

Complex systems

a dynamical point of view

Mauro Faccin

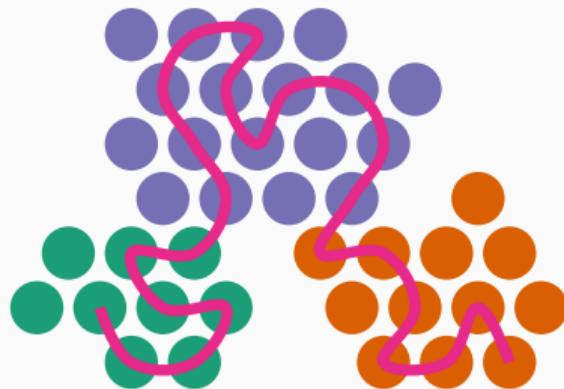
March 7, 2023

AutoInformation state aggregation

Projected Markov Chain

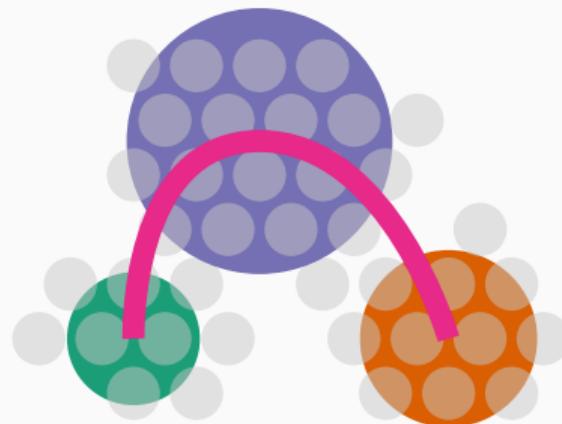
Markov Chain

$\dots, \mathbf{x}_{\text{past}}, \mathbf{x}_{\text{now}}, \mathbf{x}_{\text{future}}, \dots$



Projection

$\dots, \mathbf{y}_{\text{past}}, \mathbf{y}_{\text{now}}, \mathbf{y}_{\text{future}}, \dots$



Non-linear correlations

AutoInformation

$$I(y_t; y_{t-\tau})$$

Non-linear *correlation* between successive time-steps

$$I(y_t; y_{t-\tau}) = I(y_t; y_{t-\tau}, \dots) - I(y_t; y_{t-2\tau}, \dots | y_{t-\tau})$$

where τ represents a time-scale parameter.

We ask to:

- 1 maximize predictability of the dynamics;
- 2 minimize non-Markovianity (effective memories from the projection).

M.F. et al, Journal of Complex Networks, 2018

M.F. et al, PRL, 2021

Links to the literature

Modularity

χ_c characteristic function of class c

$$Q = \sum_c \text{Cov}(\chi_c(t), \chi_c(t+1))$$

Auto-covariance of the dynamics on the partition space.

Linear correlation between consecutive time-steps.

Shen et al. (2010) PRE, 82, 016114

DC-SBM

$$\mathcal{S} \propto \frac{1}{2} \sum_{cd} e_{cd} \log \frac{e_{cd}}{e_c e_d}$$

Fitting a generative model (e.g. DC-SBM) to the data through log-likelihood maximization can be seen as maximizing the AutoInformation for paths of lenght $\tau = 1$ (e.g. links).

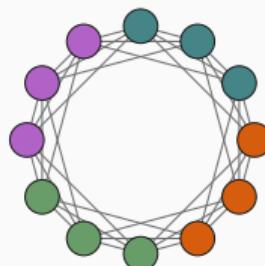
Karrer and Newman (2011), PRE 83, 016107.

Example: A cycle

A regular ring lattice with **N** nodes, each connected with **k** neighbours.

How many classes?

Adj:

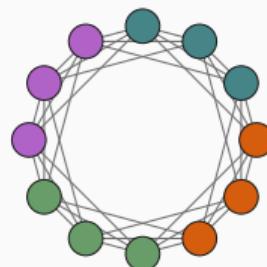
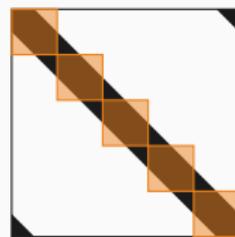


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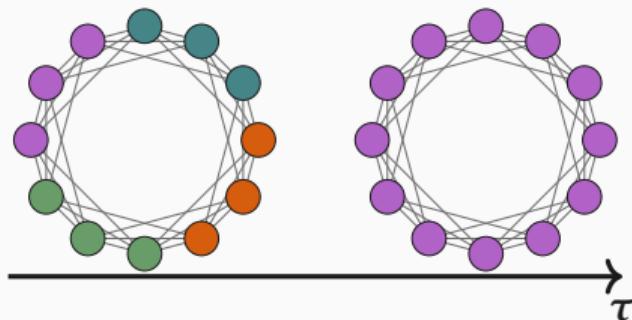


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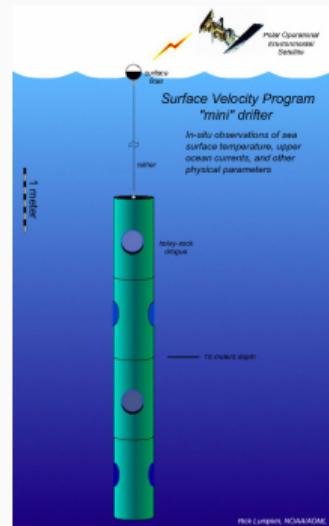
Adj:



