

1 Phase Transitions in Information Spreading on Structured  
2 Populations

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## Abstract

Mathematical models of social contagion that incorporate networks of human interactions have become increasingly popular, however, very few approaches have tackled the challenges of including complex and realistic properties of socio-technical systems. In this work we define a framework to characterize the dynamics of the Maki-Thompson rumor spreading model in structured populations, and analytically find a previously uncharacterized dynamical phase transition that separates the local and global contagion regimes. We validate our threshold prediction through extensive Monte Carlo simulations. Furthermore, we apply this framework in two real-world systems, the European commuting and transportation network and the Digital Bibliography and Library Project (DBLP) collaboration network. Our findings highlight the importance of the underlying population structure in understanding social contagion phenomena and have the potential to define new intervention strategies aimed at hindering or facilitating the diffusion of information in socio-technical systems.

The mathematical modeling of contagion processes is crucial in gaining insight into a broad range of phenomena from the spreading of infectious diseases to social collective behavior. While this avenue of research has a long tradition both in the biological and social sciences, in recent years there have been significant advancements triggered by increasing computational power and data availability characterizing socio-technical systems. These advances are particularly evident in the area of infectious disease forecasting where current models now incorporate realistic mobility and interaction data of human populations [1, 2, 3, 4, 5, 6]. Analogously, social contagion phenomena that were initially modeled using the same mathematical framework as epidemics [7, 8, 9, 10] are now described by *complex contagion* models [11, 12, 13] aimed at specifically characterizing processes such as the establishment of shared social norms and beliefs [14, 15, 16], the diffusion of knowledge and information [17, 18], and the emergence of political consensus [19]. These models consider complex factors such as reinforcement and threshold mechanisms [20, 21, 22, 23] and the loss of interest mediated by social interactions [24, 25, 26]. Furthermore, many of these theoretical approaches have put networks at the center of our understanding of social contagion phenomena and the information spreading process [27, 28, 8, 17, 29, 30, 31, 32, 12, 33]. However, most theoretical and numerical work on the dynamics of social contagion focuses on highly stylized models, trading off the realistic features of human interactions for analytical transparency and computational efficiency. As a result, social contagion models able to integrate the effects of human mobility, community structure, and time varying behavioral patterns are largely unexplored.

In this paper, we consider the classic rumor spreading model [24, 25] to study the effects of structured populations on the global diffusion of a rumor or piece of information. More specifically, we model the spatial structure of realistic populations and the behavior of individuals in virtual social networks through a reaction-diffusion model in a metapopulation network, and an activity-driven model with communities, respectively. We first identify analytically the necessary conditions for the social contagion to spread to a macroscopic fraction of the population. This analysis shows that although the rumor model is lacking any critical threshold, the population structure introduces a dynamical phase transition (global invasion threshold [34]) which is a function of the interactions between subpopulations. We validate the analytical results with large-scale numerical simulations on synthetic networks with different topological structures. Additionally, we recover the global threshold of the contagion process in data-driven models of the European transportation network and the Digital Bibliography and Library Project (DBLP) collaboration network.

Understanding how the social structure in both the physical and virtual worlds affects the emergence of contagion phenomena has the potential to indicate novel ways to utilize the network connectivity to develop efficient network-based interventions. The framework developed here opens a path to study the effects of communities and spatial structures in other complex contagion processes which can incorporate agent memory [35] and social reinforcement [22], or introduce other heterogeneous features such as age-dependent contact patterns and socio-economic conditions [36].

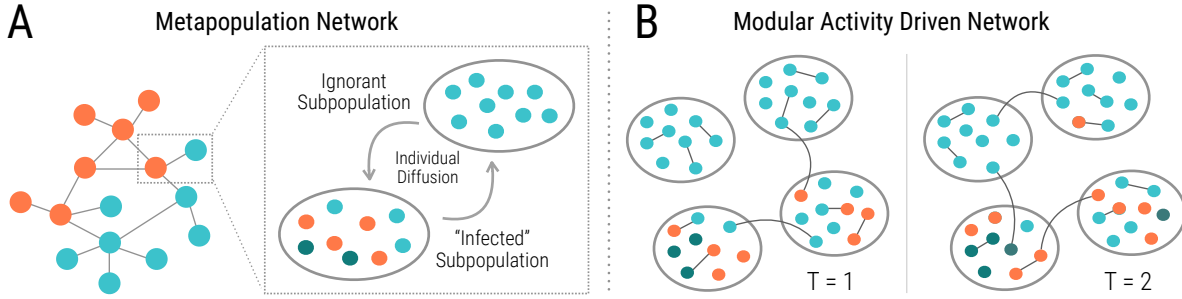


Figure 1: Types of structured populations considered in the modeling framework. **(A)** A schematic representation of a reaction-diffusion process on a metapopulation network, where individuals homogeneously interact within their current subpopulation, and then diffuse through the network constrained by the global structure. **(B)** A schematic representation of a modular activity-driven network at two points in time where individuals are confined to a single community, but when activated choose to form links to those outside of their current community based on a probability of inter-community interaction. Each instantaneous network is generated independently of prior networks.

