

## MODELLING OF EPIDEMICS ON GRAPHS.

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Why graphs? For the contact structure.

What is a graph? A set of nodes  $V$  and a set of edges  $E$  connecting various nodes. Each node  $v$  has a node degree: the number of edges connecting node  $v$  to other nodes in the graph. If  $u \rightarrow v_1 \rightarrow \dots \rightarrow v_k$  exists and is the shortest path between nodes  $u$  and  $v_k$ , then we say that the (minimal) distance between  $u$  and  $v_k$  is  $k$ .

Examples of graphs?

- Random graphs. Given  $N$  nodes, we assign edges between any two nodes at random.
- Lattice: a 2D grid.

How do we want our graph to be? What properties do “real graphs” possess?

- Geometry
- Sparsity.
- Clustering. The friend of my friend is likely my friend.
- Giant component. Everyone is connected to everyone.
- Node degree distribution. Node degrees are not all equal, they follow a power law distribution.  
 $p(k) = C \cdot k^{-(\tau+1)}$
- Small world phenomena. All  $k$ -th moments of node degrees are infinite, for  $k > \tau - 1$ . Average distances in the graph are small, less than  $\log(N)$ , where  $N$  is the graph size. If  $2 < \tau < 3$ , the variance of the node degree is infinite, and the average distance is less than  $\log(\log(N))$ . Such case is called ultra-small world. For social network that value is less than 10. That means you can “reach” any person in the world with less than 10 acquaintances of acquaintances.

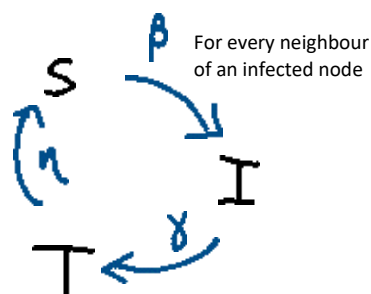
## THE MODEL

The epidemic is modelled as an ODE model for comparison and on three different types of graphs.

- A lattice (2D grid on a torus, so that there’s no boundary behaviour). It is geometric, but all nodes have the same degree.
- The configuration model (a random graph, which is constructed such that the nodes have a power law degree distribution). It has clusters, the small world (or ultra-small world) property, but it has no geometry.
- The Geometric inhomogeneous random graph (GIRG), a mixture between the previous two. It has all the properties we require. Construction: given  $N$  nodes, each node  $u$  is assigned a weight  $w_u$  according to a power-law distribution, and a uniformly random position  $x_u$ . Given nodes  $u, v$ , values  $w_u, x_u, w_v, x_v$  and parameter, the likelihood that an edge is assigned between  $u$  and  $v$  is:

$Prob(\text{edge between } u \text{ and } v) = \left( \frac{w_u \cdot w_v}{dist(x_u, x_v)^2} \right)^\alpha$ . Increasing  $\alpha$  we decrease the probability of having long edges.

$$\text{SITS model: } \begin{cases} s' = -\beta \cdot s \cdot i + \eta \cdot t \\ i' = \beta \cdot s \cdot i - \gamma \cdot i \\ t' = \gamma \cdot i - \eta \cdot t \end{cases}$$



On graphs, each node can be in one of three states: susceptible (S), infective (I) and temporary immune (T). Each (discrete) time step each infective node can infect each susceptible neighbour with probability  $\beta$ , it can itself heal with probability  $\gamma$ ; temporary immune nodes can lose their immunity with probability  $\eta$ .

As already seen, in the ode model, the disease either dies out if  $R_0 < 1$ , or it has an initial exponential increase, followed by damped oscillations until it reaches stability.

The graphs, having a stochastic nature, allow the disease to die out with non-zero probability also for  $R_0 > 1$ . In addition, all three graphs have an additional possible outcome: if  $\eta$  (loss-of-immunity parameter) is big enough there's one peak and then the disease dies out (SIR situation). There is a sharp transition of parameters between the three possible phases.

- Note: ODE, Configuration model & GIRG (with many hubs and long connections) have all initial exponential increase, when  $R_0 > 1$ . In the lattice & GIRG (without many hubs and long connections) it's linear.

## RESTRICTION MODELLING

- Travel restrictions. Decrease long-range connections by:
  1. Increasing parameter  $\alpha$ .
  2. For each edge between nodes  $u, v$ , set a cut-off value  $L$  and if  $L < d(x_u, x_v)$  then we delete the edge.
- Set a maximum number of contacts per individual.  $\rightarrow$  Set a maximal node degree  $M$  for all nodes  $v$  that have a higher node degree we randomly delete edges connected to  $v$  until its degree is within  $M$ .
- Keeping physical distance  $\rightarrow$  randomly remove edges from the graph.

## RESULTS

All interventions decrease the height of the first peak. On the other hand, they extend the time period of the peak. The critical  $\eta$  is shortened, so it is possible that intervention cause the disease to survive instead of dying out after the first peak. Travel restrictions (2) are the most effective in lowering the first peak.

## REFERENCES

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