

An SIS-type Marketing Model on Random Networks

Reinhard Illner and Junling Ma

Department of Mathematics and Statistics, University of Victoria,
Victoria, BC V8W 2Y2

¹rillner@uvic.ca

²junlingm@uvic.ca

May 22, 2015

Abstract

Marketing on random networks displays similarities to epidemiological models in the sense that “word-of-mouth” information passes between individuals and may “infect” susceptible buyers such that they end up buying the product. The difference to epidemics is that there are usually many competing products (rather than just one disease), and in addition to word-of-mouth transmission, products are also advertised by the producers, which can be thought of as external nodes connected to the network. In this paper we develop a model in which these various transmission pathways compete, and, in addition, where product fatigue and product switching are possible. This is a genuine and realistic extension of the model developed in [4], where a customer would never abandon a product after purchase. The model presented here is similar to and was inspired by SIS epidemiological models.

Keywords: random network, SIS marketing model, product fatigue, product switch

Mathematics Subject Classification (2010): 92D60

1 Introduction

In [4] the classical Bass model, which models propagation of a new product due to advertising and word-of-mouth marketing in a completely connected and homogeneous population, was generalized to populations modelled as a random graph of configuration type [7], and to several competing products. The underlying assumptions in this model were that a potential buyer (a “susceptible”) is subject to independent efforts by the competing companies (efforts realized by advertising) and to the independent influences transmitted by his/her neighbours or acquaintances, who will praise the virtues of the product they have already bought. Once the susceptible buyer has in fact purchased a product, he/she is assumed to stick with this product forever.

These assumptions allow to generalize the ideas first brought forward by Miller and Volz ([5, 8]) in the context of epidemiology to marketing. However, buyers may not stay loyal to their product, as always happens in reality. Products age, break down, become obsolete, or a customer may simply be intrigued by a competing product, abandon what he/she already owns and go shopping for a new product.

This more realistic scenario has a counterpart in epidemiology known as SIS (susceptible-infected-susceptible) models, in contrast to SIR (susceptible-infected-recovered) models. In SIS models, an individual will recover from the disease but will again be susceptible; there is no recovery to an immune state. SIS models on random graphs have been introduced in [3]. Here we adapt the ideas from this work to the marketing problem, with product fatigue (“recovery”), where the recovered customer returns to being a susceptible customer.

Our model is complicated yet uses some contentious simplifying assumptions (in particular something we call “edge chaos”, a concept borrowed from the kinetic theory of gases; see Section 2). Without such assumptions, the modelling would become rather intractable, as will be explained in due time. Further, we will assume that a customer who has bought a product will not abandon this product because of aggressive advertising by the competition or by his contacts; rather, we assume that the customer will abandon the product due to general product fatigue (like a car, or a computer, or a refrigerator getting so old that it needs to be replaced), and that the customer will then again be susceptible to all competing products. Generalizations to more sophisticated behaviours are certainly possible but will complicate the model further.

The fundamental idea, similar to the methodology used in the references on epidemiology, is to study edge dynamics on random networks. Our model will investigate changes in edge types depending on the status of neighbouring edges. This is difficult, and it is in this context that an edge chaos assumption enters into our discussion.

We design the model for two competing products, similarly to the scenario discussed in [4]. The setup of the model is done in Section 2, and we provide some discussion and analysis. In the subsequent section we compare predictions of the model with microscopic numerical simulations based on Gillespie’s algorithm [1, 2]. The agreement between the two approaches is excellent, although high fluctuations on some types of scale-free random networks impose limitations on the applicability of the model in such cases.

2 Prerequisites, Notation, and the Model

We consider a population modelled as an undirected random network of configuration type. Such a network is defined as a random graph with N nodes (individuals), where N is large and a priori not known. The network is completely described by its edge distribution $\{P_k\}_{k=0,1,2,\dots}$, where P_k is the probability that a randomly chosen node has degree k , i.e., has k neighbours.

For the marketing context we assume that each node (individual) is connected to two competing companies, denoted as A and B , which can “reach” the individual via advertising. Each company has therefore N connections into the network. The (unique) products of A and B will simply be denoted by A and B as well. For convenience we label individuals as $S = S(t)$ (susceptible, has not bought either product), or as $A = A(t)$ or $B = B(t)$ if the individual has bought A or B , respectively. Assuming that no one will buy both products, we immediately get the conservation equation

$$S(t) + A(t) + B(t) = N.$$

If we define S_k as the number of all susceptibles with degree k we have $S = \sum_{k=0}^{\infty} S_k$. If nobody has bought a product at time 0 we have the initial conditions $S_k(0) = P_k N$, $S(0) = N$. Similarly, defining N_k , A_k and B_k in the obvious way, we have

$$S_k + A_k + B_k = N_k.$$

We have suppressed the dependence on t and will continue to do this in the sequel, but the initial conditions are $A_k(0) = B_k(0) = 0$ for all k .

We summarize the small notational abuse implied by the above: A , B denote two competing companies, their (unique) products, and the number of individuals that hold product A or B at a given time. We will assume that nobody will hold both products at the same time.