## 1 Model Definition

Here we use a variant of the original rumor model [24], known as the Maki-Thompson (MT) model [25], to describe the spread of information through a population based on interactions between agents [37]. Similar to epidemic models, individuals can be classified into three compartments, *ignorants*: those who do not know the rumor, *spreaders*: those who know and are actively sharing the rumor, and *stiflers*: those who know the rumor but are no longer spreading it. The contagion process evolves through interactions between individuals in a population. If a spreader contacts an ignorant individual, with a probability  $\lambda$ , the ignorant will transition into a spreader. However, when a spreader contacts either a stifler or another spreader, with a probability  $\alpha$ , the spreader will transition into a stifler. The stifling mechanism describes an individual's tendency to become uninterested in the rumor once the appeared novelty of the information is lost. In homogeneously mixed populations, this feature does not allow the presence of a *rumor threshold* [24, 38], meaning that for any  $\lambda > 0$  the rumor will always spread to a macroscopic proportion of the population (see methods). We investigate the behavior of this model on two types of structured populations that incorporate the complexity observed in socio-technical systems (Fig. 1).

- Rumor model in spatially structured populations.** We first consider a population where spatially defined groups of individuals (subpopulations) are coupled together by a mobility rate (Fig. 1a) [39, 33]. This structure, also called a metapopulation network, is used to model species persistence in ecosystems [40], the evolution of populations [41], and the global spreading of infectious diseases [42]. Specifically, we consider a metapopulation network with  $V$  subpopulations, each with an average population size of  $\bar{N}$  individuals. Reaction-diffusion processes are used to characterize both the local interaction and global mobility dynamics. Individuals first *react* within their current subpopulation according to the rumor model dynamics and then *diffuse* between subpopulations based on a Markovian diffusion process. The probability that an individual will leave her current subpopulation and travel to a specific neighbor is  $p/k$ , where  $p$  is the mobility parameter and  $k$  is the number of neighboring subpopulations.
- Rumor model in virtual structured populations.** In contrast to the reaction-diffusion scheme, rather than moving between communities, individuals may belong to a specific virtual community such

84 as an interest or disciplinary group, online forum, or political affiliation etc., but interact occasionally  
85 with individuals in other virtual communities through collaborations, forum posts, or direct messages  
86 etc.(Fig. 1b). We model these interaction dynamics using a modular activity-driven network scheme  
87 [43, 44]. In particular, we consider a population with  $V$  communities whose sizes ( $s$ ) follow a specific  
88 distribution,  $P(s)$ . Every individual is assigned an activity potential  $a_i\Delta t$  that is sampled from a  
89 preset distribution  $F(a)$ . The activity of an individual corresponds to the probability with which the  
90 individual becomes active during a given time step [43]. Each active individual will form a single  
91 connection to another individual in the population creating an instantaneous network. To induce  
92 a community structure, an activated individual will choose to form a link to a randomly selected  
93 individual outside of her home community with a probability  $\mu$ , otherwise, an intra-community link  
94 will be formed. The parameter  $\mu$  allows us to tune the interaction between communities. After each  
95 single iteration of the rumor model, the network resets and a new instance is generated in the same  
96 manner.

## 97 2 Invasion threshold in structured populations

98 Although the rumor model in a single homogeneous population does not exhibit a spreading threshold, the  
99 presence of a subpopulation structure fundamentally alters the contagion dynamics. This can be clearly  
100 seen for the rumor model in virtual structured populations by examining two limits of the inter-community  
101 interaction term,  $\mu$ . When  $\mu = 0$ , individuals will only interact with others in their community. Thus, the  
102 rumor will never escape the seed community. However, in the limit where  $\mu = 1$ , individuals effectively  
103 do not belong to any community and will always choose to form an external connection. Therefore, the  
104 rumor will certainly reach a macroscopic fraction of the population. The same reasoning can be applied to  
105 the limits of the mobility parameter,  $p$ , in the case of spatially structured populations. In both modeling  
106 frameworks, the population structure induces a transition point separating a dynamical regime where only  
107 local spreading is possible from a regime where the rumor spreads globally through the network.

