Erdós-Rényi phase transition in the Axelrod model on complete graphs

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The Axelrod model has been widely studied since its proposal for social influence and cultural dissemination. In particular, the community of statistical physics focused on the presence of a phase transition as a function of its two main parameters, F and Q. In this work, we show that the Axelrod model undergoes a second-order phase transition in the limit of $F \to \infty$ on a complete graph. This transition is equivalent to the Erdős-Rényi phase transition in random networks when it is described in terms of the probability of interaction at the initial state, which depends on a scaling relation between F and Q. We also found that this probability plays a key role in sparse topologies by collapsing the transition curves for different values of the parameter F.

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I. INTRODUCTION

The Axelrod model, originally proposed for cultural dissemination [1], is grounded in two key dynamical features: social influence, through which people become more similar when they interact; and homophily, which is the tendency of individuals to interact preferentially with similar ones. Specifically, the agents are described by a vector of F components called cultural features, which can take one of Q integer values called cultural traits. The dynamics of the model is based on an imitation rule: A random agent adopts a cultural trait of another one with a probability proportional to the number of shared features.

Despite its simplicity, the Axelrod model attracted the attention of the statistical physics community due to the emergence of a phase transition from a monocultural to a multicultural state [2,3]. The phase transition takes place by varying the number of cultural traits Q for a given fixed F. If the number of cultural traits is low, the probability of interaction is high, leading the system to a monocultural state. If Q is high, the mentioned probability is low and after few interactions the system evolves to a stationary multicultural state.

This phase transition is usually studied by taking the size of the biggest fragment as the order parameter. The transition was reported to be continuous for one-dimensional networks and discontinuous for two dimensions (2D) when F > 2 [4], although a continuous transition is recovered when the topology becomes small world [3]. On the other hand, for F = 2, the type of the transition is the opposite: continuous for 2D, and discontinuous for small-world networks [5]. The case of F = 2 is important due to the possibility of taking an analytical approach to study the model [6].

Several scaling relationships have been found in the Axelrod model. For instance, in [7] and [5] a finite-size scaling analysis is performed for F=2 in square lattices and smallworld networks. A scaling relation between Q and size N can be found in scale-free networks for F=10 [3], a scaling relationship between the density of active bonds and time in a one-dimensional network in [8], and the finding of an effective noise rate is explored in [9].

Among the reported scaling relations, there is a particular one reported in [10] and [11] where the transition curves in one-dimensional networks collapse when the control parameter is F/Q. This ratio has an immediate interpretation as the mean value of shared features given two agents in the initial state, suggesting that the initial distribution contains key information about the final outcome of the Axelrod model.

In this work, we review the Axelrod model in terms of the initial interaction probability between agents $(p_{\rm int})$. In particular, we found that the second-order phase transition in the limit of $F \to \infty$ on a complete graph is equivalent to the phase transition observed in Erdős-Rényi random networks [12]. Notably, we also observed that this interaction probability between agents $(p_{\rm int})$ plays a key role in describing the Axelrod model on sparse topologies, by making the transition curves collapse for different values of F.

II. MODELS

A. Axelrod model

The Axelrod model [1] describes each agent by a vector of F components, which represents a set of cultural features. The initial state of the system is set by assigning with equal probability one of Q integer values to each feature. The value of Q represents the number of different cultural traits that a particular feature adopts. Once the initial condition is set, the dynamics of the system is based on a pairwise interaction mechanism, which relies on two fundamental hypotheses:

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- (1) Homophily: The probability of interaction between two individuals is proportional to their cultural similarity, that is, the number of features they share.
- (2) Social influence: After each interaction, the agents become more similar. It means that one of the agents copies a feature from the other which they previously did not share.

This model shows a nonequilibrium phase transition from a monocultural to a multicultural state by varying the value of Q for a fixed F. When $Q < Q_c$, the probability that two agents can interact since the initial state is high, so all agents end with the same cultural vector. On the other hand, when $Q > Q_c$, the probability of interaction at the initial state is low so the final state shows a coexistence of regions with different cultural states.

The transition is usually characterized by measuring the size of the biggest fragment. We define a fragment as a group of topologically connected agents with at least one feature in common. At the final state, the biggest fragment is made up only by agents with the same cultural state.

