# Time Evolution of SIS epidemics in the Complete Graph

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Delft University of Technology v1.: May 2013 v2: Oct 2016 v3: April 2017

#### Abstract

We show that, at the time being, the probability  $\Pr[M(t) = k]$  that the number of infected nodes M(t) at time t equals k in the Markovian continuous-time  $\varepsilon$ -SIS process on the complete graph cannot be determined exactly.

### 1 Introduction

In spite of the simplicity of the Markovian continuous-time SIS model, there does not seem to exist an exact time-dependent solution for any graph. Most analytic results are known for the complete graph as shown in [12, Sec. 17.6]. Before elaborating on the exact analytic solution of the Markovian continuous-time SIS model on the complete graph  $K_N$  containing N nodes, we briefly review the classical mean-field approximation.

For the complete graph  $K_N$ , mean-field approximations are accurate [3, 16]. Very likely – although there does not seem to be a rigorous proof – among all graphs, mean-field approximations are the most accurate in the complete graph. In the *N*-intertwined mean-field approximation (NIMFA) [15, 11], the governing equation for the probability v(t) of infection in a node at time t in a regular graph Gwith degree r equals

$$\frac{dv(t)}{dt} = r\beta(t)v(t)(1-v(t)) - \delta(t)v(t)$$
(1)

where the infection rate  $\beta(t)$  and the curing rate  $\delta(t)$  are general non-negative real functions of time t. The probability v(t) at time t changes due to two possible actions: (a) if the node is healthy with probability 1 - v(t), its r infected neighbors – each neighbor is infected with the same probability v(t) (due to symmetry) – can infect the node with instantaneous rate  $\beta(t)$ ; (b) when the node is infected, which happens with probability v(t), a curing processes with instantaneous rate  $\delta(t)$  can heal the node. Since the rates are time-varying, the infection and curing process are independent, inhomogeneous Poisson processes [12]. The differential equation (1) can be solved exactly [13], resulting in

$$v(t) = \frac{\exp\left(\int_0^t \left(r\beta\left(u\right) - \delta\left(u\right)\right) du\right)}{\frac{1}{v_0} + r\int_0^t \beta\left(s\right) \exp\left(\int_0^s \left(r\beta\left(u\right) - \delta\left(u\right)\right) du\right) ds}$$
(2)

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where  $v_0$  is the initial fraction of infected nodes.

As shown in [6] for regular graphs, the governing differential equations are precisely the same for NIMFA and the heterogeneous mean-field (HMF) approximation [8] of Pastor-Satorras and Vespignani. Hence, the equation (1) constitutes a general SIS mean-field approximation for regular graphs. An interesting feature of (1) is its independence on the size of the network, which avoids (or ignores) finite-size effects that often complicate studies of phase transitions. For regular graphs, the NIMFA average fraction of infected nodes y(t) = v(t) and y(t) is coined the order parameter in statistical physics. Equation (1) with constant rates,  $\beta(t) = \beta$  and  $\delta(t) = \delta$ , has been investigated earlier by Kephart and White [5]. Many variations on and extensions of the epidemic Kephart and White model have been proposed (see e.g. [9, 15, 7]). In fact, the differential equation (1) with constant rates has already appeared in earlier work before Kephart and White (see e.g. [2, 4]) and is also known as the logistic differential equation of population growth, first introduced by Verhulst [17] in 1845.

# **2** The number of infected nodes in $K_N$

We consider the time-dependent  $\varepsilon$ -SIS process on the complete graph, where a positive self-infection rate  $\varepsilon$  is crucial for the existence of a non-trivial steady state as shown in [12, Chapter 17]. The number of infected nodes M(t) at time t in the complete graph  $K_N$  is described by a continuous-time Markov process on  $\{0, 1, \ldots, N\}$  with the following rates:

$$M \mapsto M + 1$$
 at rate  $(\beta M + \varepsilon) (N - M)$   
 $M \mapsto M - 1$  at rate  $\delta M$ .