Example: Ocean buoys

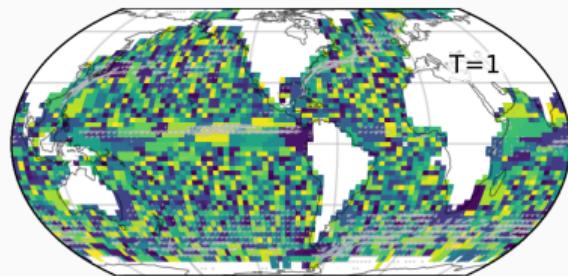


VOS Crew Deploy Next Generation SVP Drifter
Photo by: GDP



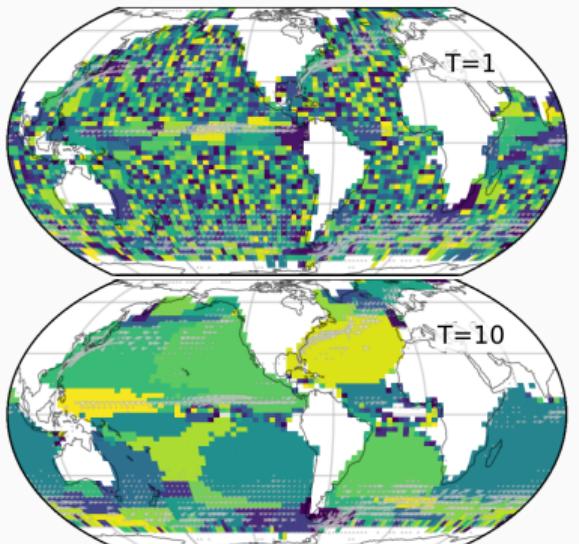
Global Drifter Program

Example: Ocean buoys

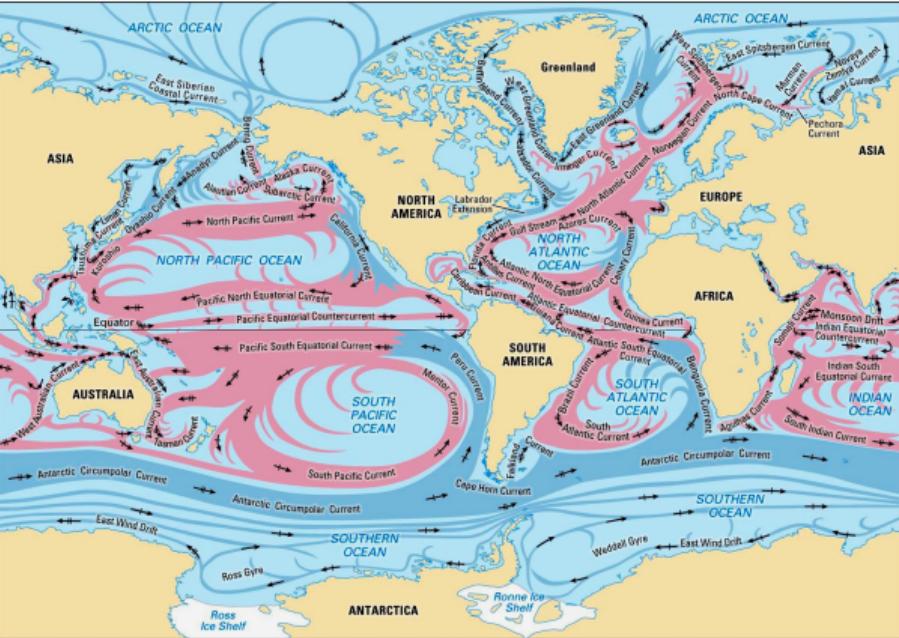


Each time step lasts 16 days.

Example: Ocean buoys



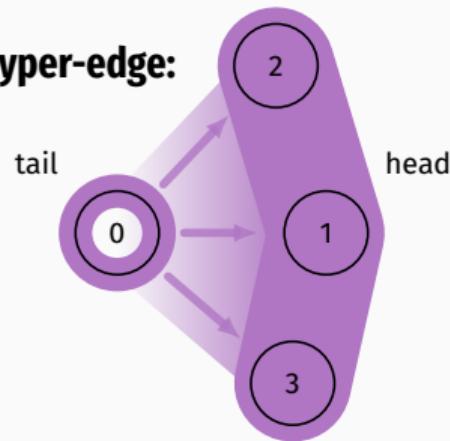
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Dynamics on hypergraphs

Hyper-edge:



node	role
0	tail
1, 2, 3	head

Hypergraph: $\{N, E\}$: nodes and hyperedges

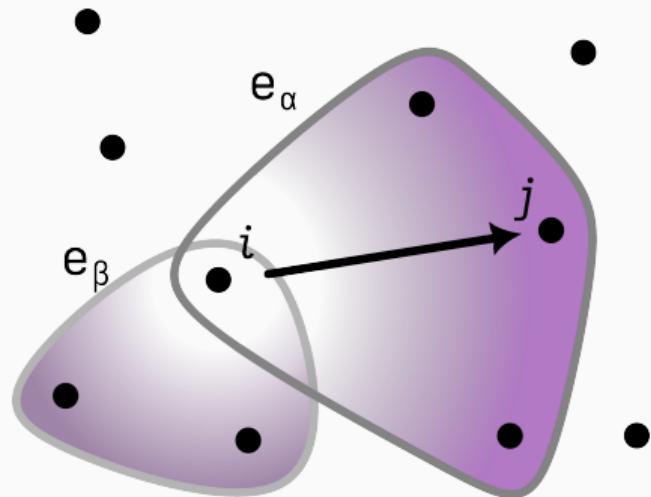
Nodes: same as before

Hyperedges: $e_\alpha = \{\text{tail}, \text{head}\} \in E$

Random walker on a hypergraph

The walker

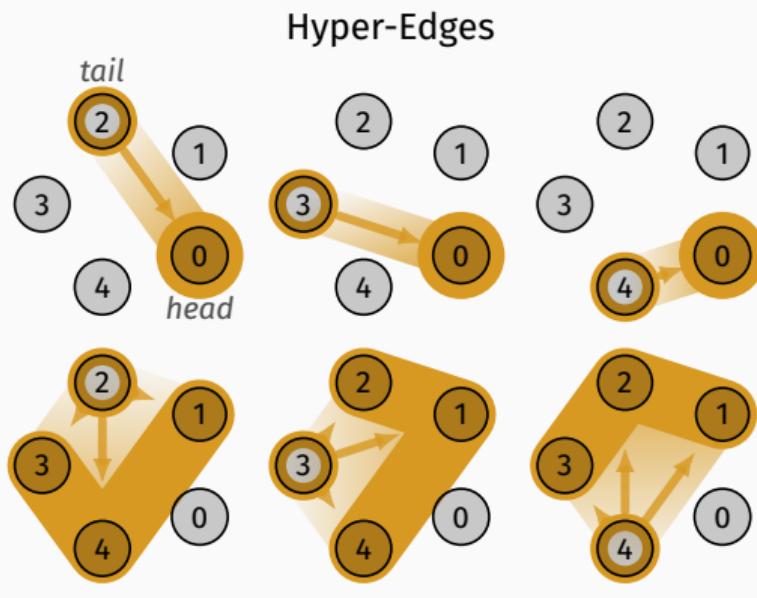
- sits on a node i
- chooses a hyperedge e_α incident on i in its tail (user i tweeted α) with probability dependant on the hyperedge size (with parameter τ);
- chooses an exit node j from the head of e_α (α get retweeted by user j) with flat probability.



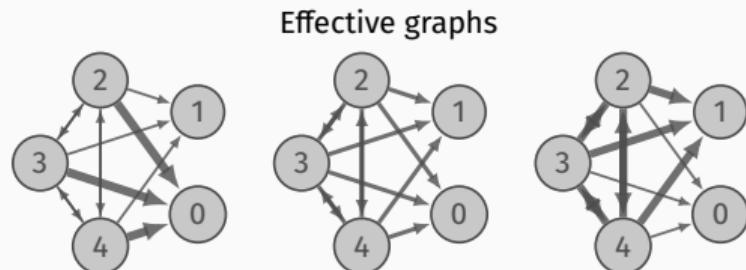
⌚ Yes but why?

