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Integrability of a SIS model

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Abstract

We prove that the classical model of an infectious disselve, which never kills and which does not induce autoimmunity, is integrable. This model can be written as x' = -bxy - mx + cy + mk, y' = bxy - (m + c)y with parameters $b, c, k, m \in \mathbb{R}$. We provide the explicit expression of its first integrals and of the set of all its invariant algebraic curves. © 2008 Elsevier Inc. All rights reserved.

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1. Introduction

In this paper we consider a simple epidemiological model which is a particular case of the classical SIS model introduced by Kernack and McKendrick in [7] and which is given by the following differential equation (see [2] for details):

$$x' = -bxy - mx + cy + mk, \qquad y' = bxy - (m + c)y,$$
(1)

where $b, c, k, m \in \mathbb{R}$ and the prime indicates derivative with respect to the time. We note that x = x(t) is the susceptible component of the population, y = y(t) is the infected component of the population, mk is the constant birth rate, m is the proportionate death rate, b is the infectivity coefficient of the typical Lotka–Volterra interaction term, and c is the recovery coefficient. We note that the disease is assumed to be nonfatal so that the standard term removing deceased infectaves -ay in reference [2] is omitted.

Of course the polynomial differential system (1) is defined in the whole plane \mathbb{R}^2 . We only consider the differential system (1) with $b \neq 0$ and $m \neq 0$, because if b = 0 the system is linear and it can be solved explicitly, and if m = 0, then the system has the invariant straight line y = 0 of singular points and doing a rescaling of the time variable it becomes again linear. Moreover these two cases, b = 0 or m = 0, have no interest as SIS models.

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