108 To characterize this transition point quantitatively, we use a branching process framework to describe the  
109 rumor spreading dynamics across subpopulations [42, 45]. Let us consider a system that is structured into  
110  $V$  subpopulations, each consisting of  $\bar{N}$  individuals, on average, at any given time. Within the homogeneous  
111 population structure, we assume that all nodes are statistically equivalent and the connections formed  
112 between pairs of nodes are uncorrelated. Let  $D_n$  be the number of affected subpopulations where the rumor  
113 is known by at least one individual at generation  $n$ . We use a tree-like approximation to write an expression  
114 that captures the number of subpopulations that know the rumor at each generation of the spreading process,  
115 obtaining:

$$116 \quad D_n = D_{n-1} \left( 1 - \frac{\sum_{m=0}^{n-1} D_m}{V} \right) C\Phi. \quad (1)$$

117 The above equation assumes that every affected subpopulation in the  $(n-1)^{th}$  generation ( $D_{n-1}$ ), may  
118 seed each one of its  $(1 - \sum_{m=0}^{n-1} D_m/V)C$  unaffected neighbors with a probability  $\Phi$ , where  $C$  indicates the  
119 average number of neighboring subpopulations and  $1 - \sum_{m=0}^{n-1} D_m/V$  is the probability that the neighboring  
120 subpopulation is not already aware of the rumor during the  $(n-1)^{th}$  generation. In a structured population  
121 model, a rumor epidemic occurs when each affected subpopulation, early in the contagion process, spreads  
122 the rumor on average to at least *one* fully ignorant subpopulation. Using the above expression, this global  
123 contagion condition reads as  $D_n/D_{n-1} \geq 1$ . Given that we are interested in the early time dynamics of  
124 the process, we assume that  $\sum_{m=0}^{n-1} D_m/V \ll 1$ , defining the *global contagion threshold*  $D_n/D_{n-1} \simeq C\Phi \geq$   
125 1. This effectively defines the *subpopulation reproductive number*  $R^* = C\Phi$ ; i.e. the average number of  
126 communities becoming aware of the rumor from a single subpopulation. Analogously to the reproductive  
127 number in biological epidemics, in order for information to spread globally,  $R^*$  must be greater than or equal  
128 to one [42, 45]. The terms,  $C$  and  $\Phi$ , depend explicitly on the type of structured population model as well as

129 the contagion process. In the following sections we provide expressions for these parameters and the rumor  
 130 invasion thresholds for both spatially structured and virtual populations.

## 131 2.1 Spatially structured populations

132 In a homogeneous metapopulation network the number of possible subpopulations that could be seeded by  
 133 each affected subpopulation is  $C = \langle k \rangle - 1$ ; i.e. the average number of neighboring subpopulations minus the  
 134 one which originally seeded the contagion process. Due to the lack of a local rumor threshold within a single  
 135 subpopulation, the probability  $\Phi$  is simply given by the probability that at least one spreader will decide  
 136 to leave an affected community and travel to a neighboring subpopulation where it will start spreading the  
 137 rumor:

$$138 \quad \Phi = 1 - \left(1 - \frac{p}{\langle k \rangle}\right)^\beta, \quad (2)$$

139 where  $\beta$  is the number of spreaders in an affected subpopulation that can travel out of their current sub-  
 140 population during the rumor epidemic. This value  $\Phi$ , is calculated by considering one minus the probability  
 141 that none of the spreaders will travel to a new community,  $(1 - \frac{p}{\langle k \rangle})^\beta$ . Here,  $\beta = \frac{2(1+\lambda/\alpha)\bar{N}}{\alpha}$ , is the product  
 142 of the total number of individual spreaders generated by the contagion process within a single population  
 143 and the average amount of time they are actively spreading the rumor (details in methods). Using the above  
 144 expressions and considering small mobility probabilities, such that  $p/\langle k \rangle \ll 1$ , we can approximate the  
 145 probability  $\Phi \simeq \beta p/\langle k \rangle$ . In this limit we obtain an explicit expression for the rumor invasion threshold as:

$$146 \quad R^* = \frac{\langle k \rangle - 1}{\langle k \rangle} 2 \left(1 + \frac{\lambda}{\alpha}\right) \frac{p\bar{N}}{\alpha} \geq 1. \quad (3)$$

147 From the above expression it is possible to rewrite the necessary threshold condition to find the critical  
 148 mobility  $p_c$  in the system required for a global spreading of the rumor as:

$$149 \quad p_c = \frac{\langle k \rangle}{(\langle k \rangle - 1)} \frac{\alpha}{2(1 + \frac{\lambda}{\alpha})\bar{N}} \quad (4)$$

