{\theta}) \right)^{-1} r(\hat{\theta}) r(\hat{\theta})' \left(H(\hat{\theta}) \right)^{-1}$$

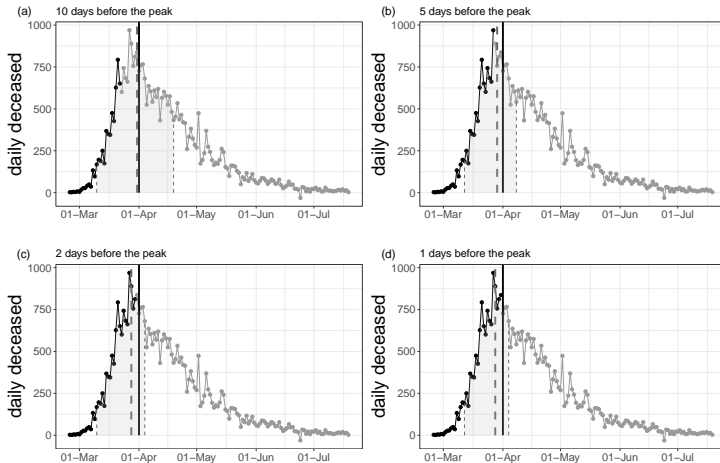
- Mean and prediction **intervals** obtained by bootstrap.

Model results - Fitting



Model fitting - Daily positives - Negative Binomial.

Model validation - peak detection



Estimation of the date of the peak for *daily deceased* at different steps-before

Introducing dependence in Mingione et al. (2021)

Disease mapping

$$Y_{gt} | \mu_{gt} \sim \text{Pois}(\mu_{gt})$$

$$\log(\mu_{gt}) = \log(E_g) + \log(m_{gt}), \quad g = 1, \dots, G, \quad t = 1, \dots, T$$

- E_g is an offset accounting for area-specific exposures levels
- m_{gt} is a relative measure of the risk of area g at time t

Disease mapping

$$Y_{gt} | \mu_{gt} \sim \text{Pois}(\mu_{gt})$$

$$\log(\mu_{gt}) = \log(E_g) + \log(m_{gt}), \quad g = 1, \dots, G, \quad t = 1, \dots, T$$

- E_g is an offset accounting for area-specific exposures levels
- m_{gt} is a relative measure of the risk of area g at time t

The relative risk

$$\log(m_{gt}) = \log\left(\tilde{\lambda}_{\gamma_g}(t)\right) + \mathbf{x}_{gt}^{\top} \boldsymbol{\beta} + \phi_{gt}$$

- $\tilde{\lambda}_{\gamma_g}(t)$ logistic growth temporal trend
- $\mathbf{x}_{gt}^{\top} \boldsymbol{\beta}$ a linear predictor based on K covariates
- ϕ_{gt} is a random effect for the g -th area at time t

Spatial dependence (Stern and Cressie, 1999)

- Neighborhood graph W s.t. $w_{ii} = 0$, $w_{ij} > 0$ iff $i \sim j$
- **CAR proper prior**

$$\phi_t \sim N_G \left(\mathbf{0}, \sigma^2 (D - \alpha W)^{-1} \right)$$

Spatial dependence (Stern and Cressie, 1999)

- Neighborhood graph W s.t. $w_{ii} = 0$, $w_{ij} > 0$ iff $i \sim j$
- **CAR proper prior**

$$\phi_t \sim N_G \left(\mathbf{0}, \sigma^2 (D - \alpha W)^{-1} \right)$$

Temporal dependence (Rushworth et al., 2014)

- 1-st order dependence on the past
- **AR(1)** over the space-vectors $f_t, g_{t=1}^T$

$$\phi_t | \phi_{1:t-1} \sim N_G \left(\rho \phi_{t-1}, \sigma^2 (D - \alpha W)^{-1} \right)$$



Common factors

- $\log(E_g) = \log(\text{residents}_g)$ scaled by a factor of 10^5
- Number of total weekly swabs (standardised) as covariate

Common factors

- $\log(E_g) = \log(\text{residents}_g)$ scaled by a factor of 10^5
- Number of total weekly swabs (standardised) as covariate

“Global VS Individual” growth

- One **common Richards** for all regions $\lambda_\gamma(\)$
- **Individual Richards** for each region $\lambda_{\gamma_g}(\)$

