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**Predicting reintroduction outcomes
using data from multiple populations**

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Abstract

Predicting reintroduction outcomes before populations are released is inherently challenging. Reintroductions typically involve small data sets from specific locations, making it difficult to know whether results from individual case studies are more widely applicable. However, a number of species have now been reintroduced to multiple sites, providing an opportunity to move beyond the inferences possible from single-site studies. I present a novel approach where data from multiple reintroduced populations are modelled simultaneously, allowing *a priori* predictions that account for random variation among sites to be made before new reintroductions are attempted. I construct models using data from multiple reintroductions of the North Island robin (*Petroica longipes*) to identify important factors influencing population establishment, vital rates and growth across existing reintroduction sites, and use the best supported models to make predictions for a candidate reintroduction site under alternative management scenarios. My results indicate that rat tracking rate (an index of rat density) and the surrounding landscape at reintroduction sites are important for both establishment and growth of reintroduced robin populations, and that sourcing founders from habitat similar to that at the reintroduction site (forest type and predators present) is also important for post-release establishment. I then extend the multi-population approach to integrate data from multiple species, and use the resulting model to predict growth of a reintroduced population at a range of predator densities when the candidate species for reintroduction (the North Island saddleback, *Philesturnus rufusater*) has never been observed in the presence of those predators. I predict saddleback population growth at different rat tracking rates using the relationship modelled for North Island robins, with the strength of the relationship adjusted to account for the greater vulnerability of saddlebacks to predation. The relative vulnerability to predation of saddlebacks (and 24 other New Zealand forest bird species) is estimated by measuring range contraction following the arrival of introduced mammalian predators on New Zealand's mainland. My results suggest that saddlebacks could be successfully reintroduced to sites with very low rat densities. This study illustrates how an integrated approach to modelling reintroductions improves the information available to managers, providing guidance about site suitability and appropriate management measures. For species reintroduced to multiple sites, integrated models provide an ideal opportunity to develop understanding over time of the key drivers of reintroduction success.

*I dedicate this thesis to
my Mum, who never ceases to inspire
and
my Dad, whose love and support know no limits*

Preface

I have had a love of nature for as long as I can remember. I think, perhaps, it stemmed from the time I spent as a child on our family's beautiful bush-clad property on Great Barrier Island. To this day, I am filled with a sense of peace and awe whenever I am in native forest, along with a deep knowledge that this is my happy place. It was this love of the natural world that led me to do a Bachelor of Science majoring in Zoology. I still remember the day I was sitting in a lecture, listening to someone called Doug Armstrong talk about something called "reintroduction". That hour changed my life. I knew instantly that I wanted to be involved with reintroductions – they fulfilled all of my idealistic dreams of conserving native species and helping to redress the effects of humans on the world. I was sold. And so I embarked on a Masters project to study one of the first North Island robin reintroductions to a New Zealand mainland site. After completion of my Masters, I worked in conservation jobs in New Zealand and overseas; but always dreamt of doing a reintroduction-related PhD. Dream became reality when I was awarded a Doctoral scholarship from Massey University, meaning I was lucky enough to have the autonomy to develop my own research topic. I knew I wanted to contribute towards increasing reintroduction success, so I set out to develop an approach that could help us learn from past reintroductions to improve the outcomes of those carried out in the future. This was a somewhat ambitious undertaking, as I had little quantitative modelling experience; but I figured PhDs are all about learning, and modelling was the means to the end I was after. I have since come to greatly enjoy the challenge of writing code although, I must confess, not quite as much as I enjoy the challenge of finding a well hidden nest.

So here it is, after all these years, my attempt at contributing to the field of reintroduction biology. I hope that in some way, the work in this thesis will benefit reintroduced individuals and populations, and will make the hard decisions faced by the dedicated people who manage them a bit easier.

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Co-authors

While this thesis is my own original work, all four research chapters include input from one or both of my supervisors that warrants co-authorship.

Doug Armstrong (Ecology Group, Massey University)

As my primary supervisor, Doug has worked extensively with me to develop my research ideas and provide guidance on data analysis and interpretation. He reviewed all chapters (including manuscripts for publication), and has collaborated with me on every part of this research. He is consequently co-author on Chapters 2-5.

John Innes (Landcare Research)

As my secondary supervisor, John provided advice on my research concepts and his suggestions for Chapter 4, in particular, were invaluable. John reviewed Chapters 1, 4 and 6, and is a co-author on Chapter 4.