150 Below the critical value,  $p_c$ , the amount of individual mobility restricts the global propagation of the ru-  
 151 mor. In this subcritical regime, spreaders in affected communities are generally unable to travel to a new  
 152 subpopulation before they transition into stiflers, which consequently causes the rumor to go extinct in the  
 153 early stages. This critical mobility is a function of both the network structure and the rumor model param-  
 154 eters,  $\lambda$  and  $\alpha$ . However, for homogeneous networks with sufficiently large average degrees  $\langle k \rangle$ , the effect  
 155 of the network structure is relatively insignificant. In the Supplementary Information we derive the critical  
 156 mobility for metapopulation networks with heavy tailed degree distributions and find that the analytical  
 157 expression depends not only the average degree of the network  $\langle k \rangle$ , but also the second moment  $\langle k^2 \rangle$  of  
 158 the degree distribution. Heterogeneous networks are characterized by having degree distributions with high  
 159 variance (large  $\langle k^2 \rangle$ ), thus considerably affecting the value of the mobility threshold. We also see that  $p_c$  is  
 160 linearly dependent on  $\alpha$ . When  $\lambda$  is small relative to  $\alpha$ , the critical mobility is controlled predominantly by  
 161 the stifling probability. Recall that the stifling probability characterizes the tendency for an individual to  
 162 become disinterested in the rumor (i.e. transition into a stifter) when interacting with others that know the  
 163 rumor. This finding is a feature worth remarking for the global spread of a rumor in a spatially structured  
 164 environment, that places the emphasis not on how *appealing* a rumor is, but rather on the rate at which  
 165 people decide that the rumor is not worth spreading.

## 166 2.2 Virtual structured populations

167 Now let us consider a modular activity-driven network where we assume discrete time,  $\Delta t = 1$ , a homogeneous  
 168 activity rate ( $a$ ) for all individuals in the network, and a homogenous distribution of community sizes. In  
 169 this model, if an individual chooses to form an inter-community link, by construction it can choose any of

170 the other  $C = V - 1$  communities. The probability  $\Phi$ , that at least one spreader from an affected population  
 171 will choose to connect another individual outside of their current community and successfully transmit the  
 172 rumor can be written as

$$173 \quad \Phi = 1 - \left(1 - \frac{\lambda\mu}{V-1}\right)^\beta. \quad (5)$$

174 In this expression  $\frac{\lambda\mu}{V-1}$  is the probability that an inter-community link successfully transmits the rumor to  
 175 one of the  $V - 1$  specific subpopulations and  $\beta$  translates to the number of potential chances that a single,  
 176 affected community has to spread the rumor to another community. As described in the methods section,  $\beta$   
 177 is not explicitly dependent on the activity assigned to each individual as long as  $a > 0$  and still remains a  
 178 function of the total number of spreaders in the community and the average amount of time they were active.  
 179 Therefore, the same equation used for spatially structured populations,  $\beta = \frac{2(1+\lambda/\alpha)\bar{N}}{\alpha}$ , holds. We can thus  
 180 calculate the rumor invasion threshold by assuming that  $\frac{\lambda\mu}{V-1} \ll 1$  in the limit of large  $V$ , obtaining:

$$181 \quad R^* = 2\left(1 + \frac{\lambda}{\alpha}\right) \frac{\lambda\bar{N}\mu}{\alpha} \geq 1. \quad (6)$$

182 The rumor invasion threshold in terms of the critical inter-community interaction rate,  $\mu_c$  reads,

$$183 \quad \mu_c = \frac{\alpha}{2\lambda\bar{N}\left(1 + \frac{\lambda}{\alpha}\right)}. \quad (7)$$

184 This expression resembles the mobility threshold of Eq. 4 except for the addition of the  $\lambda$  parameter in the  
 185 denominator, which comes from the node interaction process. In the spatially structured model, the mobility  
 186 of an individual was the only factor that controlled whether the rumor spread to a new community. However,  
 187 in the activity-driven model, active individuals do not move to another community, but rather may form a  
 188 single connection through which the rumor has to be successfully transmitted to another individual in order  
 189 to start the contagion process. This introduces a linear dependence on  $\alpha/\lambda$  rather than  $\alpha$  alone. When  
 190 the spreading probability  $\lambda$  is high, the more likely an ignorant individual will transition into a spreader  
 191 during a specific interaction. Thus, the rumor spreads more readily and does not require a high amount  
 192 inter-community interaction to globally propagate. This result shows the inherent differences between Eq. 4  
 193 and Eq. 7 and brings attention to the importance of the type of structured population used when modeling  
 194 a socio-technical system. In the SI we derive the critical interaction probability for populations with a  
 195 heterogeneous size distribution.