Common factors

- $\log(E_g) = \log(\text{residents}_g)$ scaled by a factor of 10^5
- Number of total weekly swabs (standardised) as covariate

“Global VS Individual” growth

- One **common Richards** for all regions $\lambda_\gamma(\)$
- **Individual Richards** for each region $\lambda_{\gamma_g}(\)$

Alternative spatial dependence graphs W

- $W_{Ind} = \mathbf{0}$! spatial independence
- W_{Flow} based on proximity flows (direct HV trains, flights, ferries) as in Della Rossa et al. (2020)
- W_{Geo} based on regions' geographical position

Prior setting

$$\begin{array}{ll} \log(b), \log(r) & N(\mathbf{0}, 100) \\ \log(h), \log(s) & N(0, 1) \\ p & N(T/2, T/(2 \cdot 1.96)) \\ \beta & N_K(\mathbf{0}, 100 \mathbf{I}_K) \end{array}$$

Prior setting

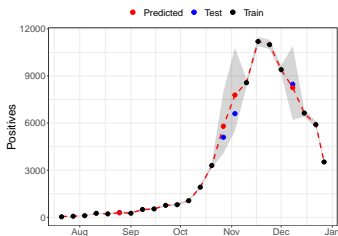
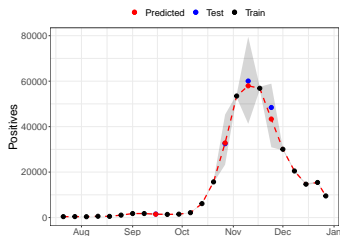
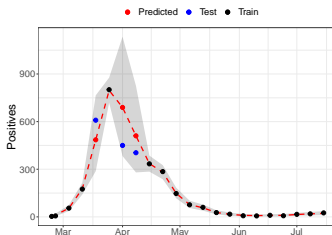
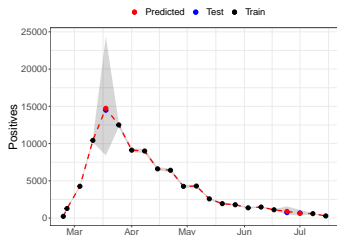
$$\begin{array}{llll} \log(b), \log(r) & N(0, 100) & \log(h), \log(s) & N(0, 1) \\ p & N(T/2, T/(2 \cdot 1.96)) & \beta & N_K(\mathbf{0}, 100 \mathbf{I}_K) \end{array}$$

Implementation using STAN (Carpenter et al., 2017)

- Hamiltonian Monte Carlo / **NUTS** (Hoffman and Gelman, 2014)
- **Exact-sparse** CAR (Joseph, 2016) for computational efficiency
- 70%/30% **in-sample/out-of-sample** split

Codes at <https://github.com/minmar94/Covid19-Spatial>

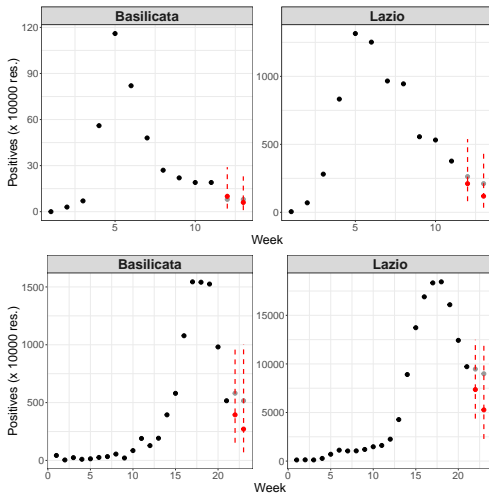
Out-of-sample predictions



(a) Lombardia

(b) Sicilia

Forecasting ability of the best model



Black dots in-sample, grey dots out-of-sample, red dots predicted values, red dashed 95% prediction intervals.