Marketing the products A and B involves both direct advertising and word-of-mouth “transmission”. In order to incorporate the structure of the network we will keep track of the status of edges. Even though the social network is assumed to be undirected, we assume that each edge consists of two directed edges of opposite direction to keep track of who markets to whom, pointing from *source* to *target*. These directed edges are denoted by pairs like SS , SA , AS , BA , AA and so on. The first letter will always denote the state of the target, the second the state of the source. M_{SS} will be the number of all directed edges with susceptible target and source; note that this number is twice the number of SS -edges, as each edge is counted twice in opposite directions. Similarly we define M_{AS} , M_{SA} , M_{SB} , M_{BA} etc. Note that transmission (advertising) under consideration can occur across M_{SA} but not across M_{AS} edges, (because A can transmit to S but not vice versa). However, we have $M_{SA} = M_{AS}$ because they count the same edges. Similarly, $M_{SB} = M_{BS}$, and $M_{AB} = M_{BA}$.

Further, M_S denotes the number of edges with target S . Hence

$$M_S = \sum_{k=0}^{\infty} kS_k = M_{SS} + M_{SA} + M_{SB} \quad (1)$$

Similarly,

$$M_A = \sum_{k=0}^{\infty} kA_k = M_{AS} + M_{AA} + M_{AB}, \quad (2)$$

$$M_B = \sum_{k=0}^{\infty} kB_k = M_{BS} + M_{BA} + M_{BB}, \quad (3)$$

We denote by β_A (and β_B) the transmission rates from A (B) buyers to susceptible nodes along an edge. In contrast, the per capita “conversion rates” of susceptibles due to outside advertising by companies A and B will be denoted by α_A, α_B . Define further

$$p_A := \frac{M_{SA}}{M_S}, \quad p_B := \frac{M_{SB}}{M_S}. \quad (4)$$

These are the probabilities that an edge with target S has source in A or B , respectively.

A last feature which we implement into our model is “recovery”. Here this means that the products A and B have a finite lifespan (this could be due to technical innovation (the product becomes obsolete), tiredness of the buyer, or malfunction). For simplicity, we assume that there is a constant “recovery” rate $\gamma > 0$, same for A, B , such that owners of A or B recover to join class S at this rate. As mentioned in the introduction, this is quite simplistic and may be modified in future refinements of the model.

These assumptions naturally lead to a first set of equations for the evolution of S_k ,

$$\frac{d}{dt}S_k = -\beta_{AP}p_A k S_k - \beta_{BP}p_B k S_k - \alpha_A S_k - \alpha_B S_k + \gamma(A_k + B_k). \quad (5)$$

While this equation is linear in terms of the unknowns S_k, A_k, B_k , it is the appearance of the p_A, p_B , defined in terms of the unknowns M_{SA} etc., which introduces nonlinearity into the system.

Similarly,

$$\begin{aligned} \frac{d}{dt}A_k &= \beta_{AP}p_A k S_k + \alpha_A S_k - \gamma A_k, \\ \frac{d}{dt}B_k &= \beta_{BP}p_B k S_k + \alpha_B S_k - \gamma B_k. \end{aligned}$$

As mentioned earlier, the initial conditions will be $S_k(0) = P_k N$, and $A_k(0) = B_k(0) = 0$, $p_A(0) = p_B(0) = 0$.

By summing over k and using $\sum A_k = A, \sum B_k = B, \sum k S_k = M_S$ we find

$$\begin{aligned} \frac{d}{dt}A &= \beta_{AP}p_A M_S + \alpha_A S - \gamma A, \\ \frac{d}{dt}B &= \beta_{BP}p_B M_S + \alpha_B S - \gamma B. \end{aligned}$$

Note that $p_A M_S = M_{SA}$ and $p_B M_S = M_{SB}$. They lead to

$$\frac{d}{dt}A = \beta_A M_{SA} - \gamma A + \alpha_A S \quad (6)$$

$$\frac{d}{dt}B = \beta_B M_{SB} - \gamma B + \alpha_B S. \quad (7)$$

Similarly,

$$\frac{d}{dt}M_A = \sum_k k \frac{dA_k}{dt} = \beta_{AP_A} \sum_k k^2 S_k + \alpha_A M_S - \gamma M_A, \quad (8)$$

$$\frac{d}{dt}M_B = \sum_k k \frac{dB_k}{dt} = \beta_{BP_B} \sum_k k^2 S_k + \alpha_B M_S - \gamma M_B. \quad (9)$$

with $M_A(0) = M_B(0) = 0$, $M_S(0) = \sum_k k S_k(0) = N \sum_k k P_k$. The number N of nodes is considered constant; it can of course be scaled out of the equations by considering fractions of susceptibles, A-buyers, B-buyers, and different types of edges rather than numbers.

Deriving equations for the numbers of different types of arcs is more involved and will require some fairly strong assumptions. In particular, we need the concept of the “average excess degree” of a susceptible node after following an edge (this node could be source or target). Specifically, observe that $\sum_{k=0}^{\infty} k S_k$ is the total number of edges starting from a susceptible. Hence, if we follow an edge and arrive at a node of class S , the probability that this node has j edges is $j S_j / \sum_{k=0}^{\infty} k S_k$. (this expression reflects the fact that it is proportionally more likely to reach nodes with more edges if one follows an edge). The *average excess degree* E of such nodes is then

$$E := \sum_{j=1}^{\infty} (j-1) \frac{j S_j}{\sum_{k=0}^{\infty} k S_k}. \quad (10)$$

The factor $j-1$ appears here because the edge we followed is not counted. This E clearly depends on the S_k , but for ease of notation we suppress this dependency in our formulas. E involves only the S_k because nodes of type A or B cannot be changed by neighbours; they can recover, but our assumptions are that the recovery does not depend on the neighbouring nodes. For (10) to be meaningful, we assume that

$$\sum_{j=1}^{\infty} (j-1) \frac{j P_j}{\sum_{k=0}^{\infty} k P_k} < \infty. \quad (11)$$