Every infected node heals with rate  $\delta$ , which explains the transition rate  $M \mapsto M - 1$ . Every healthy node (of which there are N-M at state M) has exactly M infected neighbors, each actively transferring the virus with rate  $\beta$  in addition to the self-infection rate  $\varepsilon$ . Alternatively, each of the M infected nodes can infect its N - M healthy neighbors with a rate  $\beta$  and the N - M healthy nodes can infect themselves with self-infection rate  $\varepsilon$ .

This Markov process M(t) is a birth and death process with birth rate  $\lambda_k = (\beta k + \varepsilon) (N - k)$  and death rate  $\mu_k = k\delta$  when it is in a state with M(t) = k infected nodes. The steady-state probabilities  $\pi_0, \ldots, \pi_N$ , where  $\pi_k = \lim_{t\to\infty} \Pr[M(t) = k]$ , of a general birth-death process can be computed exactly [12, p. 230],[14] as

$$\pi_k = \pi_0 \binom{N}{k} \varepsilon^* \tau^{k-1} \frac{\Gamma\left(\frac{\varepsilon^*}{\tau} + k\right)}{\Gamma\left(\frac{\varepsilon^*}{\tau} + 1\right)} \qquad (k > 0)$$
(3)

and

$$\pi_0 = \frac{1}{\sum_{k=0}^N \binom{N}{k} \tau^k \frac{\Gamma\left(\frac{\varepsilon^*}{\tau} + k\right)}{\Gamma\left(\frac{\varepsilon^*}{\tau}\right)}} \tag{4}$$

where the effective infection rate  $\tau = \frac{\beta}{\delta}$  and  $\varepsilon^* = \frac{\varepsilon}{\delta}$ . Thus,  $\pi_0$  is the steady-state probability that the complete graph  $K_N$  is infection free or overall healthy. When  $\varepsilon \to 0$  for N fixed, we observe from (3) that  $\lim_{\varepsilon \to 0} \pi_k = 0$  for k > 0 and, consequently, that  $\lim_{\varepsilon \to 0} \pi_0 = 1$ , which reflects that the steady state of the SIS process (in any finite graph) is the overall-healthy state or absorbing state.

#### **3** A generating function approach

We denote the probability that the number of infected nodes M(t) at time t equals k (or that the  $\varepsilon$ -SIS process at time t is in state k) by

$$s_k(t) = \Pr\left[M\left(t\right) = k\right] \tag{5}$$

By convention, we agree that  $s_k(t) = 0$  if k > N or if k < 0. Thus,  $s_0(t)$  is the probability that the epidemic dies out at time t or that the complete graph  $K_N$  is infection free at time t, but only remains infection free provided the self-infection rate  $\varepsilon = 0$ . Further, the steady-state probabilities

$$\pi_{k} = \lim_{t \to \infty} s_{k}\left(t\right)$$

are explicitly known in (3). The birth rate  $\lambda_k = (\beta k + \varepsilon) (N - k) = -\beta k^2 + (N\beta - \varepsilon) k + N\varepsilon$  is quadratic in k and the death rate  $\mu_k = \delta k$  is linear in k for any state  $k \in \{0, 1, 2, ..., N\}$ . The time-dependent evolution of the constant birth and death process [12, p. 239] as well as the linear birth and death process is described in [12, p. 243]. Here, we study the quadratic birth and death process, whose solution has, by the best of our efforts, not yet appeared in the literature.

Applying the differential equations of a general birth and death process to  $\varepsilon$ -SIS process yields the set

$$s_0'(t) = \delta s_1(t) - N\varepsilon s_0(t) \tag{6}$$

$$s'_{k}(t) = \left\{\beta k^{2} - \left(N\beta + \delta - \varepsilon\right)k - N\varepsilon\right\}s_{k}(t)$$
(7)

$$+\left\{-\beta\left(k-1\right)^{2}+\left(N\beta-\varepsilon\right)\left(k-1\right)+N\varepsilon\right\}s_{k-1}(t)+\delta\left(k+1\right)s_{k+1}(t)$$

where all involved rates  $\beta$ ,  $\delta$  and  $\varepsilon$  can depend upon time t. The first differential equation (6) is incorporated in the general one (7) for k = 0, since  $s_{-1}(t) = 0$  by our convention. If k = N, then  $\lambda_N = 0$  as well as  $s_{N+1}(t)$ , so that (7) reduces to

$$s'_{N}(t) = -\delta N s_{N}(t) + \{\beta (N-1) + \varepsilon\} s_{N-1}(t)$$

Since the  $\varepsilon$ -SIS epidemic must always be in one of the possible states, there holds that  $\sum_{k=0}^{N} s_k(t) = 1$ .