B. Erdős-Rényi graph

The Erdős-Rényi network is a graph model in which a set of N initially disconnected nodes are linked with probability p. The outcome of this model is a network with a binomial degree distribution of parameters N and p, with mean degree k = (N-1)p.

An interesting feature of this model is that the size of the largest connected component shows a second-order phase transition at $k_c = 1$: When $k < k_c$ the largest component has a finite size, while for $k > k_c$ it scales with the size of the system [12,13].

C. Connection between both models

Given two agents, the parameters F and Q set their initial number of shared features by sampling this quantity from a binomial distribution with parameters F and 1/Q. If we define the initial interaction probability p_{int} as the probability that two agents are able to interact at the initial state (i.e., the probability of sharing at least one feature), then

$$p_{\rm int} = 1 - \left(1 - \frac{1}{O}\right)^F. {1}$$

Given this definition, we can think of the initial state of the Axelrod model on a complete graph as an Erdős-Rényi model with parameter p_{int} . In other words, we discriminate between links with zero and nonzero homophily, being p_{int} equal to the fraction of pairs of agents that can interact at the initial state.

III. RESULTS

A. Axelrod model in complete graphs

We first analyze the Axelrod model on a complete graph of N=1024 agents. Figure 1 shows that the relative size of the biggest fragment (S_{max}/N) in the final state is smaller than its initial value for low F. However, this difference approaches to zero when F increases. This result suggests that the relative size of the stationary biggest fragment is fully determined at the initial condition in the limit of $F \to \infty$. The importance

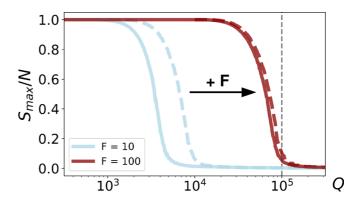


FIG. 1. Axelrod transition in a complete graph for different values of F. Solid lines show the relative size of the biggest fragment $S_{\rm max}/N$ as a function of the parameter Q for N=1024 at the final state. Dashed lines show their initial values for each F. When F increases (pointed out by the arrow), $S_{\rm max}/N$ set by the Axelrod dynamics becomes closer to its initial value. Vertical dashed lines point to the critical value $Q_c=10^5$ mentioned in the text. Light blue (light gray) corresponds to F=10 while dark red (dark gray) corresponds to F=100.

of the initial condition is reflected in the fact that two agents who initially do not share any feature cannot interact, at least until other interactions take place and eventually change their cultural states.

The suggestion of the equivalence between both transitions in the limit of $F \to \infty$ is clearer by taking p_{int} as the control parameter. Figure 2 shows this for the relative size of the biggest fragment [panel (a)] and the average finite-fragment

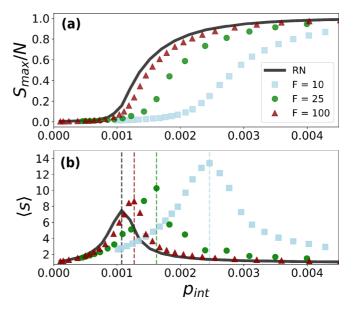


FIG. 2. Relative size of biggest fragment $S_{\rm max}/N$ and average fragment size $\langle s \rangle$ as a function of $p_{\rm int}$ for N=1024. Both figures show that the Axelrod transition tends to the Erdős-Rényi (RN) transition for increasing F. Light blue (squares) corresponds to F=10, green (circles) to F=25, dark red (triangles) to F=100, and black solid line to the random network model. Dashed lines in panel (b) point out the critical values of $p_{\rm int}^c(N)$ for different F and the random network model.

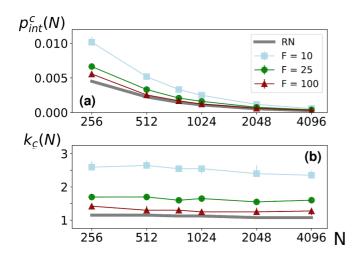


FIG. 3. Critical point and mean degree for different values of F. It can be observed that when F is increased, $p_{\text{int}}^c(N)$ tends to the random network critical point at fixed N, while k_c tends to 1 for all N. Light blue (squares) corresponds to F = 10, green (circles) to F = 25, dark red (triangles) to F = 100, and black solid line to the random network model.

size [panel (b)]. Moreover, the definition of $p_{\rm int}$ [Eq. (1)] allows one to estimate the critical value of the transition Q_c for large F. Since the biggest fragment emerges in an Erdős-Rényi network when $Np_{\rm int}^c \sim 1$ [13],

$$Q_c \sim \left[1 - \left(1 - \frac{1}{N}\right)^{1/F}\right]^{-1},$$

in the Axelrod model. Applying this analogy for N = 1024 and F = 100, it gives $Q_c \sim 10^5$, as can be seen in Fig. 1.