- **Data sources:** GitHub repository of *Italian Protezione Civile*
<https://github.com/pcm-dpc/COVID-19>
- **Detail:** collected and updated daily by the regional Health Systems
- **Type (main):**
 - **Prevalence:** current positives, intensive care occupancy (stock)

$$Y_t = Y_{t-1} + I_t - O_t$$

- **Incidence:** cumulative positives, cumulative deceased (flow)

$$Y_t^C = Y_{t-1}^C + Y_t$$

Focus on **ICU occupancy**

- $Y_{gt} \leq f_0, \dots, B_g$ **occupied beds** at time t in area g , with **capacity** B_g
- **Key for planning and allocating** health resources

Provide **1 to 5 days-ahead** predictions

- Use only data from the most **recent two weeks**
- Optimal **ensemble** of two methods
- Validate **short-term** performances on the run

First model

- **Generalized Linear Mixed Effect Model (GLMM)**
- Fit through `glmer` in the `lme4` package (Bates et al., 2007)

GLMM

$$Y_{gt} \sim \text{Poisson}(\lambda_{gt})$$

where

$$\log(\lambda_{gt}/R_g) = \beta_{g0} + \beta_{g1}t + \beta_{g2}t^2$$

and

$$(\beta_{g0}, \beta_{g1}, \beta_{g2}) \sim \text{MVN}((\beta_0, \beta_1, \beta_2), \Sigma)$$

First model

- **Generalized Linear Mixed Effect Model (GLMM)**
- Fit through `glmer` in the `lme4` package (Bates et al., 2007)

Second model

- **INGARCH** (Agosto et al., 2016; Chen and Lee, 2016)
- Fit through `tsglm` in the `tscount` package (Liboschik et al., 2015)

INGARCH

$$Y_{gt} \sim \text{Pois}(\mu_{gt})$$

$$\log(\mu_{gt}) = \alpha_{0g} + \log(\mu_{g,t-1}) + \alpha_{1g} \log(Y_{g,t-1} + 1) + \eta_{\beta}(t)$$

where

$$\eta_{\beta}(t) = \beta_{0g} + \beta_{1g}t + \beta_{2g}t^2 + \beta_{3g}t^3$$

First model

- **INGARCH** (Agosto et al., 2016; Chen and Lee, 2016)
- Fit through `tsglm` in the `tscount` package (Liboschik et al., 2015)

Second model

- **Generalized Linear Mixed Effect Model**
- Fit through `glmer` in the `lme4` package (Bates et al., 2007)

Model averaging

- **Convex combination** of predictions
- **Leave-last-out** optimal weight

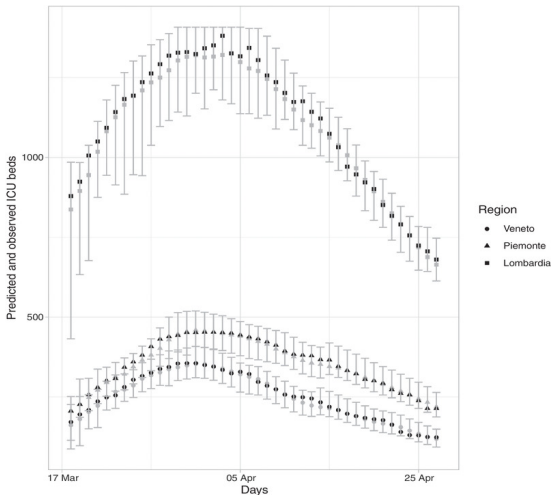
Ensemble predictor

$$\hat{Y}_{g,t+1} = w_{gt} \hat{Y}_{g,t+1}^{(INGARCH)} + (1 - w_{gt}) \hat{Y}_{g,t+1}^{(GLMM)}$$

where

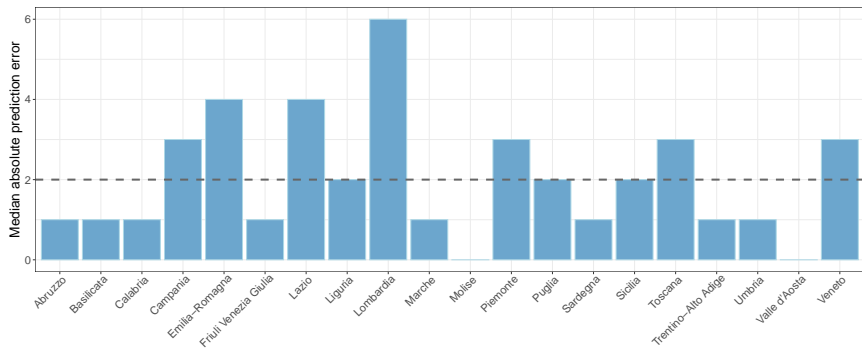
- $w_{gt} \in (0, 1)$ and $w_{gt} = 0.5$ for $t < 15$
- **Prediction intervals** are obtained as the **weighted average** of the limits of prediction intervals
- **Jensen's inequality** show that this conservatively guarantees the **nominal level**

Results (Farcomeni et al., 2021)



Predicted (grey) vs observed (black) number of ICU beds during the first outbreak in Lombardia, Veneto and Piemonte. Grey solid lines are 99% confidence intervals.