Following the general method illustrated in [12, Sec. 11.3.3-11.3.4] for the constant and linear rate birth and death process, we start by defining the probability generating function (pgf)

$$\varphi(x,t) = E\left[x^{M(t)}\right] = \sum_{k=0}^{N} s_k(t) x^k \tag{8}$$

which we can equally well write as  $\varphi(x,t) = \sum_{k=0}^{\infty} s_k(t) x^k$ , according to the convention that  $s_k(t) = 0$ if k > N or if k < 0. For any probability generating function  $\varphi_X(z) = E\left[z^X\right] = \sum_{k=0}^{\infty} \Pr\left[X = k\right] z^k$ , the radius R of convergence around z = 0 in the complex z-plane is at least equal to one, because for  $|z| \le 1$ , it holds that  $|\varphi_X(z)| \le \sum_{k=0}^{\infty} \Pr\left[X = k\right] |z|^k \le \sum_{k=0}^{\infty} \Pr\left[X = k\right] = \varphi_X(1) = 1$ .

**Theorem 1** In the time-dependent  $\varepsilon$ -SIS process on the complete graph  $K_N$ , the probability generating function  $\varphi(x,t)$  of the number of infected nodes M(t) at time t obeys the partial differential equation

$$\frac{\partial\varphi}{\partial t} = (x-1)\left\{-\beta x^2 \frac{\partial^2 \varphi}{\partial x^2} + \left\{\left[(N-1)\beta - \varepsilon\right]x - \delta\right\}\frac{\partial\varphi}{\partial x} + N\varepsilon\varphi\right\}$$
(9)

**Proof:** After multiplying both sides in (7) by  $x^k$  and summing over all  $k \ge 0$ , the first line in (7) is transformed as

$$T_1 = \sum_{k=0}^{N} \left\{ \beta k^2 - (N\beta + \delta - \varepsilon) k - N\varepsilon \right\} s_k(t) x^k$$

With  $\frac{\partial \varphi}{\partial x} = \sum_{k=0}^{N} k s_k(t) x^{k-1}$  and  $\frac{\partial^2 \varphi}{\partial x^2} = \sum_{k=0}^{N} k (k-1) s_k(t) x^{k-2}$ , we have

$$T_1 = \beta x^2 \frac{\partial^2 \varphi}{\partial x^2} - \left( (N-1)\beta + \delta - \varepsilon \right) x \frac{\partial \varphi}{\partial x} - N \varepsilon \varphi$$

Similarly, the transform of the second line in (7) taking our convention  $s_{-1}(t) = 0$  into account is

$$T_{2} = \sum_{k=1}^{N} \left\{ -\beta \left( k - 1 \right)^{2} + \left( N\beta - \varepsilon \right) \left( k - 1 \right) + N\varepsilon \right\} s_{k-1}(t) x^{k}$$

leading to

$$T_2 = -\beta x^3 \frac{\partial^2 \varphi}{\partial x^2} + \left( (N-1)\beta - \varepsilon \right) x^2 \frac{\partial \varphi}{\partial x} + N\varepsilon x\varphi$$

Finally, the transform of the third and last line in (7) is, with  $s_{N+1}(t) = 0$ ,

$$T_3 = \delta \sum_{k=0}^{N} (k+1) s_{k+1}(t) x^k = \delta \sum_{k=1}^{N+1} k s_k(t) x^{k-1} = \delta \sum_{k=1}^{N} k s_k(t) x^{k-1} = \delta \frac{\partial \varphi}{\partial x}$$

Equating the three right-hand side contributions  $T_1 + T_2 + T_3$  and the transform of the left-hand side in (7) yields

$$\frac{\partial\varphi}{\partial t} = -\beta x^2 \left(x-1\right) \frac{\partial^2 \varphi}{\partial x^2} + \left\{ \left(N-1\right) \beta x \left(x-1\right) - \varepsilon x \left(x-1\right) - \delta \left(x-1\right) \right\} \frac{\partial\varphi}{\partial x} + N\varepsilon \left(x-1\right) \varphi$$

Thus, we find the partial differential equation (9).