1. Critical point and exponents

As can be observed in Fig. 2, the critical point tends to the Erdős-Rényi one for large F. Figure 3 shows that $p_{\text{int}}^c(N)$ effectively tends to the corresponding value in random networks when F is increased at fixed N [panel (a)]. Although the differences between F values seems to vanish when $N \to \infty$, if we define the mean degree

$$k_c = (N-1)p_{\rm int}^c(N),$$

as usual for the Erdős-Rényi model, we can see in panel (b) that $k_c(F)$ is stable as a function of N and tends to the theoretical value $k_c = 1$ for large F (see Sec. II B). In the case of the Axelrod model, k is interpreted as the average number of neighbors that a given agent can interact at the initial state.

The equivalence between Erdős-Rényi's and the Axelrod model phase transition's for large F can be completed by the calculation of the critical point and critical exponents in the thermodynamics limit. Due to the closeness of the transition curves observed in Fig. 2, we estimate them for F = 100 by applying finite-size scaling according to the procedure sketched in [14]. We provide details of these calculations in the Appendix.

Table I shows that our estimations of the critical exponents are consistent with the theoretical values predicted for the Erdős-Rényi model [13]. We can observe that in all cases, the

TABLE I. Critical exponents and critical point. Theoretical values for the Erdős-Rényi phase transition and estimations from the Axelrod model with F=100, with their respective 95% confidence intervals (CI). We also report the 95% confidence intervals of the random network (RN) model applying finite-size scaling within the same range of N.

	Theoretical	Axelrod	95% CI	RN 95% CI	
		Critical	exponents		
ν	1	0.88	0.85-1.01	0.87 - 1.02	
β/ν	1/3	0.42	0.11-0.52	0.20-0.38	
γ/ν	1/3	0.36	0.33-0.46	0.33-0.35	
τ	5/2	2.45	2.41-2.61	2.49 - 2.53	
	Critical point				
p_{int}^c	0	7×10^{-5}	$(-7-10) \times 10^{-5}$	$(-5-9) \times 10^{-5}$	

theoretical value is included in the 95% confidence interval of our estimation. Moreover, applying the finite-size scaling methodology to finite random networks within the same range of N, we found that the matching between both models is even stronger.

Finally, our estimation of the critical point was $p_{\text{int}}^c(N = \infty) = 7 \times 10^{-5}$, which is consistent with the theoretical value of 0 within a 95% confidence interval (see Table I).

2. Other topological features

It should be noted that the equivalence between the Erdős-Rényi and the Axelrod model phase transition for large F (described by similar critical exponents) will not necessarily be present in other topological features given the dynamical evolution of the Axelrod model. Figure 4 shows the time evolution of S_{max}/N , the relative multiplicity RM, defined as the number of fragments normalized by N, and the average clustering coefficient of the biggest fragment $\langle C \rangle$, for different values of F at fixed p_{int} . We also show the expected value of the Erdős-Rényi model at the same probability. It can be noticed that for large values of F, S_{max}/N and RM tend to be constant and closer to their initial value which is similar to the random graph model (pointed out by dashed lines in the figure). On the other hand, $\langle C \rangle$ always ends up reaching the value of 1 (in this case, $\langle C \rangle \sim 0$ for Erdős-Rényi). The observed behavior of S_{max}/N and RM tells us that the fragments at the initial state tend to break apart during the dynamics, while the increment of the clustering coefficient points out that the connected components become cliques at the final state.

The evolution of these quantities could be understood as follows. First, we say that a link is active if it connects two agents with at least one feature in common (i.e., a nonzero homophily link). Given an active link between two agents, the Axelrod model always tends to increase their similarity. An active link can only become inactive by third-party interactions. When the value of F increases, the probability that an active link becomes inactive decreases and goes to zero when $F \to \infty$. In this limit, the initial active links define the sizes of the connected components (as in the Erdős-Rényi model) and the only effect of the dynamics is to transform all the connected components in cliques of the same size.