Also, observe that from the definition

$$p_A M_{SB} = p_B M_{SA}. \quad (12)$$

The dynamics of M_{SA} involves no fewer than 10 terms on the right hand side. Here is the equation.

$$\begin{aligned}
\frac{d}{dt}M_{SA} = & -\underbrace{\beta_A M_{SA}}_1 - \underbrace{\gamma M_{SA}}_2 + \underbrace{\gamma M_{AA}}_3 + \underbrace{\gamma M_{BA}}_4 \\
& + \underbrace{\beta_A E p_A M_{SS}}_5 - \underbrace{\beta_B E p_B M_{SA}}_6 - \underbrace{\beta_A E p_A M_{SA}}_7 \\
& + \underbrace{\alpha_A M_{SS}}_8 - \underbrace{\alpha_A M_{SA}}_9 - \underbrace{\alpha_B M_{SA}}_{10}.
\end{aligned} \tag{13}$$

A chart illustrating the meaning of these 10 numbered gain and loss terms on the right-hand side that represent the flows of edges to and from the M_{SA} class is shown in Figure 1. We discuss these numbered terms one at a time.

Terms 1-4 are easily understood. First, an arc of type SA can leave this class because the target end (S) is recruited (infected) by the source. Second, the source (A) might recover, and the arc leaves the class. Third and fourth, the target might have been of type A or B and recovered, thus adding to the class SA .

Terms 5-7 are a lot harder to explain and include, in fact, an “edge chaos” assumption. For example, let us focus on the gain term 5: This term is there because an arc of type M_{SS} can turn into type M_{SA} because its source can be converted into type A by another neighbour; such neighbours are available with average excess degree E (as defined above) and are of type A with probability p_A (this latter statement is a hidden independence (chaos) assumption; it is not automatically given that if we follow an SS edge from target to source, then neighbours of the source will be type A with probability p_A ; making this assumption means that we neglect possible correlations between the edges.)

Similarly, terms 6 and 7 are present because the target S in an SA arc could be recruited (or infected) by edges other than the source, and such edges could be of type A or B . The terms 4-7 can be written in alternate ways by using Eq. (12). Again, hidden chaos assumptions are made here, to be avoided only at the expense of going to much more complicated models, where two-arc distribution densities would have to be introduced, and the model would lose closure.

Terms 8-10 are much simpler: Term 8 is present because the source of an arc of type SS may be recruited externally by advertising, adding the arc to class M_{SA} . Similarly, terms 9 and 10 are present because the target of an arc

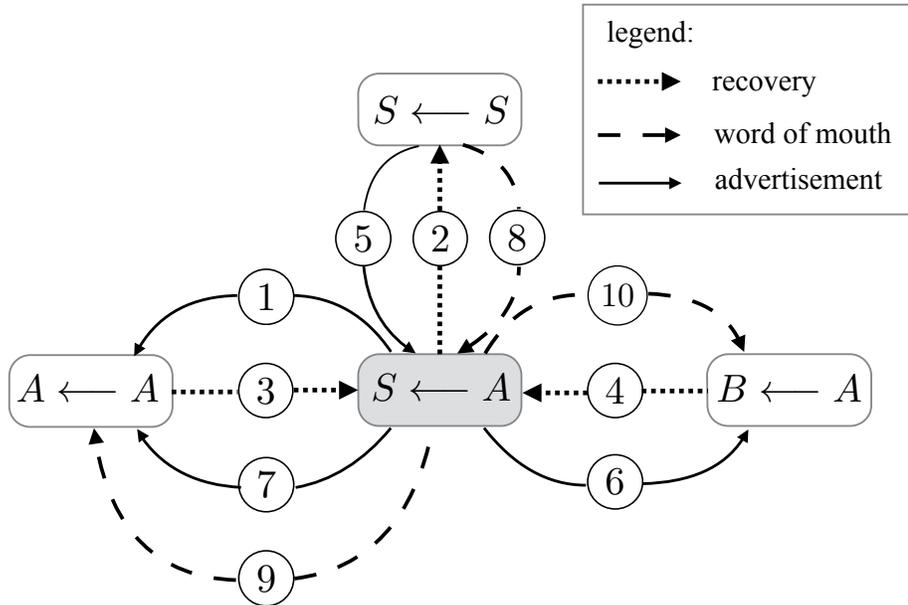


Figure 1: A flow chart for the gain and loss of the edges in the M_{SA} class (for edges $S \leftarrow A$, pointing from the source node A to the target node S), whose dynamics is given in (13). The circled number on each flow corresponds to the such numbered term in (13). Flows 2, 3, and 4 represent gains and losses due to recovery of a buyer on one end of an edge; flows 8, 9, and 10 represent the gains and losses due to the conversion of a susceptible to a buyer by advertisement; the other flows represent the conversion of a susceptible to a buyer by word-of-mouth.

of type SA may be recruited by company A or B, moving the arc into class M_{AA} or M_{BA} , respectively.