The factor (x-1) at the right-hand side of (9) is a consequence of the conservation of probability at any time t, namely that  $\varphi(1,t) = \sum_{k=0}^{\infty} s_k(t) = 1$ , implying that the  $\varepsilon$ -SIS stochastic process is surely in one of the possible states. Furthermore,  $\frac{\partial \varphi}{\partial x}\Big|_{x=1} = \sum_{k=0}^{\infty} k s_k(t)$  is the average number of infected nodes at time t. Hence, the average fraction of infected nodes at time t equals

$$y(t;\tau) = \frac{1}{N} \left. \frac{\partial \varphi(x,t)}{\partial x} \right|_{x=1}$$
(10)

Initial condition. The  $\varepsilon$ -SIS process can start with a certain probability distribution, which then requires that the initial state vector  $s(0) = (s_0(0), s_1(0), \ldots, s_N(0))$  is given. When precisely mnodes in  $K_N$  are infected initially at t = 0, then the boundary condition  $\varphi(x, 0) = \sum_{k=0}^{\infty} \delta_{km} x^k = x^m$ . Clearly, the value of m > 0 must exceed zero, because  $\varphi(0, t) = s_0(t)$  is the probability that the complete graph is infection free at time t and, on the long run,  $\lim_{t\to\infty} \varphi(0, t) = \pi_0$  is given by (4).

**Confinement.** In the sequel, we limit ourselves to constant rates: none of the infection rate  $\beta$ , self-infection rate  $\varepsilon$  or curing rate  $\delta$  is a function of time t. In addition, we assume that the  $\varepsilon$ -SIS process starts at t = 0.

### 3.1 The steady-state probability generating function $\varphi_{\infty}(x)$

The steady-state probability generating function (assuming constant rates) equals with (3)

$$\lim_{t \to \infty} \varphi(x, t) = \sum_{k=0}^{\infty} \lim_{t \to \infty} \Pr[M(t) = k] x^k = \sum_{k=0}^{\infty} \pi_k x^k = \varphi_{\infty}(x)$$

where

$$\varphi_{\infty}(x) = \pi_0 + \frac{\varepsilon^* \pi_0}{\tau \Gamma\left(\frac{\varepsilon^*}{\tau} + 1\right)} \sum_{k=1}^N \binom{N}{k} \Gamma\left(\frac{\varepsilon^*}{\tau} + k\right) (\tau x)^k \tag{11}$$

Thus, if  $\varepsilon = 0$ , then  $\pi_0 = 1$  and there holds that  $\lim_{t\to\infty} \varphi(x,t) = \varphi_{\infty}(x) = 1$ . If  $\varepsilon > 0$ , the steadystate probability generating function  $\varphi_{\infty}(x)$  is a polynomial of degree N in x, which is more elegantly written as

$$\varphi_{\infty}(x) = \frac{\pi_0}{\Gamma\left(\frac{\varepsilon^*}{\tau}\right)} \sum_{k=0}^{N} \binom{N}{k} \Gamma\left(\frac{\varepsilon^*}{\tau} + k\right) (\tau x)^k \tag{12}$$

and the general relation for any pgf,  $\varphi_{\infty}(1) = 1$ , also follows from (4). Finally,  $\varphi_{\infty}(x)$  is a function of three parameters

$$\varphi_{\infty}(x) = \varphi_{\infty}(x; \tau, \varepsilon^*, N)$$

The partial differential equation (9) simplifies, in the steady state for  $t \to \infty$  and  $\frac{\partial \varphi}{\partial t} = 0$ , to

$$-\beta x^2 \frac{\partial^2 \varphi_{\infty}}{\partial x^2} + \left\{ \left[ (N-1)\beta - \varepsilon \right] x - \delta \right\} \frac{\partial \varphi_{\infty}}{\partial x} + N \varepsilon \varphi_{\infty} = 0$$
(13)