Results



Median absolute prediction error by region for ICU occupancy since March 12, 2020.

Concluding remarks

Conclusion

- ✗ Development of a coherent framework for the growth dynamic of counts
- ✗ Inclusion of desirable space-time dependence in the residuals
- ✗ Development of a reliable ensemble predictor of ICU occupancy

Conclusion

- ✗ Development of a coherent framework for the growth dynamic of counts
- ✗ Inclusion of desirable space-time dependence in the residuals
- ✗ Development of a reliable ensemble predictor of ICU occupancy

Work in progress

Growth model for prevalence indicators using INAR perspective

Include space-time dependence in the ICU ensemble model

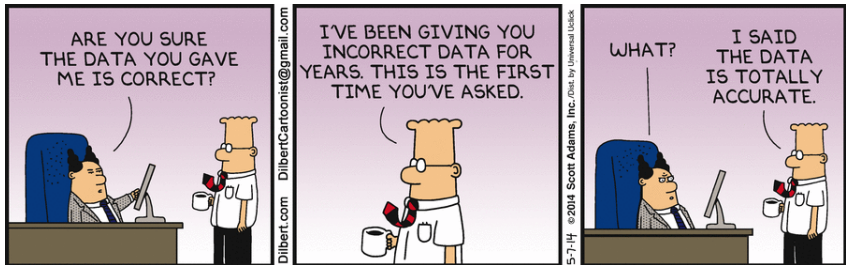
Include the effects of external policies in the model

Main references

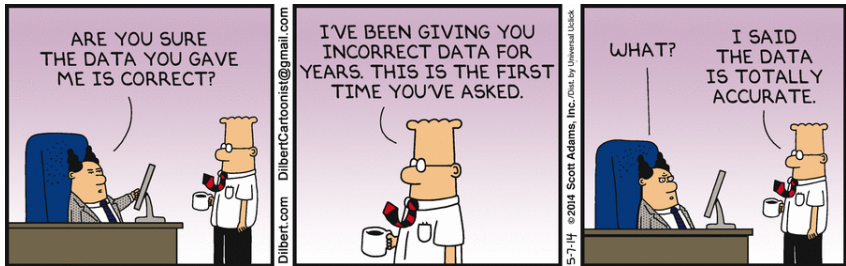
- A. Agosto, G. Cavaliere, D. Kristensen, and A. Rahbek. Modeling corporate defaults: Poisson autoregressions with exogenous covariates (parx). *Journal of Empirical Finance*, 38:640–663, 2016.
- P. Alaimo Di Loro, F. Divino, A. Farcomeni, G. Jona Lasinio, G. Lovison, A. Maruotti, and M. Mingione. Nowcasting covid-19 incidence indicators during the italian first outbreak. *Statistics in Medicine*, 40(16):3843–3864, 2021.
- D. Bates, D. Sarkar, M. D. Bates, and L. Matrix. The lme4 package. *R package version*, 2(1):74, 2007.
- B. Carpenter, A. Gelman, M. D. Hoffman, D. Lee, B. Goodrich, M. Betancourt, M. A. Brubaker, J. Guo, P. Li, and A. Riddell. Stan: a probabilistic programming language. *Grantee Submission*, 76(1):1–32, 2017.
- C. W. Chen and S. Lee. Generalized poisson autoregressive models for time series of counts. *Computational Statistics & Data Analysis*, 99:51–67, 2016.
- G. Chowell, D. Hincapie-Palacio, J. Ospina, B. Pell, A. Tariq, S. Dahal, S. Moghadas, A. Smirnova, L. Simonsen, and C. Viboud. Using phenomenological models to characterize transmissibility and forecast patterns and final burden of zika epidemics. *PLoS currents*, 8, 2016.
- F. Della Rossa, D. Salzano, A. Di Meglio, F. De Lellis, M. Coraggio, C. Calabrese, A. Guarino, R. Cardona-Rivera, P. De Lellis, D. Liuzza, et al. A network model of Italy shows that intermittent regional strategies can alleviate the COVID-19 epidemic. *Nature communications*, 11(1):1–9, 2020.
- O. Diekmann, H. Heesterbeek, and T. Britton. *Mathematical tools for understanding infectious disease dynamics*, volume 7. Princeton University Press, 2012.
- A. Farcomeni, A. Maruotti, F. Divino, G. Jona-Lasinio, and G. Lovison. An ensemble approach to short-term forecast of COVID-19 intensive care occupancy in italian regions. *Biometrical Journal*, 63:503–513, 2021.
- M. D. Hoffman and A. Gelman. The No-U-Turn sampler: adaptively setting path lengths in Hamiltonian Monte Carlo. *J. Mach. Learn. Res.*, 15(1):1593–1623, 2014.
- J. P. Ioannidis, S. Cripps, and M. A. Tanner. Forecasting for COVID-19 has failed. *International Journal of Forecasting*, 2020. ISSN 0169-2070. doi: <https://doi.org/10.1016/j.ijforecast.2020.08.004>. URL <http://www.sciencedirect.com/science/article/pii/S0169207020301199>.
- M. Joseph. Exact sparse CAR models in Stan, 2016. URL <http://mc-stan.org/users/documentation/case-studies/mjoseph-CARStan.html>, 2016.
- T. Liboschik, K. Fokianos, and R. Fried. *tscout: An R package for analysis of count time series following generalized linear models*. Universitätsbibliothek Dortmund Dortmund, Germany, 2015.
- M. Mingione, P. Alaimo Di Loro, A. Farcomeni, F. Divino, G. Lovison, G. J. Lasinio, and A. Maruotti. Spatial modelling of covid-19 incident cases using richards' curve: an application to the italian regions. *arXiv preprint arXiv:2106.05067*, 2021.
- A. Rushworth, D. Lee, and R. Mitchell. A spatio-temporal model for estimating the long-term effects of air pollution on respiratory hospital admissions in Greater London. *Spatial and spatio-temporal epidemiology*, 10:29–38, 2014.
- H. Stern and N. A. Cressie. Inference for extremes in disease mapping. 1999.