The initial conditions for the quantities appearing in this and subsequent equations are

$$\begin{aligned} M_{SA}(0) &= 0, & M_{SB}(0) &= 0, & M_{AA}(0) &= 0, \\ M_{BB}(0) &= 0, & M_{AB}(0) &= 0, \end{aligned}$$

and

$$M_{SS}(0) = M_S(0) - M_{SA}(0) - M_{SB}(0) = N\Psi'(1).$$

A corresponding equation, with the same caveats, holds for M_{SB} . Because of the symmetry between A and B , this equation can be derived by exchanging A and B in (13).

$$\begin{aligned} \frac{d}{dt}M_{SB} &= -\beta_B M_{SB} - \gamma M_{SB} + \gamma M_{BB} + \gamma M_{AB} \\ &\quad + \beta_B E p_B M_{SS} - \beta_A E p_A M_{SB} - \beta_B E p_B M_{SB} \\ &\quad + \alpha_B M_{SS} - \alpha_B M_{SB} - \alpha_A M_{SB}. \end{aligned} \quad (14)$$

We further recall that $M_S = \sum k S_k$. This means, from the equations for S_k , we know the dynamics of M_S , and as $M_{SS} = M_S - M_{SA} - M_{SB}$,

$$M_{SS} = \sum_k k S_k - M_{SA} - M_{SB}. \quad (15)$$

We do not need to write equations for M_{AS} and M_{BS} , because $M_{AS} = M_{SA}$, and $M_{SB} = M_{BS}$.

It remains to derive equations for M_{AA} , M_{BB} and M_{AB} (which also gives M_{BA}). We begin with M_{AB} , as the other two can be derived from it similarly to the derivation of M_{SS} .

$$\frac{d}{dt}M_{AB} = -\underbrace{2\gamma M_{AB}}_1 + \underbrace{\beta_A E p_A M_{SB}}_2 + \underbrace{\beta_B E p_B M_{SA}}_3 \quad (16)$$

$$+ \underbrace{\alpha_B M_{AS}}_4 + \underbrace{\alpha_A M_{BS}}_5 \quad (17)$$

Term 1 here is present because either end of the edge can recover (hence the factor 2); Terms 2 and 3 are present because the S - end of an SB - or

a SA - edge may become recruited by another neighbour (with the previous caveats regarding independence). And last, terms 4 and 5 account for external recruitment (by advertisements) of the S - end of SA - and SB - edges. We then have

$$M_{AA} = M_A - M_{AS} - M_{AB} = \sum_k k A_k - M_{SA} - M_{AB}, \quad (18)$$

$$M_{BB} = M_B - M_{BS} - M_{BA} = \sum_k k B_k - M_{SB} - M_{AB}. \quad (19)$$

In summary, the full model consists of the following equations:

$$M_S = \sum_{k=0}^{\infty} k S_k; \quad E = \frac{\sum_{j=1}^{\infty} (j-1)j S_j}{M_S}; \quad (20a)$$

$$\frac{d}{dt} S_k = -\beta_{AP_A} k S_k - \beta_{BP_B} k S_k - \alpha_A S_k - \alpha_B S_k + \gamma(N_k - S_k); \quad (20b)$$

$$\frac{d}{dt} A = \beta_A M_{SA} - \gamma A + \alpha_A S; \quad (20c)$$

$$\frac{d}{dt} B = \beta_B M_{SB} - \gamma B + \alpha_B S; \quad (20d)$$

$$\frac{d}{dt} M_A = \beta_{AP_A} \sum_k k^2 S_k + \alpha_A M_S - \gamma M_A; \quad (20e)$$

$$\frac{d}{dt} M_B = \beta_{BP_B} \sum_k k^2 S_k + \alpha_B M_S - \gamma M_B; \quad (20f)$$

$$\begin{aligned} \frac{d}{dt} M_{SA} = & -\beta_A M_{SA} - \gamma M_{SA} + \gamma(M_A - M_{BA}) \\ & + \beta_A E p_A M_{SS} - \beta_B E p_B M_{SA} - \beta_A E p_A M_{SA} \\ & + \alpha_A M_{SS} - \alpha_A M_{SA} - \alpha_B M_{SA}; \end{aligned} \quad (20g)$$

$$\begin{aligned} \frac{d}{dt} M_{SB} = & -\beta_B M_{SB} - \gamma(M_B - M_{SB}) + \gamma M_{AB} \\ & + \beta_B E p_B M_{SS} - \beta_A E p_A M_{SB} - \beta_B E p_B M_{SB} \\ & + \alpha_B M_{SS} - \alpha_B M_{SB} - \alpha_A M_{SB} \end{aligned} \quad (20h)$$

$$\begin{aligned} \frac{d}{dt} M_{AB} = & -2\gamma M_{AB} + \beta_A E p_A M_{SB} + \beta_B E p_B M_{SA} \\ & + \alpha_B M_{SA} + \alpha_A M_{SB}. \end{aligned} \quad (20i)$$

3 Comparison with stochastic simulations

Our model aims to describe, in simplified terms, the following stochastic marketing process. On a random network, each node is labeled by its buyer status: a potential buyer (or susceptible, denoted as S), a product A buyer (denoted as A), or a product B buyer (denoted as B). The status transition in the absence of word-of-mouth contact is described by the following matrix (which rows and columns are in the order of S , A , and B).

$$T = \begin{bmatrix} -\alpha_A - \alpha_B & \alpha_A & \alpha_B \\ \gamma & -\gamma & 0 \\ \gamma & 0 & -\gamma \end{bmatrix}.$$

The first row here represents the conversion from an S node to either A or B by advertisement, the second row represents that an A -node recovers (i.e., loses the buyer status) and becomes an S node. That last row represents the recovery process of a B node. Along each edge connected to an A source node (or a B source node), contact events occur as a Poisson process with rate β_A (or β_B). Upon contact, if the target node is an S node, the target is converted to A (or B). Otherwise the contact event is ignored. The contact process along an edge stops when the source node recovers.

