Exercise 2 – Determining the spreading of epidemics

Deadline: April 23/2019. **Upload in the moodle**: 1. A presentation of the project in .pdf (no more than 6 pages). 2. The source code through which you generated the presented results

The files name should be surname_name_ex2.pdf and surname_name_code2.XX If the project has been developed by a group (max 3 persons), please indicate the name of the other authors in the presentation of the project (.pdf)

The Dataset:

- A) Files net1.txt and net2.txt contain two undirected networks representing two different mobility (i.e. contact) networks among individuals.
- B) File cases_100locations_first77days.txt describes the early stages of an outbreak of Chicken Pox in 100 different locations (see file's header for a description of the fields).

Tasks

- 1. Using the two networks provided in (A), calculate the epidemic threshold in the SIS dynamics for the homogenous, heterogenous and quenched mean-field approximations for mu=0.5.
- 2. Implement the (stochastic) SIS model in the two networks and simulate the entire phase diagram for mu=0.5. Compare the three theoretical thresholds with the numerical results. *Suggestion:* you need to simulate many (e.g. 100) simulations for each network and compare mean and std with the different analytical approximations.
- 3. Characterize the two networks in terms of their connectivity, degree distribution, page-rank distribution, clustering coefficient. Comment the results at the light of point 2 (e.g. how structural properties of the networks and results of the epidemic dynamics are related?)

- 4. (Optional). If you can segregate 1% of the individuals (i.e. remove 1% of the networks nodes), which node would you remove and why? What happen if you run again the SIS in network 1 and 2 with the nodes that you removed? Briefly discuss your results.
- 5. Plot the the early stages of infected for the 100 locations. Assuming that the disease spreading can be approximated by a SIR dynamics, use the method described in https://towardsdatascience.com/infection-modeling-part-1-87e74645568a to estimate the parameters of the SIR model. Will the chicken pox outbreack become viral?