## Theoretical frameworks for the SIRS model in complex networks with different localization patterns

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In the present work, we investigate the performance of theoretical approaches in the prediction of properties of the SIRS epidemic model, which involves immunity periods  $1/\alpha$  and whose acronym indicates the allowed states (*susceptible*, *infected* and *recovered*) as well as the transitions among them.

As a special case, we have the SIS model  $(1/\alpha = 0)$ , whose activation mechanisms in power-law degree distributed networks,  $P(k) \sim k^{-\gamma}$ , are well known. They involve self-sustained activity by means of feedback mechanisms in subextensive subgraphs [1], thus spreading to the rest of the network, with the epidemic threshold vanishing in the thermodynamic limit [2, 3]. The quenched mean-field theory [4] predicts this behavior qualitatively, with the threshold  $\lambda_c = \frac{1}{\Lambda_A}$  vanishing in the thermodynamic limit, where  $\Lambda_A$  is the Largest Eigenvalue (LEV) of the adjacency matrix  $A_{ij}$ . An enhancement of the QMF theory is the pair-quenched mean-field theory (PQMF), which takes into account the dynamic correlations at a pairwise level [5].

In the SIR limit  $(1/\alpha = \infty)$ , there is a transition to a state in which the fraction of recovered agents is finite, with a vanishing epidemic threshold in power-law distributed networks with  $\gamma \leq 3$  and a finite one when  $\gamma > 3$ . This behavior is well captured by the *heterogeneous mean-field theory* and the exact threshold is predicted by the *message passing* approach [6] on top of tree-like networks.

It is not clear which theoretical approach would be more suitable for the SIRS model with finite immunity periods  $1/\alpha$ , what is the main concern of our present work [7]. To investigate their properties, we developed PQMF equations for the SIRS model on networks and the rDMP equations reported in [8], whose predictions for threshold and Inverse Participation Ratio (IPR) [9, 10] were compared with extensive stochastic simulations. For more details, we refer the reader to [7].

Our results show that PQMF theory outperforms other approaches in networks without considerable localization effects [7]. Fig. 1 shows how localization affects mean-field theory predictions, by introducing a *hub* in a *random regular* network. PQMF theory undergoes strong localization on the hub and its neighborhood, while rDMP theory shows less localized behavior, agreeing qualitatively with simulation results, specially in dynamics with longer immunity times  $1/\alpha$ . Other results shown

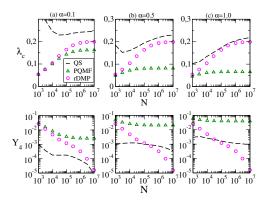


FIG. 1. Finite size scaling for SIRS model in random regular networks ( $k_{RR} = 6$ ) with hub ( $k_{hub} = 10^3$ ). Upper row: threshold  $\lambda_c$ . Lower row: IPR at the threshold. Immunity rates  $\alpha$  indicated at the column heading.

in our paper [7] present the same tendency of localization for the PQMF theory in the asymptotic limit, predicting a vanishing epidemic threshold, while simulations and rDMP theory predict delocalization and a finite threshold in power-law degree distributed networks with  $\gamma > 3$  [7].

**Keywords**: Epidemiology - Mathematical models. Networks (Mathematics). SIRS model. Immunity period. Mean-field theory.

## ACKNOWLEDGMENTS

The authors thank the financial support given by: Coordenação de Aperfeiçoamento de Pessoal de Nível Superior (CAPES), Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq), Fundação de Amparo à Pesquisa do Estado de São Paulo (FAPESP) and Fundação de Amparo à Pesquisa do Estado de Minas Gerais (FAPEMIG) - Brazil.

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