Measure the dynamics through its transition matrix instead of extending to the hypergraph framework with the corresponding complexity overhead.

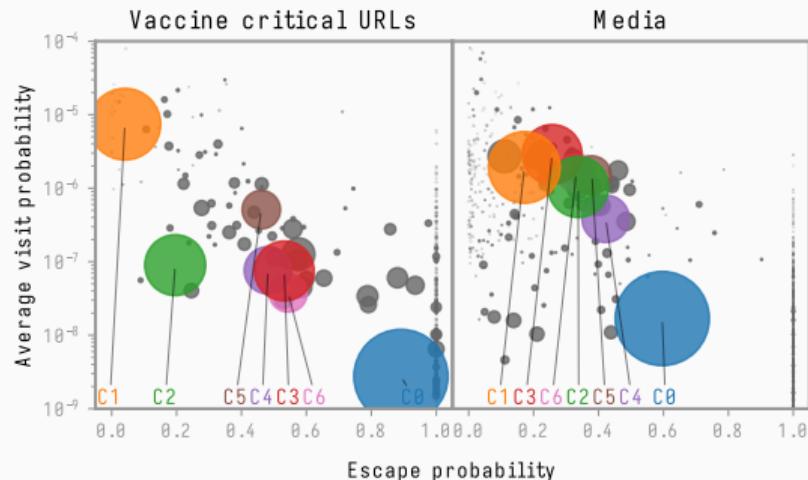
↓ Ranking nodes



$\tau = -1$		$\tau = 0$		$\tau = 1$	
ranking	prob	ranking	prob	ranking	prob
0	0.325	1	0.224	1	0.251
1	0.186	0	0.224	2	0.2
2	0.163	2	0.184	3	0.2
3	0.163	3	0.184	4	0.2
4	0.163	4	0.184	0	0.149



Communities and roles



Visiting probability probability of being visited by a random walker (a retweet)

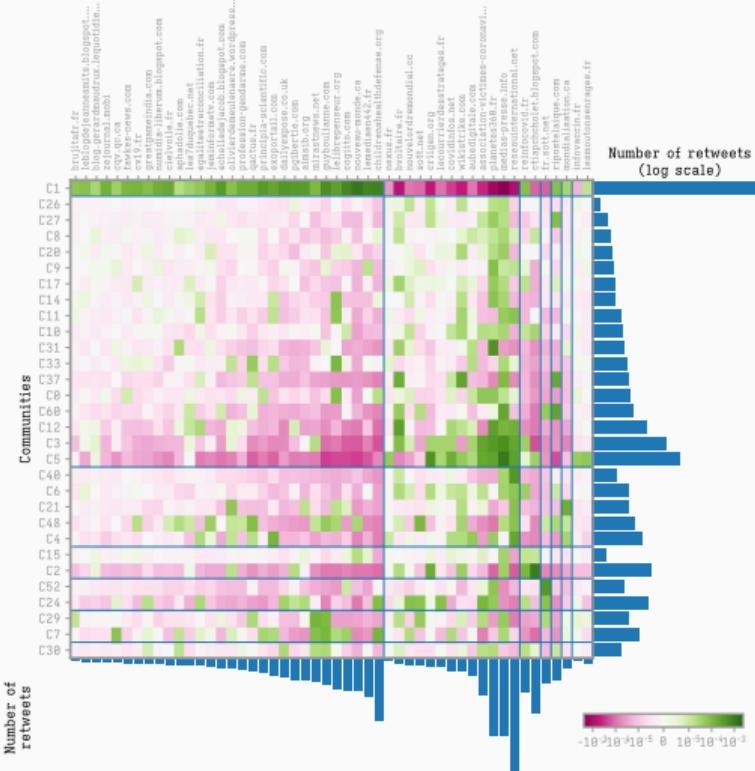
Escape probability probability of reaching other communities (being retweeted outside one's bubble)

Comm. Interpretation

- C₀ media aggregators or web influencers.
 - C₁ Far right groups.
 - C₂ health institutions and MDs
 - C₃ French news media.
 - C₄ international news media.
 - C₅ Far left and trade unions.
 - C₆ government representatives.
 - C₇ Canada
-

C₁ and C₅ are the main actors in spreading vaccine-critical content (high visiting and escape probability).

Community URL usage pattern.



Clustering of communities by URL usage pattern.

right wing C₁ use an original body of URLs.

left wing and news media use a similar set of sources.

health institutions use an original set of sources.

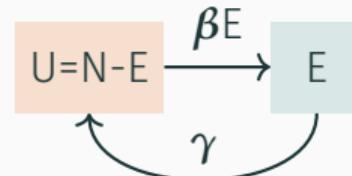
 **Compartmental models**

Engagement

Engaged users share a URL from the set within a time window $\tau = 3$ days.