THANK YOU FOR YOUR ATTENTION!



THANK YOU FOR YOUR ATTENTION!



Questions?

APPENDIX

Poisson

$$\hat{d}_i^{\text{Poi}} = \text{sgn}(y_t - \mu_{\hat{\theta}}(t)) \sqrt{2y_t \log\left(\frac{y_t}{\mu_{\hat{\theta}}}\right) - (y_t - \mu_{\hat{\theta}}(t))}$$

Negative Binomial

$$\hat{d}_i^{\text{NB}} = \text{sgn}(y_t - \mu_{\hat{\theta}}(t)) \sqrt{2 \left[y_t \log\left(\frac{y_t}{\mu_{\hat{\theta}}(t)}\right) - (y_t + \nu) \log\left(\frac{y_t + \nu}{\mu_{\hat{\theta}}(t) + \nu}\right) \right]}$$



$$\hat{\phi}_t = \frac{y_t - \hat{y}_t}{\widehat{\text{Var}}[Y_t]}, \quad t = 1, \dots, T.$$

where:

$$\widehat{\text{Var}}_{\text{Poi}}[Y_t] = \mu_{\hat{\theta}}(t), \quad \widehat{\text{Var}}_{\text{NB}}[Y_t] = \mu_{\hat{\theta}}(t) + \frac{\mu_{\hat{\theta}}(t)^2}{\hat{\nu}}$$

Model validation

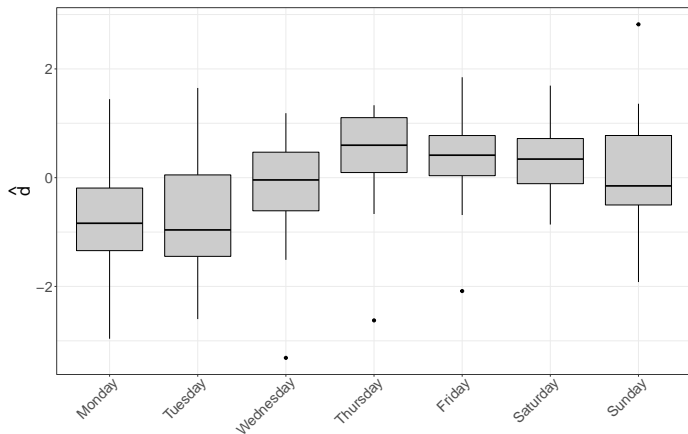


Figure: Deviance residuals distribution aggregated by day of the week for *daily positives*.