We use the Gillespie method [1, 2] to simulate this stochastic process. The random network is generated from the configuration model as described in [6]. This model needs a given degree distribution and a given number of nodes. For each node, a random number k is drawn from the distribution and assigned as its degree. Then k stubs (half edges) are attached to the node. Two random stubs that are not from the same node or nodes that are already neighbours are randomly selected and connected to form an edge. This connection process is repeated until no such edges can be formed. Leftover stubs are discarded.

Initially, each node is initially labeled S . On a randomly generated network, 100 simulations are conducted, and their averages are compared with the solutions of our differential equation model. The degree distribution in our model is computed from the randomly generated network.

Our experiments are for Poisson networks ($P_k = \frac{\lambda^k}{k!} e^{-\lambda}$) where the expectation and variance of connections are both $\lambda > 0$. These arise in the limit $N \rightarrow \infty$ if each pair of nodes has identical probability $p = \lambda/N$ of being connected. Other relevant examples are so-called scale-free networks with $P_k = Ck^{-r}$. (C is a normalizing constant). These latter networks are

popular in epidemiological and social applications and arise when nodes are added to a growing network and new edges attach to an existing node with a probability proportional to its degree: more popular nodes attract more connections. Depending on the parameter r and the size of N , the expected number of edges and their variance can be huge. For example, with $r = 2$ and $N = \infty$ the expected value $\langle k \rangle$ is infinite!

Figure 2 shows that, on randomly generated Poisson and scale-free networks with $N = 10^4$ nodes, our model is in excellent agreement with the ensemble average of the stochastic simulations. The minimum degree of the scale-free network was chosen to be 2 to avoid isolated components.

Figure 3 shows that, on a scale-free network with exponent $r = 3$, with average degree $\langle k \rangle = 3.2$ and variance $\text{Var}[k] = 17$, the stochastic simulations behave similarly to their mean shown in Figure 2. However, on a scale-free network with exponent $r = 2$, with $\langle k \rangle = 10$ and $\text{Var}[k] = 3144$, the stochastic simulations show huge variances around their mean. We conclude that, on such a scale-free network, the ensemble mean is not a good representation of the behaviour of the stochastic marketing process. In fact, our simulations suggest that the large variance of such a network makes reliable predictions of market penetration by products almost impossible.

4 Market share

We begin with a remark on the elementary case where there is no word-of-mouth effect. In this case, the model reduces to the much simpler linear system

$$\frac{d}{dt}S = -(\alpha_A + \alpha_B)S + \gamma(A + B) \quad (21)$$

$$\frac{d}{dt}A = \alpha_A S - \gamma A \quad (22)$$

$$\frac{d}{dt}B = \alpha_B S - \gamma B \quad (23)$$

and it is straightforward to compute the asymptotic values of S, A, B as

$$S_\infty = \frac{\gamma N}{\alpha_A + \alpha_B + \gamma}, \quad A_\infty = \frac{\alpha_A N}{\alpha_A + \alpha_B + \gamma}, \quad B_\infty = \frac{\alpha_B N}{\alpha_A + \alpha_B + \gamma}. \quad (24)$$