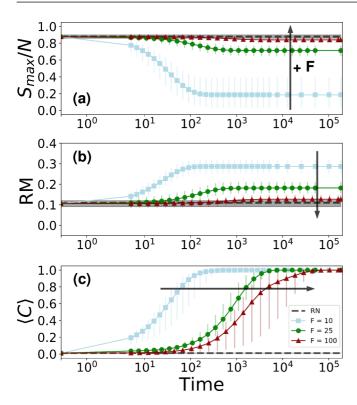


FIG. 4. Relative size of the biggest $S_{\rm max}/N$ [panel (a)], relative multiplicity of fragments RM [panel (b)], and average clustering coefficient $\langle C \rangle$ of the biggest fragment [panel (c)]. These simulations belong to N=1024 and $p_{\rm int}=0.0024$. Similar results were observed for different values of $p_{\rm int}$. Light blue (squares) corresponds to F=10, green (circles) to F=25, dark red (triangles) to F=100, and black dashed lines to the random network model (gray bands denote error bars). Arrows point out increasing F and time is measured per N interactions.

B. Axelrod model in 2D lattices

Let us now explore the Axelrod transition in a twodimensional lattice as a function of the new control parameter $p_{\rm int}$. Figure 5 shows the transition curves for different values of F as a function of $p_{\rm int}$, in addition to the initial relative size of the biggest fragment. The inset of this figure shows the same curves as a function of Q, where it can be seen that the transition shifts to larger values of Q when F increases.

Figure 5 shows that all transition curves collapse to one within the error bars. This collapse is also found in other sparse topologies as random regular networks with equivalent mean degree (not shown). In contrast to the observed behavior in complete networks (see Fig. 1), the collapsed curve does not match that corresponding to the initial state.

The collapse as a function of p_{int} is essentially the same pointed out by [10] for one-dimensional networks. As was mentioned in Sec. II C, F and Q set the number of shared features for a pair of agents by sampling this quantity from a binomial distribution with parameters F and 1/Q. These quantities also set the value of p_{int} . In the limit of large F and Q, this binomial distribution can be well approximated by a Poisson one with parameter F/Q, which is the mean shared features by two random agents and the control parameter introduced in [10] for one-dimensional networks.

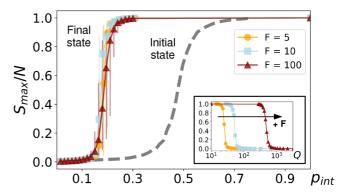


FIG. 5. Axelrod transition in a 2D lattice. Relative size of the biggest fragment as a function of $p_{\rm int}$ for both the initial (dashed line, the same for all F) and the final state for different F (markers with error bars). Same quantity as a function of Q for different values of F are shown in the inset (arrow points out increasing F). System size is N=1024. Orange (circles) corresponds to F=5, light blue (squares) to F=10, and dark red (triangles) to F=100.

We can conclude here that, for sparse topologies, the Axelrod model can be described by a control parameter related to the similarities among agents at the initial state, in particular to the fraction of nonzero homophily links (p_{int}) . This is related to what we have observed for complete graphs when F is large. The perspectives of these observations are sketched in the conclusions of our work.

IV. CONCLUSIONS

In this work, we show that the Axelrod model on complete graphs displays a phase transition equivalent to the one observed in the Erdős-Rényi model when the order parameter is plotted as a function of p_{int} , which is the probability that two agents share at least one feature in the initial state. This happens in the limit of $F \to \infty$. This claim is supported by the calculation of critical exponents and the critical point following the approach of finite-size scaling sketched in [14].

This equivalence can be understood by observing that when F increases, the size of the fragments as well as the amount of them are preserved during the dynamics. In other words, the probability that the fragments set in the initial state break apart goes to zero for large F. However, the Axelrod dynamics is displayed by leading to a stationary state where the connected components also become cliques. In summary, it means that in the limit $F \to \infty$, the initial fragments which are similar to an Erdős-Rényi realization with parameter $p_{int}(F, Q)$ are preserved but transformed in cliques. An implication of this equivalence is that the final state of the Axelrod model is highly predictable from its initial state for large F. A possible future research line could be to explore if this conclusion is extensible to other dense topologies such as random regular networks with high mean degree.