Introducing the integral for the Gamma function  $\Gamma(s) = \int_0^\infty u^{s-1} e^{-u} du$ , valid for Re(s) > 0, into (12) yields

$$\varphi_{\infty}(x) = \frac{\pi_0}{\Gamma\left(\frac{\varepsilon^*}{\tau}\right)} \sum_{k=0}^N \binom{N}{k} (\tau x)^k \int_0^\infty u^{\frac{\varepsilon^*}{\tau} + k - 1} e^{-u} du$$
$$= \frac{\pi_0}{\Gamma\left(\frac{\varepsilon^*}{\tau}\right)} \int_0^\infty u^{\frac{\varepsilon^*}{\tau} - 1} e^{-u} \left\{ \sum_{k=0}^N \binom{N}{k} (u\tau x)^k \right\} du$$

Invoking Newton's binomium leads to an integral representation<sup>1</sup> of the steady-state probability generating function for  $\varepsilon > 0$ ,

$$\varphi_{\infty}(x;\tau,\varepsilon^*,N) = \frac{\pi_0}{\Gamma\left(\frac{\varepsilon^*}{\tau}\right)} \int_0^\infty u^{\frac{\varepsilon^*}{\tau}-1} e^{-u} \left(1+u\tau x\right)^N du \tag{15}$$

<sup>1</sup>Assuming a positive real x and letting  $w = (\tau x) u$ , we find

$$\varphi_{\infty}(x) = \frac{\pi_0}{\left(\tau x\right)^{\frac{\varepsilon^*}{\tau}} \Gamma\left(\frac{\varepsilon^*}{\tau}\right)} \int_0^\infty e^{-\frac{w}{\tau x}} w^{\frac{\varepsilon^*}{\tau}-1} \left(1+w\right)^N dw$$

We conclude that the steady-state probability generating function  $\varphi_{\infty}(x)$  can be written as

$$\varphi_{\infty}(x) = \frac{\pi_0}{(\tau x)^{\frac{\varepsilon^*}{\tau}}} U\left(\frac{\varepsilon^*}{\tau}, \frac{\varepsilon^*}{\tau} + 1 + N, \frac{1}{\tau x}\right)$$
(14)

where the confluent hypergeometric function [1, 13.2.8]

$$U(a, b, z) = \frac{1}{\Gamma(a)} \int_0^\infty e^{-zt} w^{a-1} (1+w)^{b-a-1} dt$$

is one of the independent solutions of Kummer's differential equation  $x\frac{d^2f}{dx^2} + (b-x)\frac{df}{dx} - af = 0$  (see e.g. [1, Chapter 13]).

#### **3.2** General solution of the partial differential equation (9)

**Theorem 2** In the time-dependent  $\varepsilon$ -SIS process on the complete graph  $K_N$  with constant infection rate  $\beta$ , self-infection rate  $\varepsilon$  and curing rate  $\delta$ , the probability generating function  $\varphi(x,t)$  of the number of infected nodes M(t) at time t can be written as a Laplace transform

$$\varphi(x,t) = \int_0^\infty e^{-ct} g(x;c) \, dc \tag{16}$$

where the function g(x, c) obeys the differential equation

$$-x^{2}(x-1)\frac{d^{2}g}{dx^{2}} + \left\{ \left[ (N-1) - \frac{\varepsilon^{*}}{\tau} \right] x - \frac{1}{\tau} \right\} (x-1)\frac{dg}{dx} + \frac{1}{\tau} \left( N\varepsilon^{*}(x-1) + c^{*} \right) g = 0$$
(17)

and  $\tau = \frac{\beta}{\delta}$ ,  $\varepsilon^* = \frac{\varepsilon}{\delta}$  and  $c^* = \frac{c}{\delta} \ge 0$ .