$$dE_t = \alpha_t \frac{E_t(N_t - E_t)}{N_t} - \beta_t E_t$$

$$R_t = \frac{\alpha_t}{\beta_t}$$

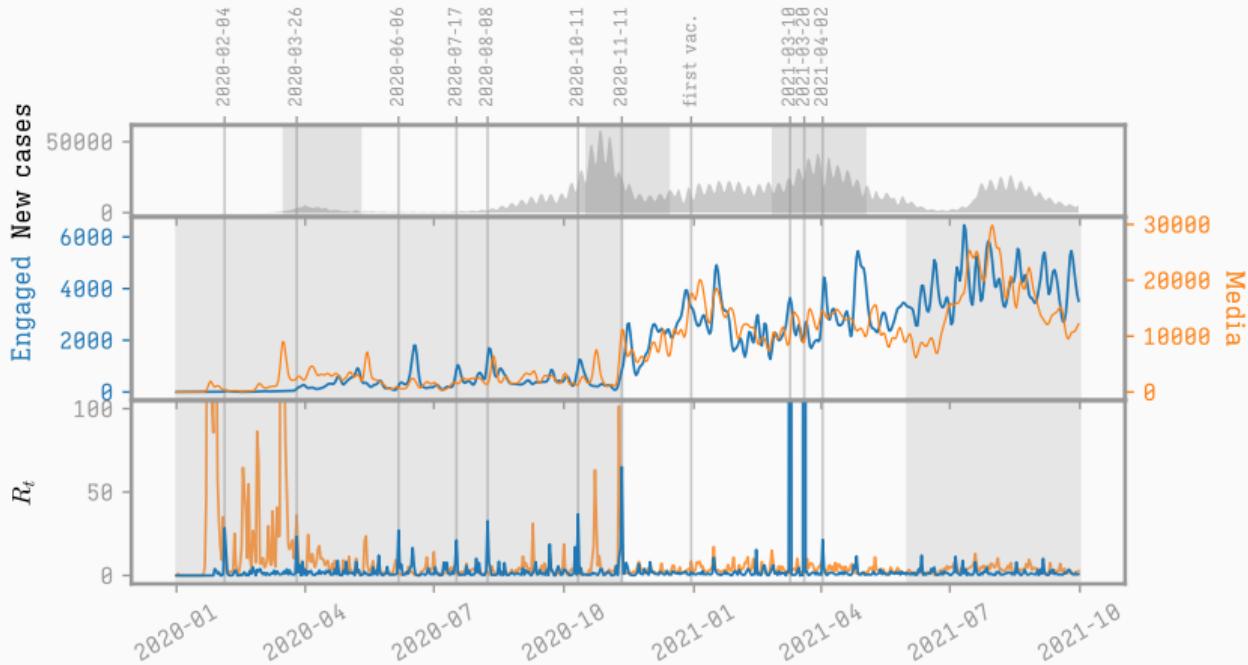


α_t engagement rate

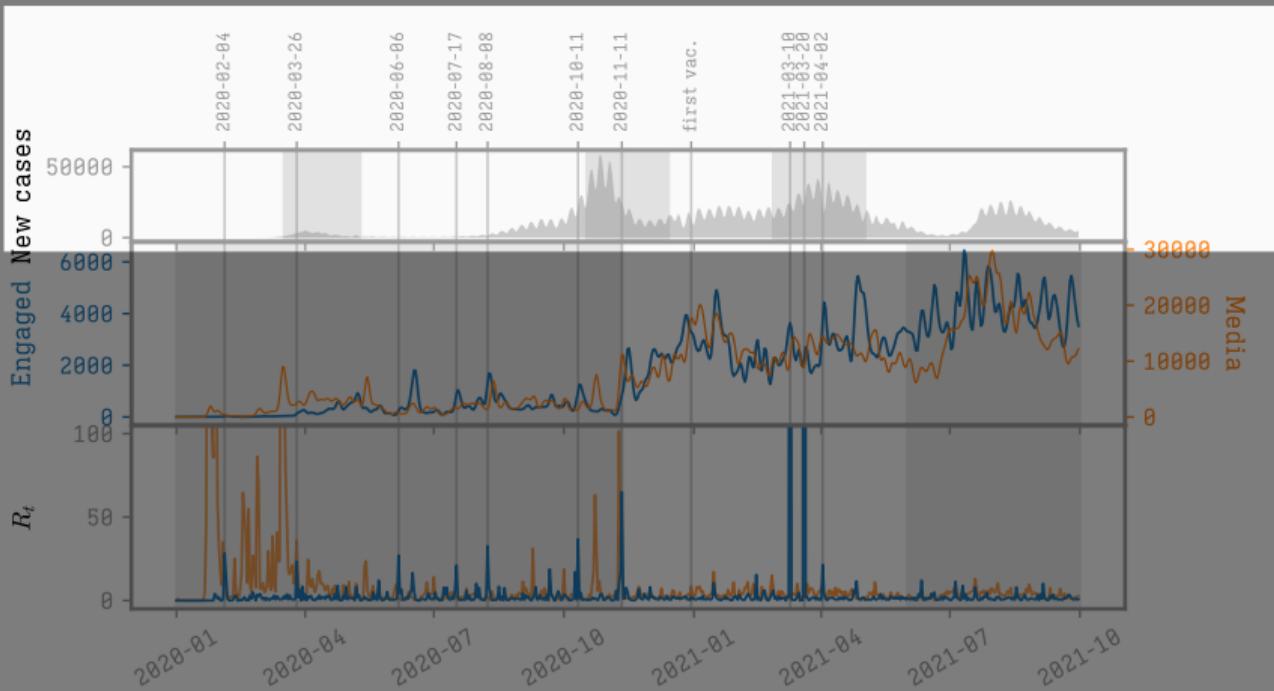
β_t unengagement rate

R_t reproduction number

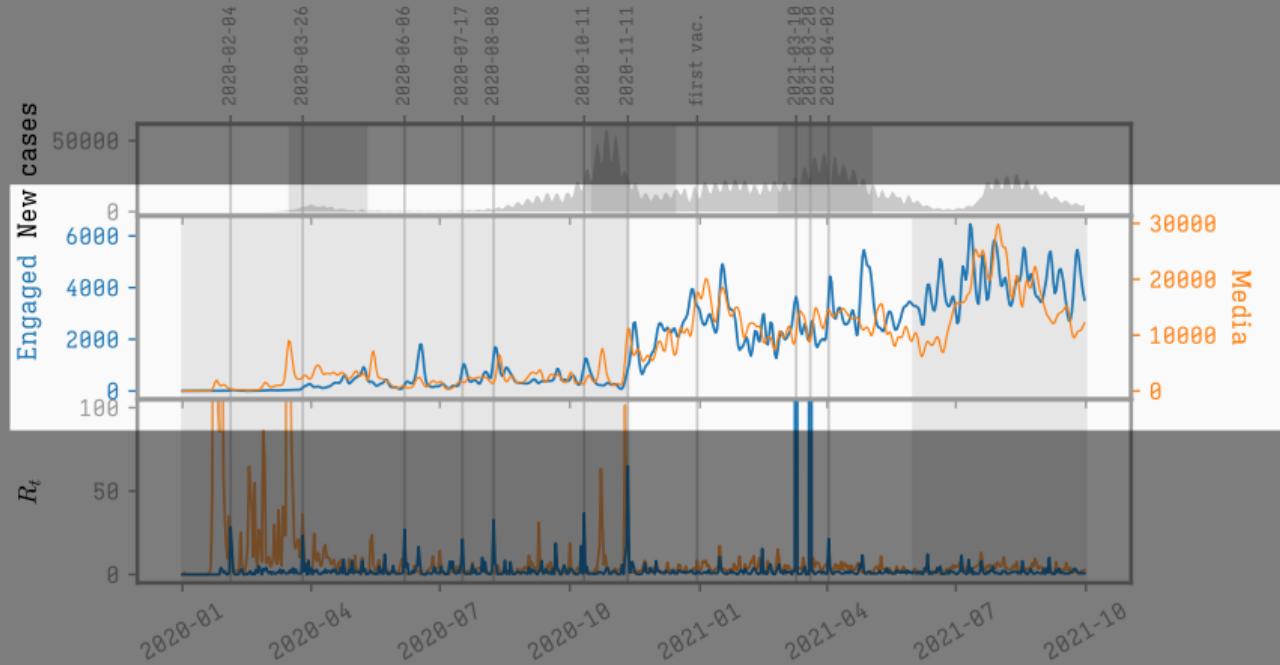
👤 + Engagement evolution



