Some Implications of Mathematical Analyses of Epidemics

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Introduction

Mathematical analysis is a powerful tool that facilitates conceptualization and understanding of epidemics of infectious diseases [1]. The time-course of cumulative case count in epidemics is well-described by the sigmoidal shape of discrete and continuous logistic functions. For example [2], the Pearson coefficient ‘r’ for correlation between the observed cumulative sequence count and the computed values of logistic-type functions varied from a minimum of 0.9495 to a maximum of 0.9991 for pandemic influenza A (H1N1) pdm09 (pH1N1) hemagglutinin (HA) sequences collected at 23 geographic locations distributed world-wide (p<2.22e-16 at each location). A normalized version of this logistic function is given below as equation 1:

\[ Y = \frac{1}{1 + e^{-at}} \]

(1)

The fourth and fifth derivatives (\(Y''''\) and \(Y'''''\)) of \(Y\) shown in figure 1 are of the type of “jerk”, “jounce” and “swing” functions associated with turbulence of fluid flow, instability of electrical circuits and irregularity of pendulum motion [4,5]. Furthermore, it has been reported that chaotic regions have been detected in SIS ODE epidemic models [6]. The biological significance of these mathematical disturbances of evolutionary trajectory remains to be determined.

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References


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Figure 1: Time-Dependence of a Continuous Logistic-Function and its Derivatives.
Values of the logistic function (Y) and the first five of its derivatives with respect to time (Y', Y'', Y''', Y'''', Y'''''') are on the ordinates. Time is denoted on the abscissas. The function (Y) is given in equation 1. Parameter (a) was varied from 0.0 to 2.0 in increments of 0.1. Time derivatives of equation 1 were obtained with Maple 15.01 (Maplesoft, a division of Waterloo Maple, Inc.). Graphs were plotted with Python 2.7.3 (EPD 7.3-1 (64-bit)), SciPy and matplotlib.