Model validation

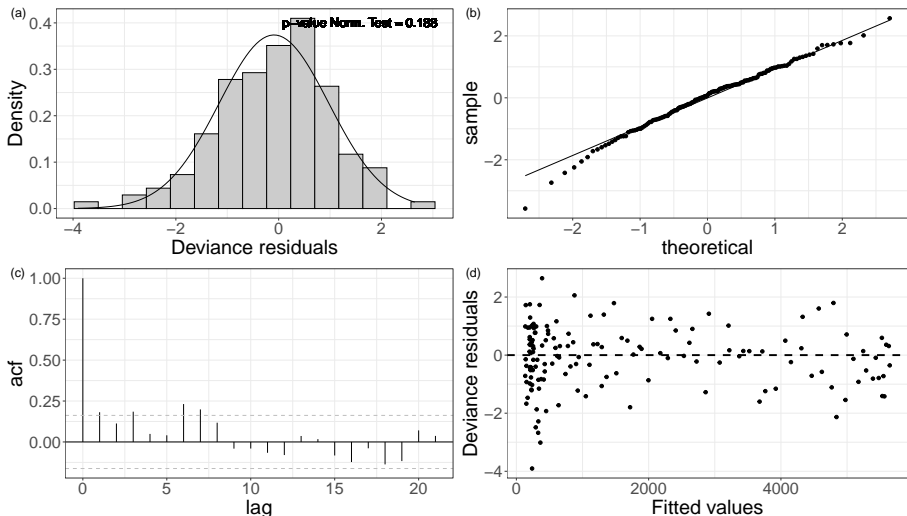


Figure: Residual diagnostic - Daily positives - Negative Binomial.

Model results

Table: Parameters' point estimates and 95% confidence intervals for the additive model on daily positives.

Parameter	Point estimate	95% Interval
β_0	5.26	(5.18, 5.34)
β_{wd}	0.46	(0.53, 0.38)
r	224.57 10^3	(224.13 10^3 , 225.01 10^3)
h	0.0289	(0.0287, 0.0291)
p	23.26	(29.64, 16.88)
s	44.42	(35.67, 124.51)
ν	22.01	(21.35, 22.70)

Parameters' estimates

Wave	Param.	M_{Ind}	M_{Flow}	M_{Geo}
I	α	–	0.14 (0.02, 0.21)	0.76 (0.71, 0.81)
	ρ	0.89 (0.87, 0.91)	0.88 (0.90, 0.93)	0.86 (0.85, 0.89)
	β	0.36 (0.26, 0.44)	0.34 (0.25, 0.42)	0.21 (0.14, 0.29)
II	α	–	0.93 (0.92, 0.95)	0.87 (0.85, 0.90)
	ρ	0.88 (0.86, 0.90)	0.87 (0.85, 0.89)	0.82 (0.80, 0.85)
	β	0.42 (0.38, 0.46)	0.27 (0.24, 0.30)	0.13 (0.09, 0.16)

Comparison of parameters' estimates for the spatial (α) and temporal (ρ) auto-correlation, and for the swabs' effect in the first and the second wave.

Model selection and validation

Wave	Metric	M_{Ind}	M_{Flow}	M_{Geo}
I	Coverage	0.98	0.98	0.98
	PIW	1535	1178	1144
	RMSE	423	184	272
	WAIC	2869	2650	2774
	LOO	3087	2849	2982
II	Coverage	0.96	0.97	0.92
	PIW	33393	4497	4046
	RMSE	12841	910	995
	WAIC	4112	3820	3971
	LOO	4393	4080	4252

Out-of-sample predictive performances for the first and the second wave.