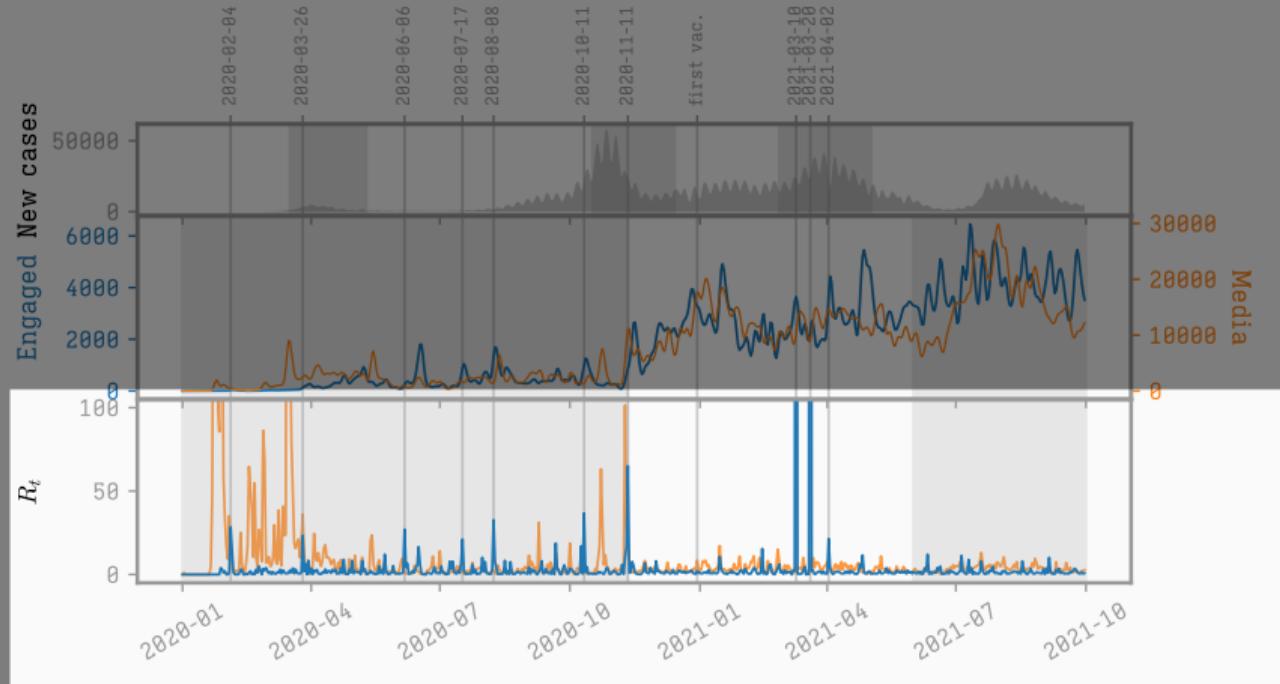
👤 Engagement evolution



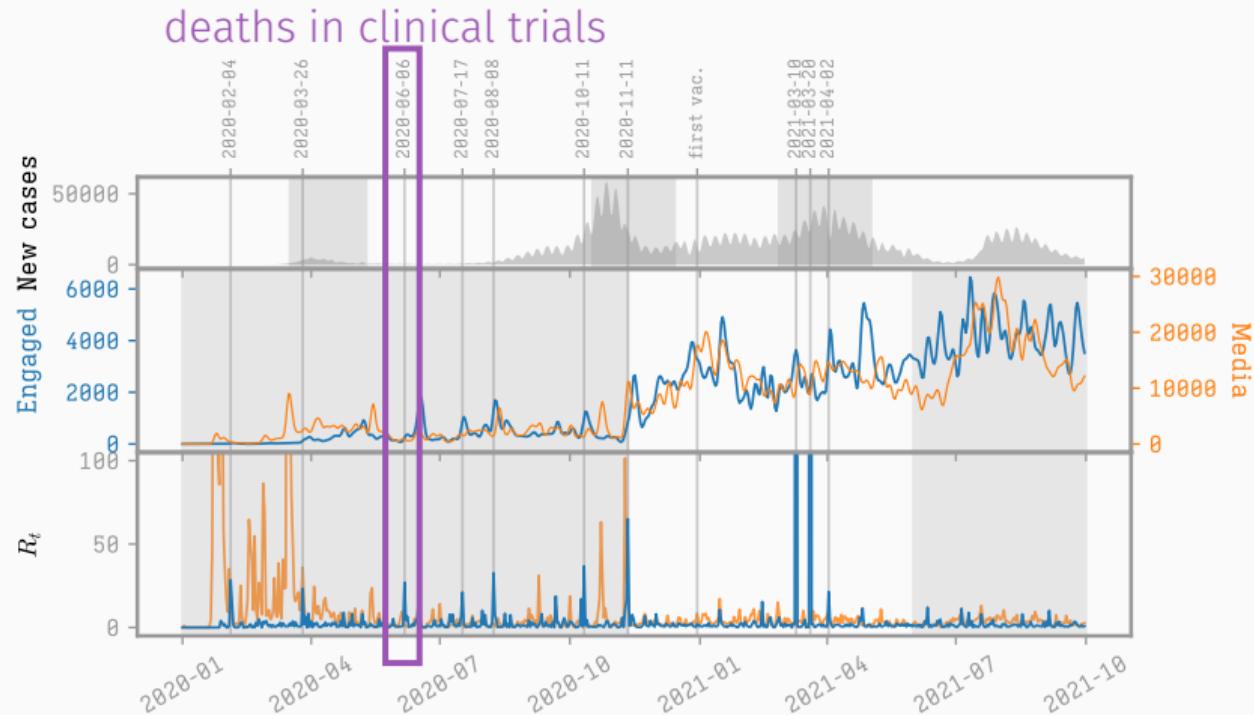
👤 Engagement evolution



👤 + Engagement evolution

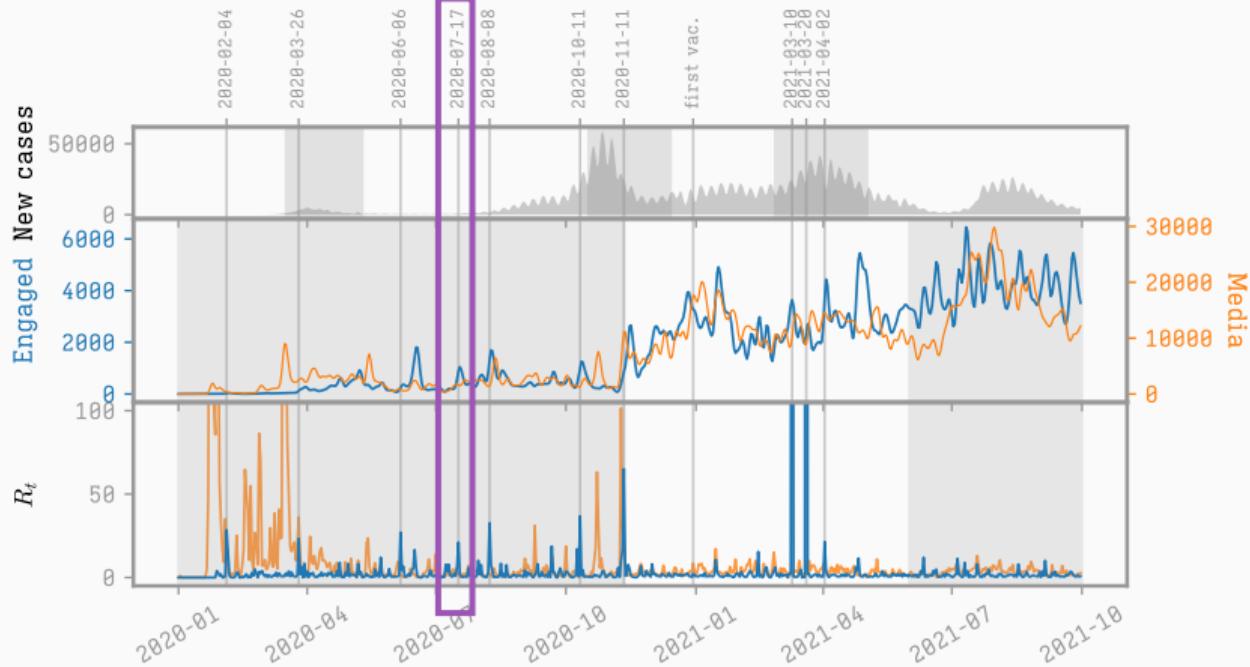


•+ Engagement evolution



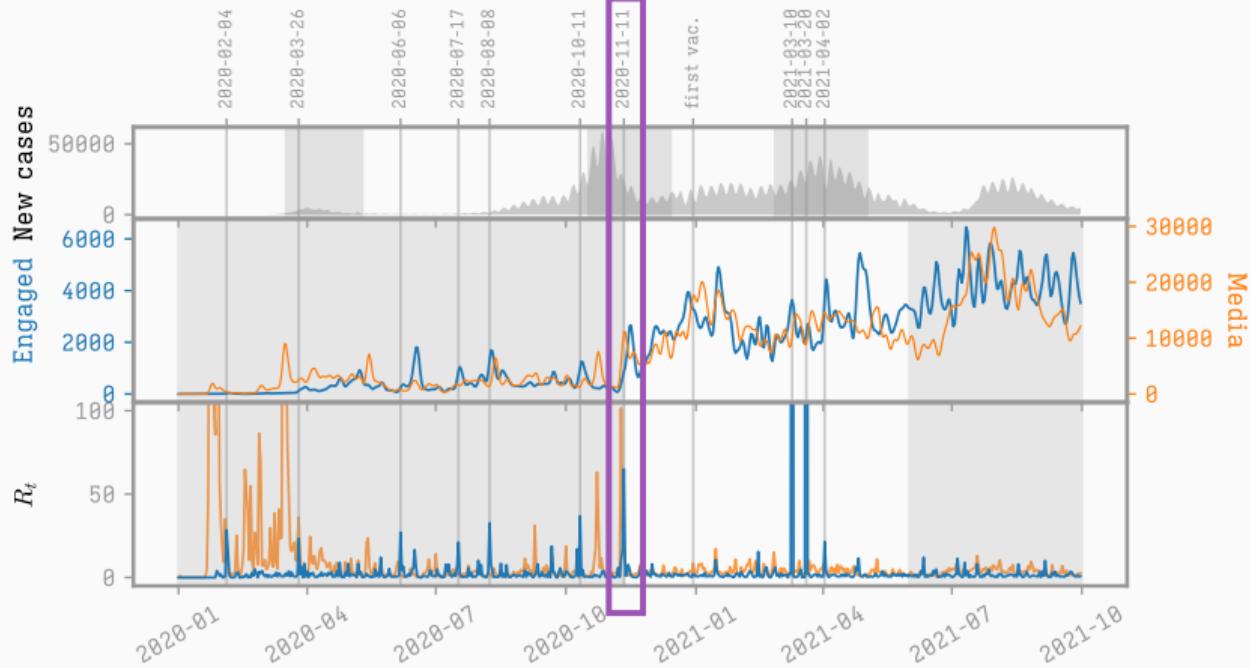
👤 + Engagement evolution

HC and remdesivir



