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Backward Bifurcation in SIR endemic models

This thesis is presented in partial fulfillment of the requirements for the Degree of

AT MASSEY UNIVERSITY, ALBANY, AUCKLAND, NEW ZEALAND.

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2008

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Abstract

In the well known SIR endemic model, the infection-free steady state is globally stable for $\mathcal{R}_0 < 1$ and unstable for $\mathcal{R}_0 > 1$. Hence, we have a forward bifurcation when $\mathcal{R}_0 = 1$. When $\mathcal{R}_0 > 1$, an asymptotically stable endemic steady state exists. The basic reproduction number \mathcal{R}_0 is the main threshold bifurcation parameter used to determine the stability of steady states of SIR endemic models.

In this thesis we study extensions of the SIR endemic model for which a backward bifurcation may occur at $\mathcal{R}_0 = 1$. We investigate the biologically reasonable conditions for the change of stability. We also analyse the impact of different factors that lead to a backward bifurcation both numerically and analytically. A backward bifurcation leads to sub-critical endemic steady states and hysteresis.

We also provide a general classification of such models, using a small amplitude expansion near the bifurcation. Additionally, we present a procedure for projecting three dimensional models onto two dimensional models by applying some linear algebraic techniques. The four extensions examined are: the *SIR* model with a susceptible recovered class; nonlinear transmission; exogenous infection; and with a carrier class.

Numerous writers have mentioned that a nonlinear transmission function in relation to the infective class, can only lead to a system with an unstable endemic steady state. In spite of this we show that in a nonlinear transmission model, we have a function depending on the infectives and satisfying certain biological conditions, and leading to a sub-critical endemic equilibriums.

Acknowledgments

I would like to thank my supervisor Professor Mick Roberts for his patience, for always being there and providing many relevant suggestions and worthy opinions throughout all stages of this thesis.

I also would like to thank to the department of IIMS at the Massey University (Albany) for giving me the opportunity to pursue my postgraduate studies and wish to thank Haydn Cooper for his valuable comments.

I am also grateful to my family, my husband Arsh and my baby Talish, for without their help and support this study would have been impossible. CONTENTS CONTENTS

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