**Proof:** The usual recipe of the separation of the variables t and x, by assuming that a solution in product form as  $\varphi(x, t) = g(x) h(t)$  exists, transforms (9) to

$$\frac{\partial \log h}{\partial t} = \frac{(x-1)}{g} \left\{ -\beta x^2 \frac{d^2 g}{dx^2} + \left\{ \left[ (N-1)\beta - \varepsilon \right] x - \delta \right\} \frac{dg}{dx} + N\varepsilon g \right\}$$
(18)

By taking the derivative of both sides with respect to x, we find with  $\frac{\partial}{\partial x} \frac{\partial \log h}{\partial t} = 0$  that

$$\frac{(x-1)}{g} \left\{ -\beta x^2 \frac{d^2 g}{dx^2} + \left\{ \left[ (N-1)\beta - \varepsilon \right] x - \delta \right\} \frac{dg}{dx} + N\varepsilon g \right\} = c_1 \tag{19}$$

where  $c_1$  is a constant that is neither a function of x nor of t, because the left-hand side in (19) is independent of t. Similarly, by taking the derivative of both sides in (18) with respect to t, we find that

$$\frac{\partial \log h}{\partial t} = c_2 \tag{20}$$

and (18) shows that  $c_1 = c_2 = -c$ .

We rewrite (19) with  $\varepsilon^* = \frac{\varepsilon}{\delta}$  and  $c^* = \frac{c}{\delta}$  to find (17).

From (20), we find  $h(t) = h(0) e^{-ct}$  for the time  $t \ge 0$ . If c were complex and  $\operatorname{Im}(c) \ne 0$ , then  $h(t) = h(0) e^{-\operatorname{Re}(c)t} (\cos t \operatorname{Im}(c) + i \sin t \operatorname{Im}(c))$  and  $\varphi(x,t) = g(x) h(t)$  is generally complex for t > 0. However, the definition (8) of the pgf  $\varphi(x,t)$  illustrates that  $\varphi(x,t)$  is real for real x at any time  $t \ge 0$ . Hence, c must be real. Moreover, since the asymptotic pgf  $\lim_{t\to\infty} \varphi(x,t) = \varphi_{\infty}(x)$  exists, c must be non-negative, otherwise  $\lim_{t\to\infty} h(t) = h(0) \lim_{t\to\infty} e^{-ct} = \infty$ . We conclude that the eigenvalue c is real and non-negative.

The general solution of the eigenvalue differential equation in c consists of a linear combination  $\sum_{c\geq 0} e^{-ct}g(x;c)$  if the eigenvalues c are discrete. Generally, one readily verifies that  $\varphi(x,t) = \int_0^\infty e^{-ct}g(x;c) dc$  satisfies the partial differential equation (9) provided that g(x;c) is a solution of the differential equation (17) as a function of the "eigenvalue" c.

In fact, we need to solve an eigenvalue problem that can be expanded in a Sturm-Liouville series [10]. For c = 0, the differential equation (17) reduces to the differential (13) and we conclude that

$$g(x,0) = \varphi_{\infty}(x)$$

The  $\varepsilon$ -SIS process on the complete graph  $K_N$  with N nodes is described by a general birth-death process by the differential equations (6) and (7). This set of linear differential equations possesses a general  $(N + 1) \times (N + 1)$  tri-diagonal matrix, whose eigenstructure is studied in depth in [12, A.6.3]. The N + 1 non-negative, real eigenvalues (and one of them is zero) imply that the eigenvalues c are a discrete set  $\{c_0 = 0, c_1, \ldots, c_N\}$ , so that the Laplace integral in (16) will reduce to a sum  $\varphi(x,t) = \varphi_{\infty}(x) + \sum_{k=1}^{N} e^{-c_k t} g(x; c_k)$  for finite size N.

The second-order differential equation (17) in the function g is of the type

$$x^{2}(1-x)\frac{d^{2}g}{dx^{2}} + (ax+b)(1-x)\frac{dg}{dx} + (\lambda+d(1-x))g = 0$$
(21)

where  $a = \frac{\varepsilon^*}{\tau} - (N-1)$ ,  $b = \frac{1}{\tau}$ ,  $d = N\frac{\varepsilon^*}{\tau}$  and  $\lambda = \frac{c^*}{\tau}$  are real numbers. Unfortunately, (21) does not seem to be of a known type. Gauss's hypergeometric function F(a, b; c; x) obeys the differential equation [1, Chapter 15]