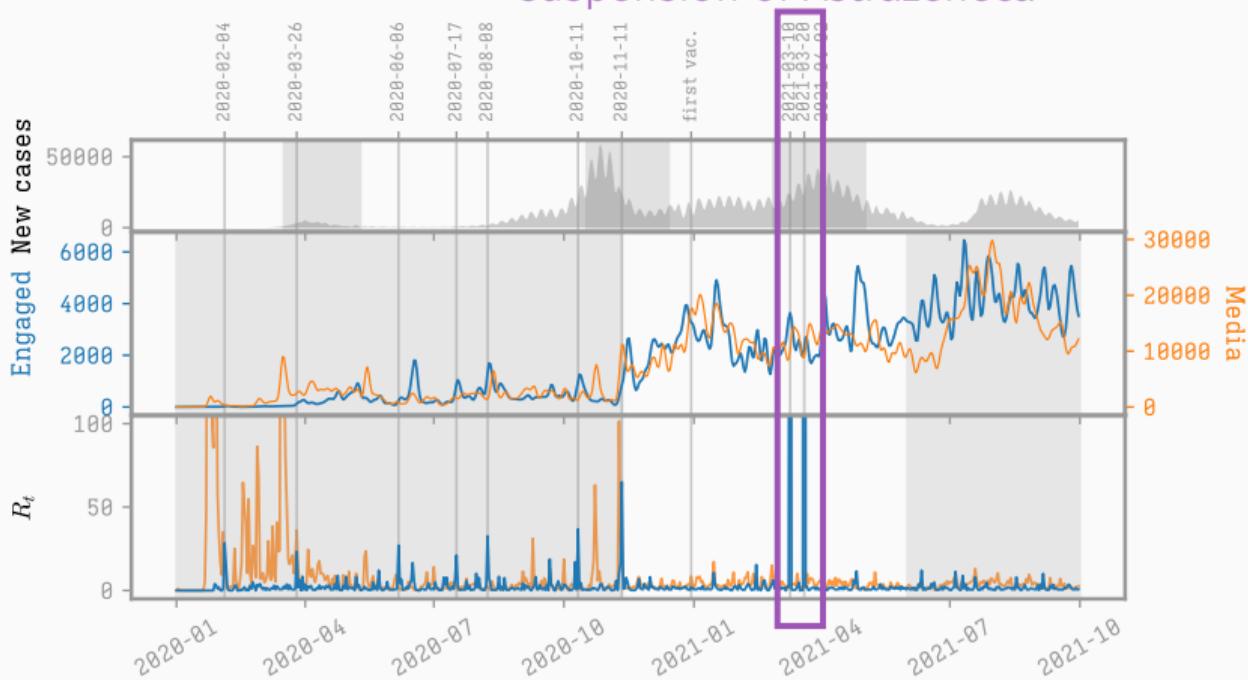
•+ Engagement evolution

Pfizer announcement / Holdup movie



•+ Engagement evolution

suspension of Astrazeneca



Tuberculosis under-detection

- 4 million of undetected/untreated TB cases, yearly [WHO]
- Hard to reach communities with high levels of TB incidence