### 196 2.3 Simulations in synthetic structured populations

197 In order to validate the analytical findings, we performed an extensive set of stochastic simulations of the  
 198 rumor model on synthetic, structured populations. We generated homogeneous metapopulation networks as  
 199 Erdős-Rényi random graphs with average degrees of  $\langle k \rangle = 12$  and an average population size of  $\bar{N} = 10^3$   
 200 individuals. To initiate the contagion process, one individual is made aware of the rumor in a single, randomly  
 201 selected community. The microscopic *reaction* dynamics are mathematically defined by chain binomial  
 202 and multinomial processes, which were used to update the stochastic transitions of individuals between  
 203 compartments (details in SI). Following the reaction process, individuals *diffuse* along a specific link to a  
 204 neighboring subpopulation with a probability  $p/k$  where  $k$  is the degree of the individual's current community.  
 205 Our simulation results show the final fraction of affected communities as a function of the mobility probability  
 206  $p$  for various  $\alpha$  parameters (see Fig. 2a), recovering the critical transition which separates the non-spreading  
 207 and global spreading dynamical regimes. The vertical lines represent the values predicted from Eq. 4 and  
 208 are in good agreement with our numerical findings. Furthermore, we see a clear dependence of the transition  
 209 point on the stifling rate  $\alpha$ . In the SI we show similar simulations for metapopulation networks with heavy  
 210 tailed degree distributions,  $p(k) \sim k^{-2.2}$ . In this scenario, the network structure significantly reduces the  
 211 threshold since, as mentioned above, it now depends on the second moment of the degree distribution which  
 212 diverges for heterogeneous networks when  $V \rightarrow \infty$ .

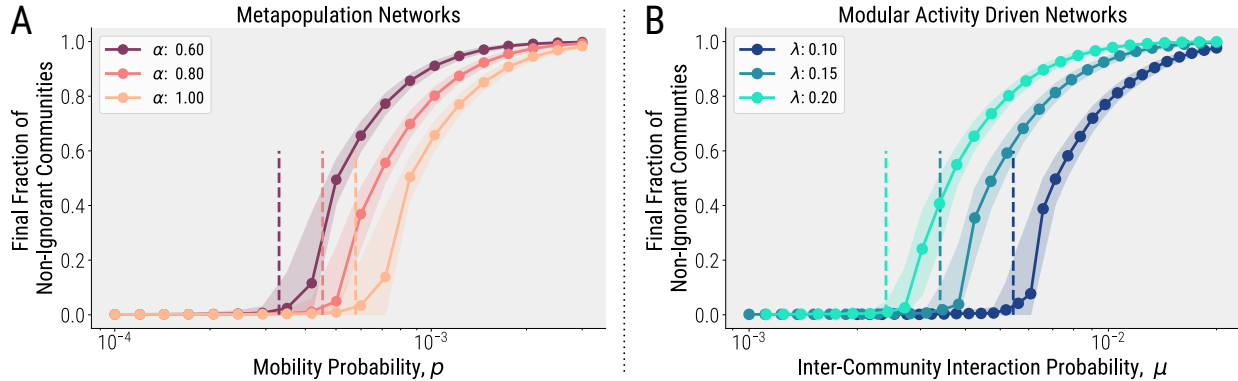


Figure 2: Results from numerical simulations of the rumor spreading process in homogeneous structured populations. **(A)** The final fraction of subpopulations where the rumor is known at the end of the rumor epidemic as a function of the mobility rate  $p$ , for varying  $\alpha$  values averaged over 4,000 simulations. The networks have  $V = 10^3$  subpopulations with an average of  $\bar{N} = 10^3$  individuals and an average degree  $\langle k \rangle = 12$ . The vertical lines represent the predicted threshold values from Eq. 4 and the value  $\lambda = 0.1$  was used for the spreading probability. **(B)** The final fraction of communities where the rumor is known as a function of the inter-community interaction probability for varying  $\lambda$  values averaged over 1,000 simulations. A total population had  $V = 10^3$  communities, each containing  $10^3$  individuals. The vertical lines represent the predicted values from Eq. 7 and a value of  $\alpha = 1$  was used as the stifling probability. The error regions represent the 90% reference range and the averages in the supercritical regime were calculated on the simulations where at least 5% of the subpopulations experienced a rumor epidemic.

213 For the second type of structured population, we generated modular activity-driven networks with  $V =$   
 214  $10^3$  total communities, each with the same number of individuals,  $\bar{N} = 10^3$ , and every individual assigned  
 215 the same activity probability,  $a = 0.1$ . To start the contagion process, a single individual from a randomly  
 216 selected community is made aware of the rumor. An instantaneous network is generated by the modular  
 217 activity-driven network model on which the rumor dynamics unfold for a single iteration. After the reaction  
 218 process, a new network instance is generated and the process repeats until all individuals are either still  
 219 ignorants or stiflers (more model details in SI). In Fig. 2b we show the final fraction of communities where  
 220 a rumor epidemic occurred as a function of the inter-community interaction parameter  $\mu$ , for multiple  $\lambda$   
 221 values. The phase diagram supports our theoretical findings, and confirms that for higher values of  $\lambda$  less  
 222 inter-community interactions are required for the rumor to globally propagate. Additionally, we also model  
 223 this system using a heterogeneous size distribution and report the results in the SI.