$$x(1-x)\frac{d^{2}g}{dx^{2}} + [c - (a+b+1)x]\frac{dg}{dx} - abg = 0$$

Slightly more general, (17) is of the type

$$p_3(x) g^{(2)}(x) + p_2(x) g^{(1)}(x) + p_1(x) g(x) = 0$$

where  $p_k(x)$  is a polynomial in x of degree k, where the hypergeometric differential equation is of the form

$$p_2(x) g^{(2)}(x) + p_1(x) g^{(1)}(x) + p_0(x) g(x) = 0$$

In conclusion, unless an analytic solution of the differential (21) can be found, we are afraid that the probability  $s_k(t) = \Pr[M(t) = k]$ , that the number of infected nodes M(t) at time t equals k in the Markovian continuous-time  $\varepsilon$ -SIS process on the complete graph  $K_N$ , cannot be determined exactly.

Acknowledgement. I am grateful to Johan Dubbeldam for checking the computations and for useful discussions.

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## A Reduction of the differential equation (17) to the standard form

We aim to transform (21) into the form [10]

$$\frac{d^2y}{du^2} + (\lambda - q(u)) y(u) = 0$$
(22)

The standard form has many interesting properties. First, the Wronskian is constant in u. Second, Titchmarsh [10] gives, at the beginning of the chapters, insight in the spectrum of  $\lambda$  and he also presents bounds to the solution y.

We make the transformation x = h(u), so that  $u = h^{-1}(x)$ . Thus, by using the chain rule and denoting f(u) = g(h(u)), we have

$$\frac{dg\left(x\right)}{dx} = \frac{d}{du}g\left(h\left(u\right)\right)\frac{du}{dx} = \frac{df}{du}\frac{1}{\frac{dx}{du}} = \frac{1}{h'\left(u\right)}\frac{df}{du}$$

and

$$\frac{d^2g\left(x\right)}{dx^2} = \frac{d}{dx}\left(\frac{dg\left(x\right)}{dx}\right) = \frac{d}{du}\left(\frac{dg\left(x\right)}{dx}\right)\frac{du}{dx} = \frac{1}{\left(h'\left(u\right)\right)^2}\frac{d^2f}{du^2} - \frac{h''\left(u\right)}{\left(h'\left(u\right)\right)^3}\frac{df}{du}$$

We obtain

$$\frac{h^2(1-h)}{(h'(u))^2}\frac{d^2f}{du^2} + \frac{(1-h)}{h'(u)}\left\{(ah+b) - h^2\frac{h''(u)}{(h'(u))^2}\right\}\frac{df}{du} + (\lambda + d(1-h))f = 0$$
(23)

Nex, we choose h such that  $h^2 (1-h) \frac{1}{(h'(u))^2} = 1$ . Thus,  $h^2 (1-h) = (h'(u))^2$  or  $\frac{dh}{du} = \pm h\sqrt{1-h}$  and integrated

$$\pm \int \frac{dh}{h\sqrt{1-h}} = u$$

As in Tichmarsh [10], we assume the positive sign and find

$$u = \log \frac{1 - \sqrt{1 - h}}{1 + \sqrt{1 - h}}$$

and, inversed,

$$x = h(u) = \operatorname{sech}^2\left(\frac{u}{2}\right)$$

Thus,  $x = h(u) = \operatorname{sech}^2\left(\frac{u}{2}\right)$  and  $u = 2\operatorname{ArcSech}(\sqrt{x})$ , which is only real and positive for  $x \in (0, 1)$ .