Efficient approach to ACF

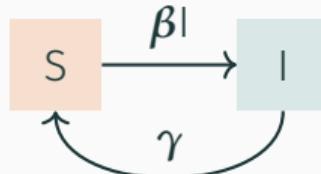
Data driven estimation of TB incidence rates to focus health-care interventions such as Active-Case Findings (ACF)

Active-Case Finding: systematic screening of the population for active TB cases.



The Model

Use of compartmental models (SIS) to disaggregate the reported cases.



Assumption

Endemic disease with slowly evolving **well mixed population**

$$\begin{cases} \frac{ds}{dt} = \gamma I - \beta \frac{Is}{S+I} \\ \frac{di}{dt} = \beta \frac{Is}{S+I} - \gamma I. \end{cases}$$

→ fit the parameters to:

- the number of cases reported by the **local health system** subunits;
- the population density as estimated by [Worldpop]

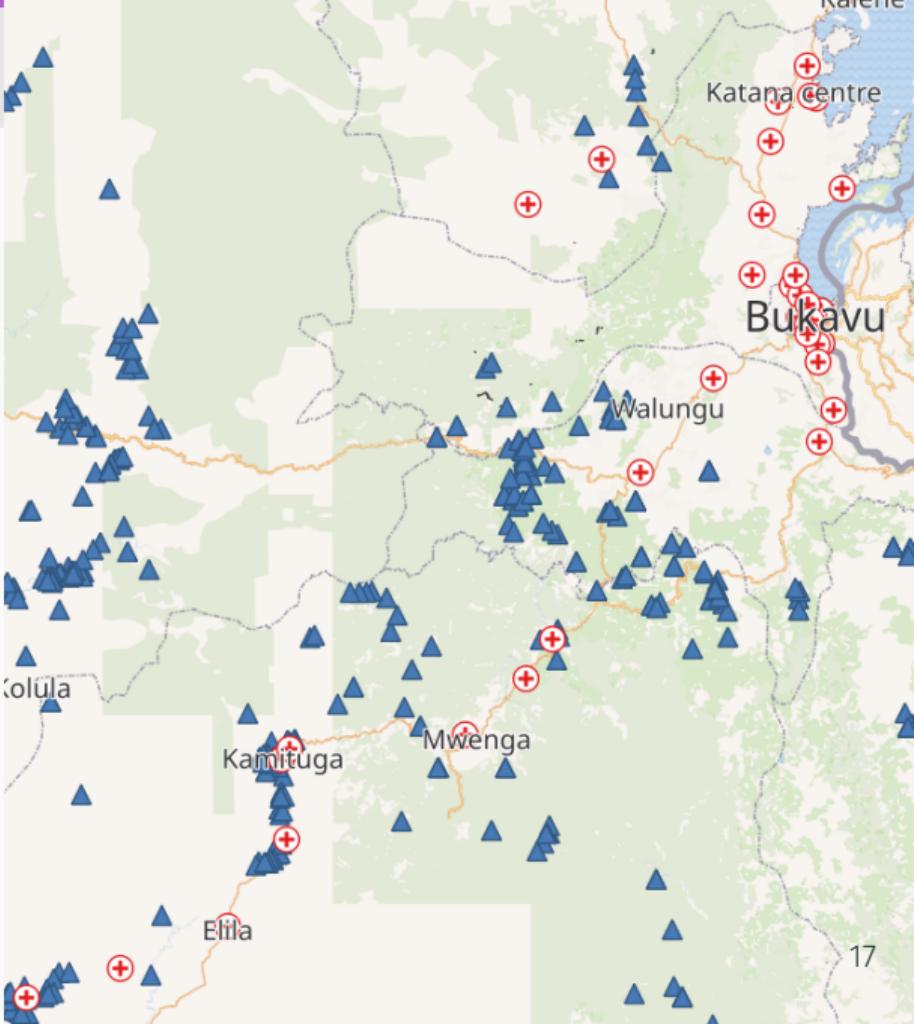
Model refinement

We include additional data to refine the estimation:

- Mine locations;
- Health facilities,

from of openly available sources:

- OpenStreetMap
- IPIS Research Project (mines)
- Healthsites

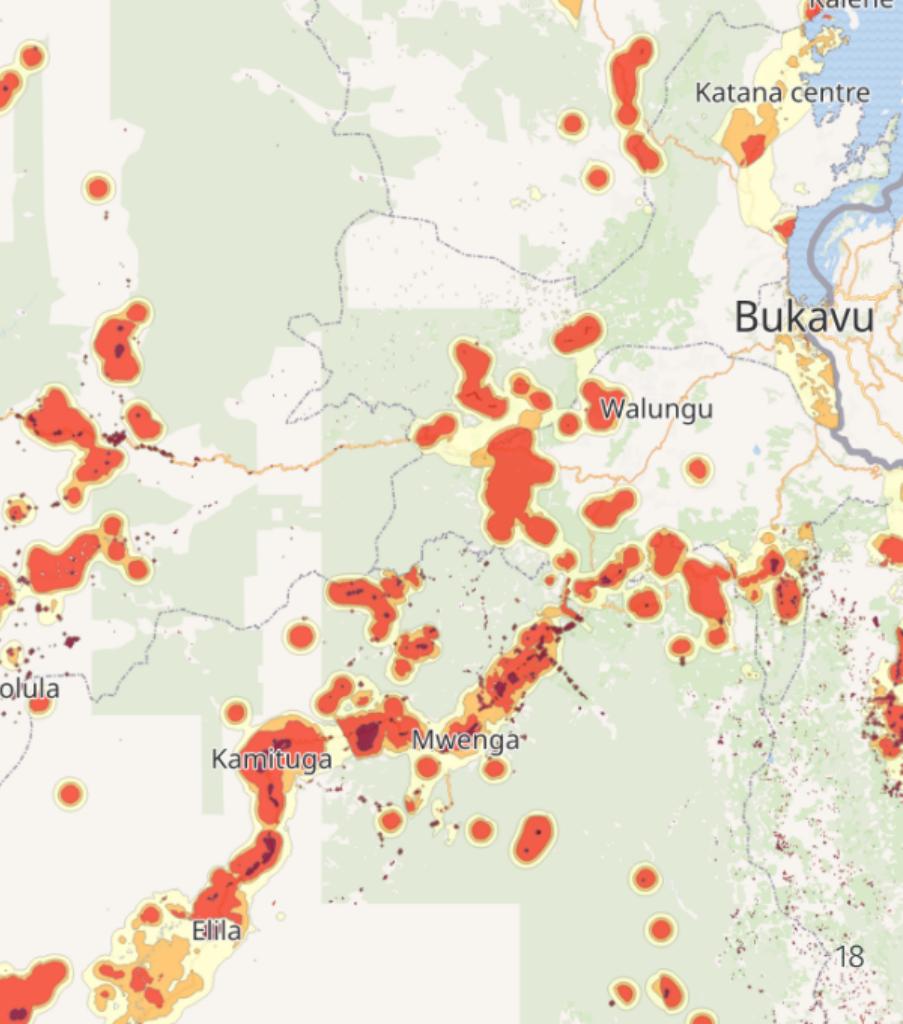


Incidence rate estimation

The estimated incidence rates highlight the location of population pockets with high risk of TB.

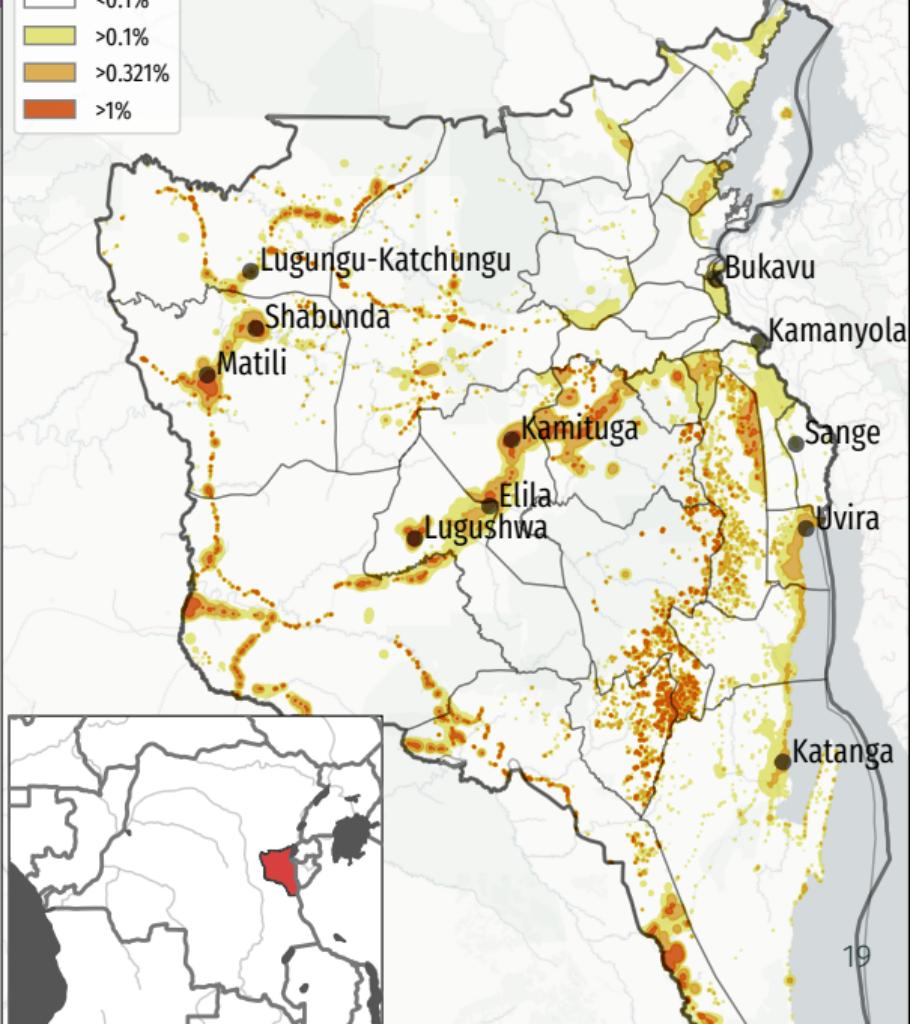
Leads to efficient ACF interventions.

color	incidence rate
yellow	>0.1%
orange	>0.32%
red	>0.5%
dark red	>1%



Multicentric clinical trial

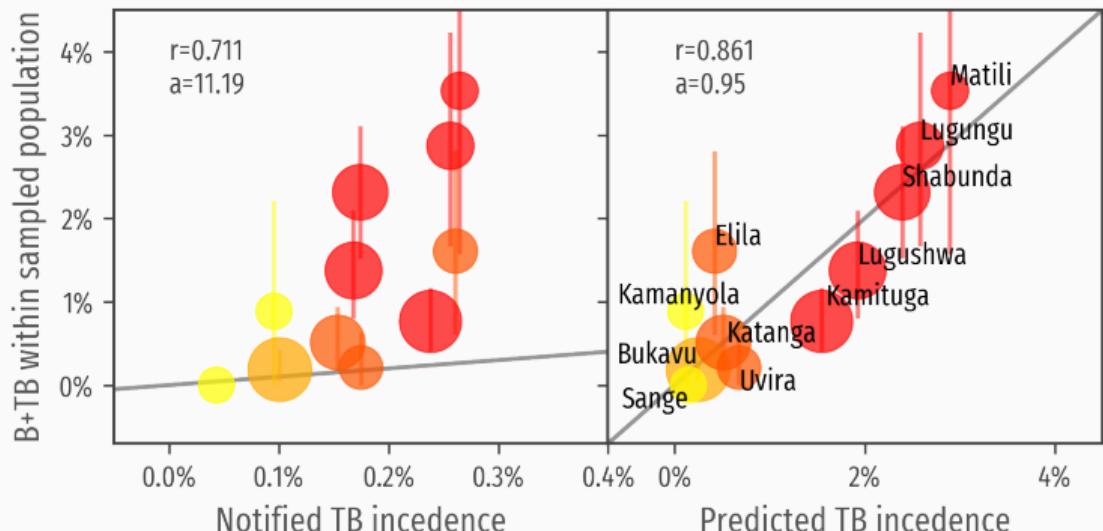
We performed a *multicentric clinical trial* addressing 11 locations with heterogeneous estimated incidence rates.



Results

screenings	13.841
lab tests	1153
positive cases	112

> 80% of positive cases originated from locations at high risk
(estimated incidence rate higher than 1%).



 Finally...

?

Questions?

Joint work with:



JC Delvenne



M Schaub



F Gargiulo



J Ward



E. André



👤 <https://maurofaccin.github.io>

✉ mauro.fccn@gmail.com

Code at:

👉 <https://maurofaccin.github.io/aisa>

👉 <https://maurofaccin.github.io/cartotb>