### 224 3 Data-driven Simulations

225 To further support the theoretical results obtained in the previous section, we analyze the rumor model  
 226 on two real-world networks. Specifically, we simulate a rumor spreading across a metapopulation network  
 227 modeling the transportation patterns in Europe and across a modular activity-driven network modeling  
 228 scholarly collaborations from the DBLP collaboration network. The mobility of individuals throughout  
 229 Europe is constructed by dividing the continent into spatial regions that are coupled together using data  
 230 about commuting patterns and long-range transportation fluxes such as airline traffic (details in SI). This  
 231 realistic, synthetic metapopulation network has been used in simulations of emerging infectious diseases as  
 232 well as in the analysis and predictions of pandemic events [46, 47, 10]. In this framework, the mobility  
 233 of individuals across subpopulations (analogous to the  $p$  parameter in spatially structured populations) is  
 234 derived from actual transportation data. To study the effects of a reduction in mobility, we rescaled the

235 proportion of individuals that travel at each time step by a factor  $\omega$ . We show the results of the rescaled  
236 mobility on the spatial diffusion of a rumor simulated over the transportation network in Fig. (3a,c).  
237 Interestingly, we see a clear transition in the mobility required for the rumor to spread. The phase diagram  
238 reveals that the critical  $\omega_c$  in this system is significantly small implying that the current human mobility  
239 pattern across Europe is orders of magnitude above the rumor threshold.

240 We also model the collaboration process of the DBLP co-authorship network using the modular activity-  
241 driven network scheme [48]. Nodes in the network are individual researchers that can form collaborations with  
242 others either within or outside of their own communities. In particular, a link represents co-authorship on at  
243 least one paper and each community is a specific publication venue. We measure the amount of interaction  
244  $\mu$  between communities by calculating the frequency of cross-community links relative to the total number  
245 of internal and external links. We simulated the rumor model to analyze how information would propagate  
246 in this system by rescaling the actual individual's tendency to link outside of their current community  
247 (analogous to the  $\mu$  parameter in the virtual structured population framework) by a factor  $\omega$  to study the  
248 effects of lowering inter-community interaction rates. In Fig. (3b,d), we show the results of this rescaling  
249 on the final fraction of affected disciplinary communities and observe a transition point characterizing the  
250 amount of inter-community collaboration needed in order for a rumor or idea to spread globally. Similar  
251 to the transportation network, the critical rescaling value  $\omega_c$  is extremely small. Both data-driven network  
252 applications extend our modeling framework by incorporating heterogeneous and non-trivial subpopulation  
253 interactions. Consequently, the assumptions of statistical equivalence of nodes and an uncorrelated network  
254 structure made in our calculations are no longer valid. Therefore, in these realistic systems, the critical  
255 value  $\omega_c$ , can not be easily computed analytically. However, we do see a similar phenomenology between the  
256 synthetic and data-driven structured populations in that there does exist a critical transition point in the  
257 amount of interaction between subpopulations or communities that is necessary for the a rumor to propagate.  
258 In both cases, the critical transition point ( $\omega_c$ ) is very small, highlighting the role of our interconnected world  
259 in facilitating the diffusion of information across geographical boundaries as well as through disciplinary  
260 communities. However, this result is not necessarily universal across all types of structured populations.  
261 Information spreading is fundamentally dependent on the strength of interactions among elements of the  
262 network, thus calling for specific case by case studies on the location of critical transition points in real world  
263 situations.

## 264 4 Discussion and Conclusion

265 In this work, by using a classic rumor spreading model lacking any critical threshold in a single homogeneous  
266 population, we show that the contagion process in structured populations exhibits a phase transition with a  
267 critical threshold dependent on the amount of interactions/coupling between subpopulations. The analytical  
268 and numerical results presented here emphasize the importance of accounting for the complex structure  
269 observed in socio-technical systems when studying social contagion processes. The features observed in real-  
270 world systems can potentially alter the theoretical picture and the understanding provided by only studying  
271 stylized models. Our results show that successful information or rumor spreading is the result of a complex  
272 interaction between the intrinsic properties of the contagion process and the dynamics of interactions between  
273 subpopulations/communities that comprise social systems. The flexibility of the framework allows for further  
274 study of different types of emergent behaviors that may be more complex than the rumor model used here.  
275 For example, in order for a contagion to spread, individuals must be contacted by multiple neighbors in  
276 their social network. Analogously, additional features can be incorporated into the interaction process and  
277 network structure such as age-dependent contact patterns, socio-economic conditions, and data-driven human  
278 mobility. These features have the potential to not only provide unexpected results of theoretical nature but  
279 also actionable insights crucial to understand and control social contagion phenomena.