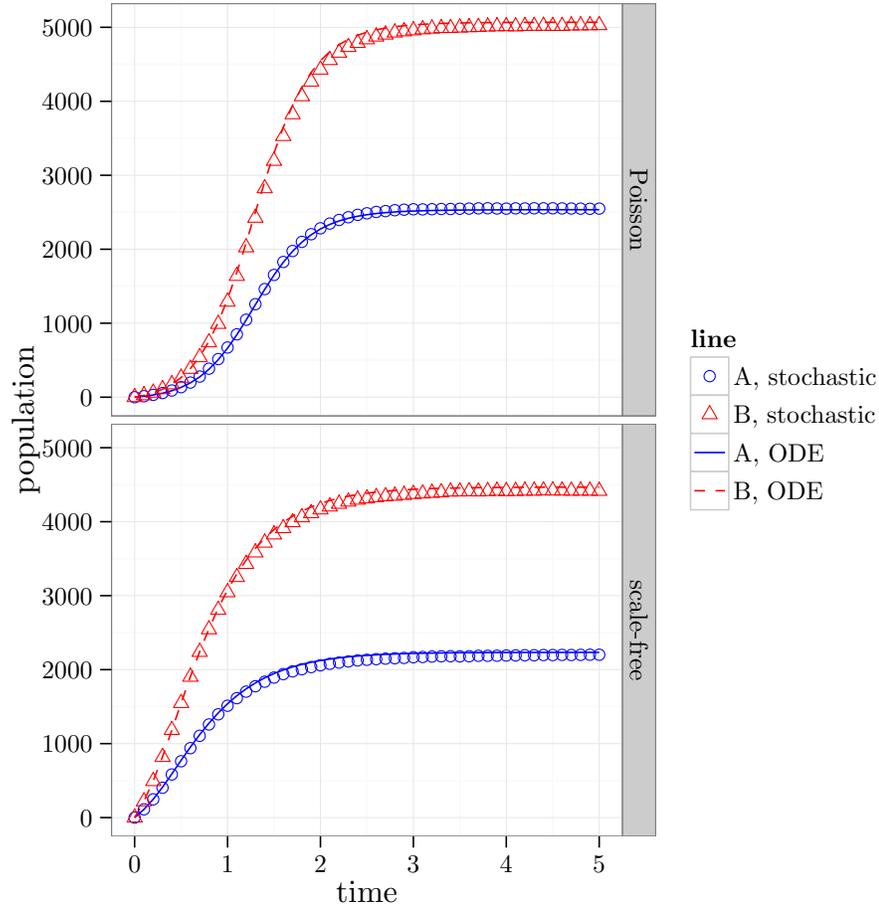


Figure 2: The comparison of the ensemble average of 100 stochastic simulations (symbols) and the solution of the ODE model (lines) on a Poisson network with average degree $\langle k \rangle = 5$, and a scale-free network with exponent $r = 3$ and minimum degree 2 (to avoid isolated pairs). Both networks have 10^4 nodes. The parameters are $\beta_A = \beta_B = 1$, $\gamma = 1$, $\alpha_A = 0.01$, $\alpha_B = 0.02$.

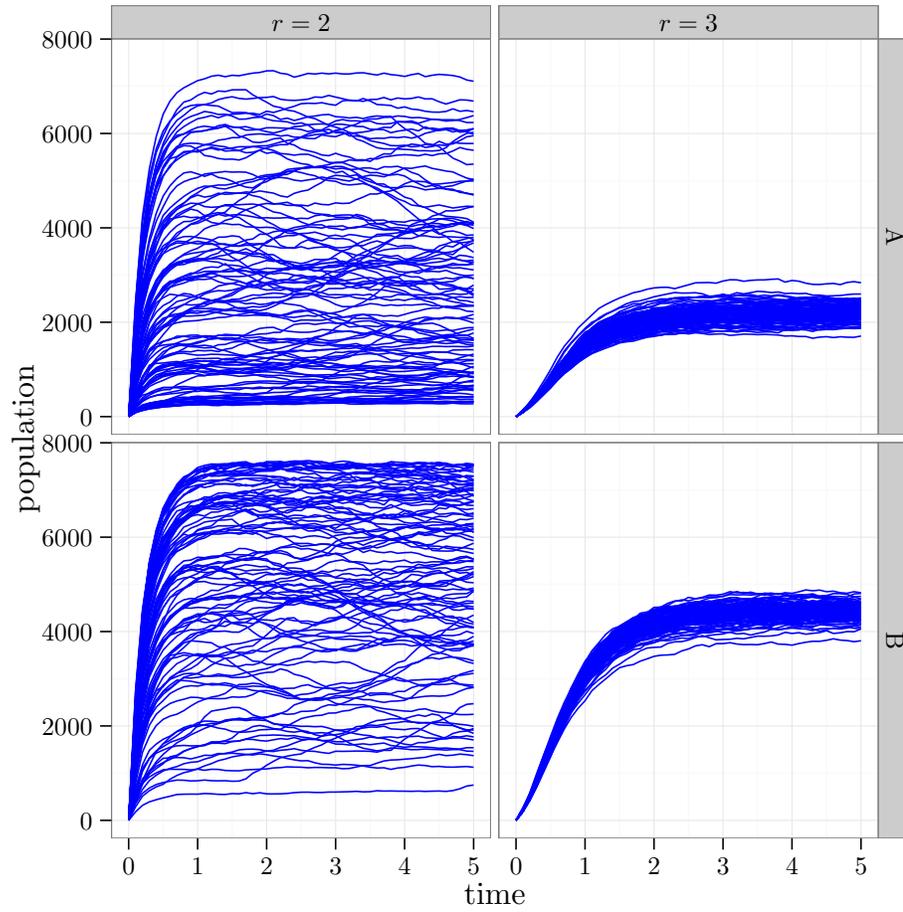


Figure 3: The comparison of 100 stochastic simulations on two scale-free networks. Both networks contain 10^4 nodes, and have minimum degree 2 to avoid isolated components. The exponents of the power-law degree distributions are $r = 2$ and $r = 3$ respectively. The marketing parameters are the same as in Figure 2.

These are stable steady solutions of the simplified model. Let $q = \alpha_B/\alpha_A$, then $B(\infty) = qA(\infty)$. In fact, since $\frac{d}{dt}(qA - B) = -\gamma(qA - B)$, starting with $A(0) = B(0) = 0$, $qA(t) = B(t)$ for all $t \geq 0$. We have similar results for the network Bass model [4], provided that $\beta_A = \beta_B$. Given that the recovery rate γ is the same for both products, our model (20) show the same behavior.

Proposition 1 *If $\beta_A = \beta_B$ and $\alpha_B = q\alpha_A$, then the solution with initial conditions $B(0) = A(0) = 0$ satisfies $B(t) = qA(t)$ and $M_{SB}(t) = qM_{SA}(t)$.*

The figures in the previous section show an example for this case ($q = 2$). The proof of the proposition is a tedious but elementary calculation, which is given in Appendix A.

Equation (24) provides a baseline for studying the effect of word-of-mouth recruitment on the equilibrium market share. For simplicity, we keep $\beta_A = \beta_B$, and so knowledge of $A(\infty)$ yields both $S(\infty)$ and $B(\infty)$ by Proposition 1 and $S(\infty) + A(\infty) + B(\infty) = N$. Figure 4 shows the influence of word-of-mouth recruitment rates with $\beta_A = \beta_B$. Not surprisingly, word-of-mouth recruitment is a significant factor in gaining market share.