After introducing  $h(u) = \operatorname{sech}^2\left(\frac{u}{2}\right)$  into (23) yields

$$\frac{d^2f}{du^2} + \left\{\frac{\cosh u - 2}{\sinh u} - a \tanh \frac{u}{2} - \frac{b}{2} \sinh u\right\} \frac{df}{du} + \left(\lambda + d \tanh^2 \frac{u}{2}\right)f = 0$$

Let

$$r(u) = \frac{\cosh u - 2}{\sinh u} - a \tanh \frac{u}{2} - \frac{b}{2} \sinh u$$
(24)

then we obtain the differential equation in  $f(u) = g\left(\operatorname{sech}^2\left(\frac{u}{2}\right)\right)$  and  $x = \operatorname{sech}^2\left(\frac{u}{2}\right)$  or  $u = 2\operatorname{ArcSech}(\sqrt{x})$ ,

$$\frac{d^2f}{du^2} + r\left(u\right)\frac{df}{du} + \left(\lambda + d\tanh^2\frac{u}{2}\right)f = 0$$

We proceed with the reduction to the standard form by considering f(u) = p(u) s(u) and the above differential equation becomes

$$0 = p''(u) + p'(u) \left\{ 2\frac{s'(u)}{s(u)} + r(u) \right\} + p(u) \left\{ \frac{s''(u)}{s(u)} + r(u)\frac{s'(u)}{s(u)} + \left(\lambda + d\tanh^2\frac{u}{2}\right) \right\}$$

The standard form requires that  $2\frac{s'(u)}{s(u)} + r(u) = 0$ , or

$$2\frac{d}{du}\log s\left(u\right) = -r\left(u\right)$$

and

$$s(u) = \exp\left(-\frac{1}{2}\int r(u)\,du\right)$$

Explicitly, we have

$$s(u) = \exp\left(-\frac{1}{2}\int r(u)\,du\right) = \frac{\tanh u\left(\cosh\frac{u}{2}\right)^a}{\sqrt{\sinh u}}e^{\frac{b}{4}\cosh u} \tag{25}$$

From  $2\frac{s'(u)}{s(u)} + r(u) = 0$  or  $\frac{s'(u)}{s(u)} = -\frac{1}{2}r(u)$ , equivalent to 2s'(u) + s(u)r(u) = 0, we find that

$$s''(u) = -\frac{1}{2}s'(u)r(u) - \frac{1}{2}s(u)r'(u)$$

which we use in

$$X = \frac{s''(u)}{s(u)} + r(u)\frac{s'(u)}{s(u)} = -\frac{1}{4}\left\{r^2(u) + 2r'(u)\right\}$$

Hence, with s(u) in (25) and obeying  $\frac{s'(u)}{s(u)} = -\frac{1}{2}r(u)$  and with  $p(u) = \frac{f(u)}{s(u)}$ , we arrive at

$$p''(u) + p(u)\left\{\lambda + d\tanh^2\frac{u}{2} - \frac{1}{4}\left\{r^2(u) + 2r'(u)\right\}\right\} = 0$$

so that

$$q(u) = \frac{1}{4} \left\{ r^2(u) + 2r'(u) \right\} - d \tanh^2 \frac{u}{2}$$
(26)

We now compute q(u). From the definition (24) of r(u),

$$r^{2}(u) + 2r'(u) = 1 - a + a(a+1)\tanh^{2}\frac{u}{2} + b\left\{\frac{b}{4}\cosh u + \frac{b}{4} - 2\right\}(\cosh u - 1) + \frac{1}{\sinh^{2}u} + a\tanh\frac{u}{2}\left(b\sinh u - 2\frac{\cosh u - 2}{\sinh u}\right)$$

which is not such an insightfull expression!

Finally, we arrive with  $p(u) = \frac{f(u)}{s(u)}$  at the standard form

$$p''(u) + \left(\lambda + d \tanh^2 \frac{u}{2} - \frac{1}{4} \left\{ r^2(u) + 2r'(u) \right\} \right) p(u) = 0$$

Explicitly, with  $\lambda' = \lambda + \frac{a}{4} - \frac{b}{2}$ 

$$p''(u) + \left(\begin{array}{c} \lambda' + \left(d - \frac{a^2}{4} - \frac{a}{4}\right) \tanh^2 \frac{u}{2} - \frac{b^2}{16} \sinh^2 u - \frac{\cosh^2 u + 2}{4\sinh^2 u} - \frac{ab}{4} \sinh u \tanh \frac{u}{2} \\ + \frac{1}{2}a \tanh \frac{u}{2} \frac{\cosh u - 2}{\sinh u} + \frac{b}{2} \cosh u \end{array}\right) p(u) = 0$$