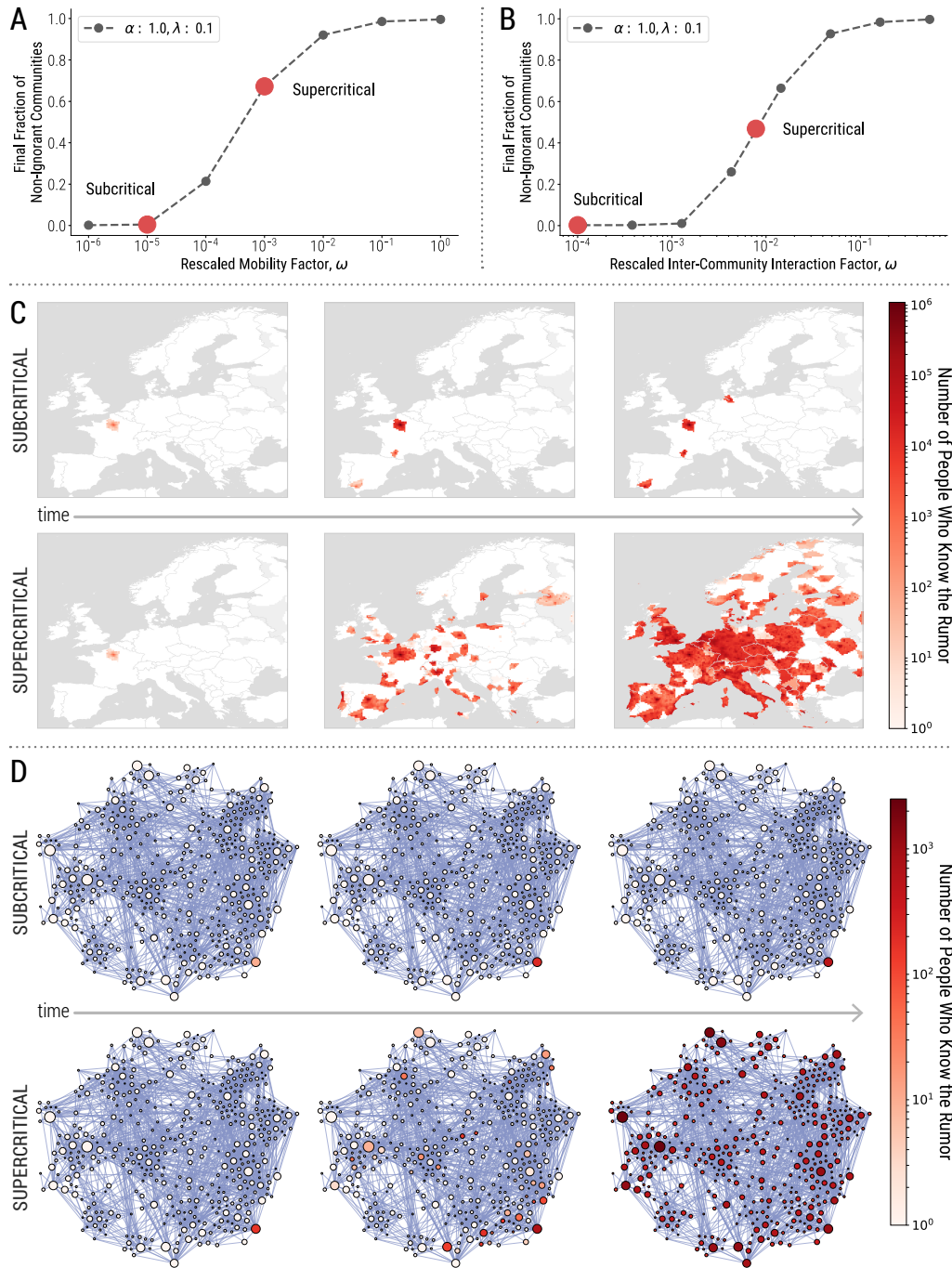


Figure 3: Results from numerical simulations of a rumor spreading in real-world networks. (A-B) The average final fraction of stiflers as a function of (A) the rescaling mobility factor in the European commuting and transportation network and (B) the rescaling factor of the inter-community interaction probability within the DBLP collaboration network. Simulations used a spreading probability of  $\lambda = 0.1$  and stifling probability of  $\alpha = 1$ . Error bars represent the 90% reference range of the simulations where at least 5% of the subpopulations experienced a rumor epidemic. (C-D) The temporal evolution of the rumor spreading process taken from individual simulations corresponding to the  $\omega$  values highlighted in (A-B) with red circles. (C) In the European commuting and transportation network, the rumor was initiated Paris, France by seeding one individual. (D) In the DBLP collaboration network, nodes represent publication venues where node size corresponds to the population size and the line thickness corresponds to the amount of inter-community interaction between each pair of communities. The activity probability per individual is  $a = 0.1$ , while the inter and intra-community probability is derived from the data.