Figure 5 shows the equilibrium market share of product B as a function of β_B , with a fixed β_A . We observe that the equilibrium market share of B starts with negligible values if $\beta_B = 0$, then increases slowly with β_B , and finally increases quickly as β_B grows beyond 0.5. However, because of the presence of advertisements there is no threshold phenomenon.

When β_B is small, even though the equilibrium market share of product B is small, it can still initially catch a significant market share, but then be out-competed by A . Eventually its market share declines to a small value as is illustrated in Figure 6. This is reminiscent of the competitions between Betamax and VHS video tape formats, and between HDDVD and BlueRay DVD formats, which showed a similar pattern.

5 Conclusion, remarks, and outlook

We have introduced a model for SIS-market competitions on random graphs based on node and edge dynamics. An “edge chaos” assumption is used to close the dynamic equations for the fractions of edge types. The model was tested for Poisson and scale free random networks and showed very good agreement with averaged microscopic simulations. Both analytical and

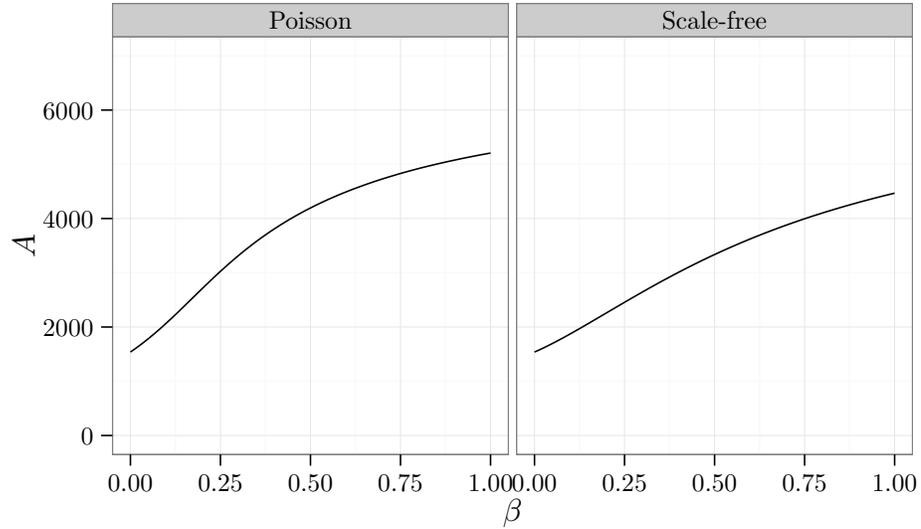


Figure 4: The dependency of the equilibrium market share of product A on β_A . Here $\beta := \beta_A = \beta_B$, $\alpha_A = 0.2$, $\alpha_B = 0.1$, $\gamma = 1$. The network is the same used in Figure 2.

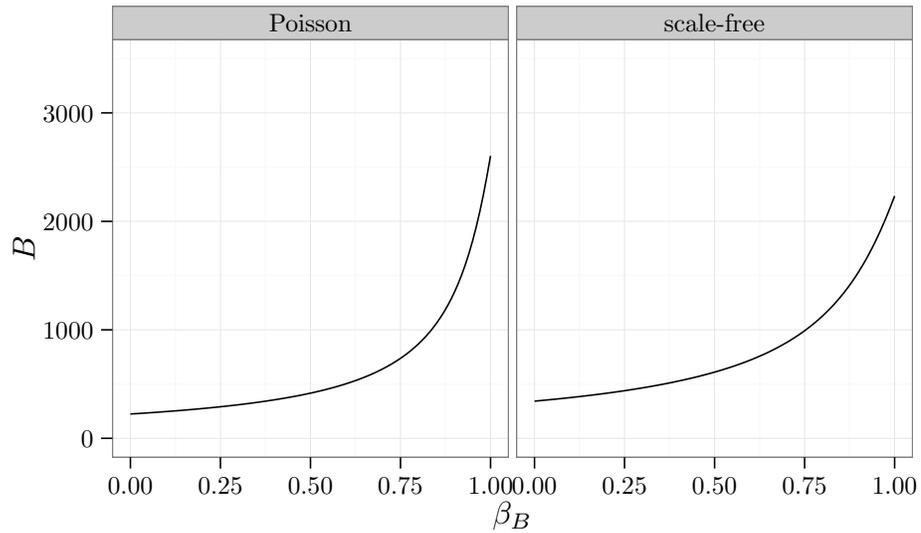


Figure 5: The dependency of the equilibrium market share of product B on β_B , with $\beta_A = 1$. The networks, α_A , α_B , and γ are the same as in Figure 4