On the other hand, when the Axelrod model is studied in terms of p_{int} on sparse graphs, we found the collapse of transition curves for different values of F. These collapsed curves do not coincide with the initial state, as they did in complete networks for $F \to \infty$. However, this means that

both the transition for complete graphs for large F and sparse topologies can be described in terms of a quantity related to the similarity between agents at the initial state, in particular to the fraction of links with nonzero homophily $p_{\rm int}$. A scope of future works will be based on rewriting the Axelrod model in terms of the similarity between agents in order to analyze the explicit dependence of this model on quantities related to the initial similarity distribution.

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APPENDIX: CRITICAL EXPONENTS AND FINITE-SIZE SCALING

The critical exponents [13] are introduced following the usual relationships. Near the critical point,

$$rac{S_{
m max}}{N} \sim \left(p_{
m int} - p_{
m int}^c\right)^{eta}, \ \langle s
angle \sim \left|p_{
m int} - p_{
m int}^c\right|^{-\gamma},$$

where p_{int}^c is the critical probability, and S_{max}/N is the relative size of the biggest fragment and the order parameter, whose fluctuations are measured by $\langle s \rangle$, which is the average finite-fragment size.

For finite systems, the critical exponents can be calculated by performing finite-size scaling following [14]. Here, the authors propose the following scaling relationships:

$$\frac{S_{\text{max}}}{N} = N^{-\beta/\nu} F_1 [(p_{\text{int}} - p_{\text{int}}^c) N^{1/\nu}],$$
$$\langle s \rangle = N^{\gamma/\nu} F_2 [(p_{\text{int}} - p_{\text{int}}^c) N^{1/\nu}],$$

where F_1 and F_2 are unknown scaling functions, but with the property that $F_{1(2)}(x) \to \text{constant when } x \to 0$ (which means, near the critical value). This implies that exactly at the critical point,

$$\frac{S_{\text{max}}}{N} \sim N^{-\beta/\nu},\tag{A1}$$

$$\langle s \rangle \sim N^{\gamma/\nu}.$$
 (A2)

These expressions can be used to estimate the relation between exponents without knowledge about F_1 and F_2 . On the other hand, the argument of these scaling functions defines

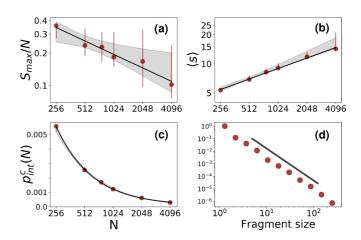


FIG. 6. Finite-size scaling and critical behavior for F = 100. Red circles are point estimates while full lines point out the fitted curves. Circle error bars and light-gray bands denote the 95% confidence interval of each estimation. Panel (d) belongs to the fragment distribution at the critical point with N = 4096.

another relationship followed by the exponent ν and the critical value p_{int}^c , which can be read in the following equation:

$$p_{\text{int}}^c(N) = p_{\text{int}}^c - bN^{-1/\nu},$$
 (A3)

where $p_{\text{int}}^c(N)$ is the pseudocritical point in which $\langle s \rangle$ takes its maximum value, being $p_{\text{int}}^c = p_{\text{int}}^c(\infty)$. Finally, the fragments size distribution f(s) near the critical point follows a power-law distribution with parameter τ [Eq. (A4)]. This relation defines this last critical exponent, which was calculated following the methodology sketched in [15].

$$f(s) \sim s^{-\tau}.\tag{A4}$$

Figure 6 shows an example of the calculus of the exponents for F=100, which were reported in Table I. The relations of Eqs. (A1)–(A4) are respectively plotted in panels (a)–(d). We also estimated the 95% confidence intervals of all quantities by bootstrapping [16], i.e., by recalculating them several times over different datasets generated by sampling with replacement of the original measures taken from our simulations. In the estimation of $p_{\text{int}}^c(\infty)$, we let this parameter fluctuate around zero including negative values despite not being conceptually correct, in order to include zero in the confidence interval in the case in which it is supported by the statistics. The same methodology was respectively implemented for each value of F for the calculus of $p_{\text{int}}^c(N)$ of Fig. 3.

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