280 **Materials and Methods**

281 **Final Rumor Size in a Single Population:** The mean field rate equations for the Maki-Thompson  
 282 (MK) rumor model in a homogeneously mixed population are listed below. The densities of spreaders (S),  
 283 ignorants (I) and stiflers (R) in a population are defined by  $s = S/N$ ,  $i = I/N$ ,  $r = R/N$ , respectively, where  
 284  $N$  is the total number of individuals in the population, yielding:

$$\begin{aligned}
 \frac{di}{dt} &= -\lambda s(t)i(t) \\
 \frac{ds}{dt} &= \lambda s(t)i(t) - \alpha s(t)(s(t) + r(t)) \\
 \frac{dr}{dt} &= \alpha s(t)(s(t) + r(t)).
 \end{aligned}
 \tag{8}$$

286 Using the initial conditions  $i(0) \approx 1$  and  $r(0) = 0$ , a solution to these differential equations can be obtained  
 287 analytically in the infinite time limit. The transcendental equation below has a trivial solution when  $r_\infty = 0$ ,  
 288 but a non-trivial solution,  $r_\infty = 1 - e^{-(1+\frac{\lambda}{\alpha})r_\infty}$ , when  $\lambda/\alpha + 1 > 1$ , confirming that a rumor will propagate  
 289 through a population and reach a macroscopic fraction of individuals [38]. Assuming that  $(1 + \frac{\lambda}{\alpha})r_\infty \ll 1$ ,  
 290 we can obtain the approximate solution:

$$r_\infty \simeq \frac{2\frac{\lambda}{\alpha}}{(1 + \frac{\lambda}{\alpha})^2} \approx 2\frac{\lambda}{\alpha}
 \tag{9}$$

292 We can see that the final density of stiflers scales with  $\lambda/\alpha$ . This relationship is verified through numerical  
 293 simulations of the rumor model for a single homogeneously mixed population as detailed in the Supplemen-  
 294 tary Material.

295 **Average Spreading Time:** The average spreading time,  $\langle \tau \rangle$ , is the time elapsed since the individual  
 296 was first told the rumor to the time the individual became a stifier. Fig. 1b in the SM shows the average  
 297 spreading time as a function of  $\frac{1}{\lambda} + \frac{1}{\alpha}$  from simulations done on a single population. A linear line is fit to  
 298 the data, producing the equation:

$$\langle \tau \rangle = \frac{1}{\lambda} + \frac{1}{\alpha} + \frac{1}{2}
 \tag{10}$$

302 **Number of potential spreaders:** The number of potential spreaders  $\beta$  that could transmit the rumor to  
 303 another population can be calculated as  $\beta = \langle \tau \rangle \bar{N} r_\infty$  where  $\langle \tau \rangle$  is the average amount of time an individual  
 304 remains a spreader, and  $\bar{N} r_\infty$  is the final average number of individuals that know the rumor at the end of  
 305 the spreading process. Using the approximated equations for the final stifier density as well as the average  
 306 spreading time, one obtains:

$$\beta = r_\infty \bar{N} \langle \tau \rangle = \frac{2(1 + \frac{\lambda}{\alpha})\bar{N}}{\alpha}.
 \tag{11}$$

310 In the modular activity-driven network model, the expression for  $\beta$  is not altered by the fact that individu-  
 311 als are "activated" with probability  $a$ . The average spreading time  $\langle \tau \rangle$  should be measured by considering  
 312 the duration of the contagion process, which should be on the order of  $1/a$  (average number of time steps  
 313 between activations), and the activity of the individual at each time step, which is  $a$ . It follows that these  
 314 terms cancel each other out, so the effective number of interactions of each spreader will not be dependent  
 315 on  $a$  as shown also numerically in the SM.

317 **Data Availability Statement**

318 The data represented in Fig 3b are available through the Stanford Network Analysis Project (SNAP) [49].  
319 All other data that supports the plots within this paper and other findings of this study are available from  
320 the corresponding author upon reasonable request.

321 **Code Availability**

322 Code is available upon request from the corresponding author.

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329 **Author contributions**

330 J.T.D., N.P., Y.M, and A.V. designed the research, J.T.D., A.V., and Q.Z., performed research and analyzed  
331 data, J.T.D., N.P., Q.Z., Y.M., and A.V. wrote the manuscript.

332 **Competing Interests**

333 The authors declare no competing financial interests.

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