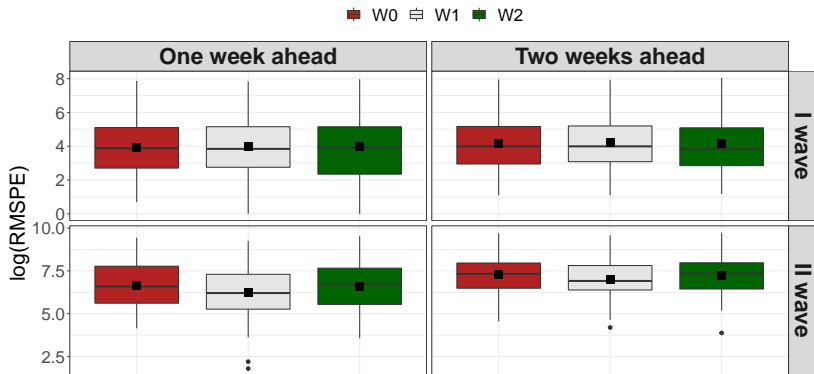
Parameters' estimates

Wave	Model	h	s
I	M_0	0.62 (0.60, 0.64)	7.8 (6.3, 9.9)
	M_1	0.62 (0.59, 0.65)	7.9 (5.5, 9.3)
	M_2	0.61 (0.58, 0.65)	7.8 (5.2, 9.3)
II	M_0	3.46 (3.26, 3.63)	0.06 (0.05, 0.07)
	M_1	2.72 (2.33, 3.08)	0.09 (0.07, 0.10)
	M_2	3.50 (3.20, 3.70)	0.06 (0.05, 0.07)

Wave	Model	b	r	p
I	M_0	0.05 (0.04, 0.06)	23 (20, 27)	2.0 (1.5, 2.5)
	M_1	0.06 (0.05, 0.07)	20 (17, 22)	2.2 (1.7, 2.8)
	M_2	0.05 (0.04, 0.06)	26 (21, 31)	2.2 (1.5, 2.9)
II	M_0	$7 \cdot 10^5 (1 \cdot 10^6, 1 \cdot 10^3)$	158 (143, 172)	23.2 (23.1, 23.3)
	M_1	$2 \cdot 10^4 (3 \cdot 10^5, 7 \cdot 10^3)$	178 (127, 215)	22.9 (22.8, 23.2)
	M_2	$4 \cdot 10^4 (3 \cdot 10^6, 1 \cdot 10^2)$	194 (163, 220)	23.1 (22.9, 23.2)

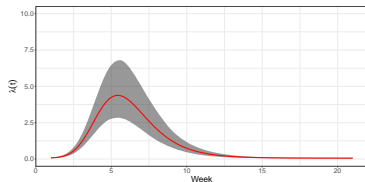
Table: Parameters' estimates of the Richards' curve for the waves I and II.

Forecasting comparisons between graphs

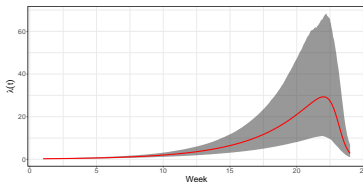


Prediction error (on the log scale) at different steps ahead, for each specification of W and for each wave.

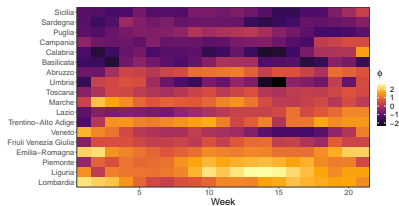
Common Richards' and latent effects



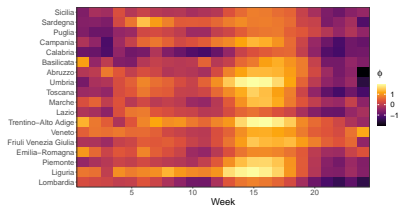
(a) M_{Flow} - I wave - $\tilde{\lambda}_\gamma(t)$



(b) M_{Flow} - II wave - $\tilde{\lambda}_\gamma(t)$



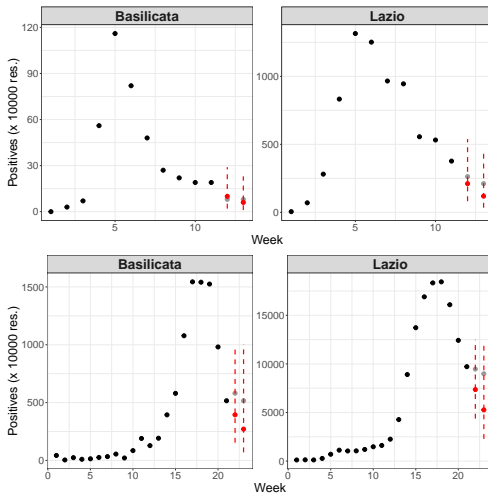
(c) M_{Flow} - I wave - ϕ_{gt}



(d) M_{Flow} - II wave - ϕ_{gt}



Forecasting ability of the best model



Black dots in-sample, grey dots out-of-sample, red dots predicted values, red dashed 95% prediction intervals.