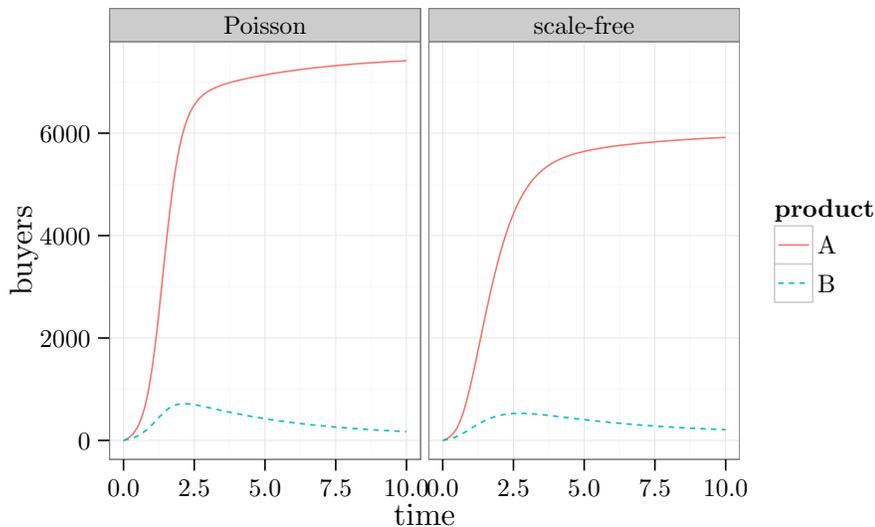


Figure 6: The time evolution of market shares of products A and B , with $\beta_A = 1$, $\beta_B = 0.7$, $\alpha_A = 0.02$, and $\alpha_B = 0.01$. The networks are the same as in Figure 6.

numerical studies on equilibrium market shares were conducted, including some predictions of final market share as a function of advertising and word-of-mouth recruitment efforts.

Apart from the initial conditions, the model equations do not distinguish between different types of random networks. This raises the question how realistic the modelling assumptions, in particular the edge chaos assumption, are for different types of networks; a relevant but probably very deep question. The simulations presented in this work suggest that the assumption is reasonable, but we considered only a few examples.

In our model (20), we assume that the recovery rates for the two products are identical. In reality, they may be different. It is straightforward to incorporate different recovery rates in our model. It may also be interesting to consider the word-of-mouth transmission rates β_A and β_B fading with time, because the interest to talk about a new product may disappear. Even though β_A and β_B are assumed constants, they represent the average transmission rates in the network. A harder but interesting question is how the variances of transmission rates affect the dynamics. More profoundly,

market dynamics may change the network itself, which may invalidate our edge chaos assumption.

As presented, our model assumes constant advertising rates. However, the number of potential buyers decreases with time. Thus advertisements becomes less efficient over time. With a limited advertising budget, a company may choose to advertise intensely to gain initial market share, then ramp down. Our model can be used to find the optimal strategy in allocating advertisement efforts.

A Proof of Proposition 1

Multiply Eq. (6) by $q = \alpha_B/\alpha_A$ and subtract from Eq. (6). With the assumption $\beta_A = \beta_B = \beta$, this gives

$$\frac{d}{dt}(B - qA) = \beta(M_{SB} - qM_{SA}) - \gamma(B - qA).$$

Thus, provided that $M_{SB} = qM_{SA}$ and $B(0) - qA(0) = 0$, this implies that $B - qA = 0$ is a solution, which is the result of the Proposition.

To show that $M_{SB} - qM_{SA} = 0$, multiply Eq. (13) by q and subtract from Eq. (14). This gives

$$\begin{aligned} \frac{d}{dt}(M_{SB} - qM_{SA}) = & -\beta(M_{SB} - qM_{SA}) - \gamma(M_{SB} - qM_{SA}) + \\ & \gamma(M_B - M_{SB} - qM_A - qM_{SA}) + \\ & \beta \frac{EM_{SS}}{M_S}(M_{SB} - qM_{SA}) - \\ & \beta \frac{E}{M_S}(M_{SB} - M_{SA})(M_{SB} - qM_{SA}) + \\ & (\alpha_B + \alpha_A)(M_{SB} - qM_{SA}) \end{aligned} \quad (25)$$

Again, provided that $M_{SB}(0) - qM_{SA}(0) = 0$, which is true given $A(0) = B(0) = 0$, $M_{SB} - qM_{SA} = 0$ is a solution provided $M_B = qM_A$.

Similarly, multiply Eq. (8) by q and subtract from Eq. (9). This gives

$$\frac{d}{dt}(M_B - qM_A) = \beta(E + 1)(M_{SB} - qM_{SA}) - \gamma(M_B - qM_A). \quad (26)$$

Note that $M_{SB} - qM_{SA} = 0$ and $M_B - qM_A = 0$ are a solution of the system (25) and (26). Thus, with the initial condition $A(0) = B(0) = 0$,

$M_{SB} - qM_{SA} = 0$ and $M_B - qM_A = 0$ hold for all time $t \geq 0$. Thus, Proposition 1 holds.

References

- [1] D. T. Gillespie. A general method for numerically simulating the stochastic time evolution of coupled chemical reactions. *J. Comput. Phys.*, 22: 403–434, 1976.
- [2] D. T. Gillespie. Exact stochastic simulation of coupled chemical reactions. *J. Phys. Chem.*, 81:2340–2361, 1977.
- [3] T. House and M. J. Keeling. Insights from unifying modern approximations to infections on networks. *Journal of The Royal Society Interface*, page rsif20100179, 2010.
- [4] M. Li, R. Edwards, R. Illner, and J. Ma. Marketing new products: Bass models on random graphs. *Commun. Math. Sci.*, 13:497–509, 2015.
- [5] J. C. Miller. A note on a paper by Erik Volz: SIR dynamics in random networks. *J. Math. Biol.*, 62:349–358, 2011.
- [6] M. Molloy and B. Reed. A critical point for random graphs with a given degree sequence. *Random Struct. Algor.*, 6:161, 1995.
- [7] M. E. J. Newman. Spread of epidemic disease on networks. *Phys. Rev. E*, 66:016128, 2002.
- [8] E. M. Volz. SIR dynamics in random networks with heterogeneous connectivity. *J. Math. Biol.*, 56:293–310, 2008.