Message #3 : connection with structured differential equation models

- mass-action assumes the risk for individuals to become infected is uniformly distributed (i.e. in-degree in the network is distributed according to a Poisson distribution) and is independent of spread
- the final epidemic size can be mapped to a mixture of Poisson in-degree distribution

$$\dot{S}_i(t) = -eta \sigma_i \sum_j M_{ij} rac{I_j}{N} S_i \,,$$
 $\dot{I}_i(t) = eta \sigma_i \sum_j M_{ij} rac{I_j}{N} S_i - \gamma I_i \,,$ $\dot{R}_i(t) = \gamma I_i \,,$

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1.0-0.8-(S)epidemic size (Kermack-McKendrick 0.2-Age-structured SIR Poiss. mixt. 0.0 3.0 1.0 2.0 reproduction number (R_0)

One-sentence conclusion: Embracing networks is an important paradigm shift, but overlooking asymmetric interactions can hide important phenomena.



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