Copyright is owned by the Author of the thesis. Permission is given for a copy to be downloaded by an individual for the purpose of research and private study only. The thesis may not be reproduced elsewhere without the permission of the Author.

## **Epidemiology of BVD in New Zealand dairy herds**

A thesis presented in partial fulfilment of the requirements for the degree of

Doctor of Philosophy

In

Veterinary epidemiology

at Massey University, Manawatu, New Zealand.

Andrew Muir Weir

2016

## Abstract

The objective of this thesis was to determine the prevalence and incidence of bovine viral diarrhoea (BVD) virus at cow and herd level, define risk factors for new infection and to quantify the impact at cow level of seroconversion during the seasonal breeding programme under the seasonal calving, pasture grazed systems in New Zealand.

A questionnaire and bulk tank milk (BTM) BVD PCR and antibody ELISA test was completed for 402 New Zealand dairy herds, and repeated in the subsequent lactation. North Island herds had a high turnover rate with 67% of virus positive herds clearing infection each lactation and being replaced with newly infected herds, while the larger South Island herds rarely cleared infection naturally (14% per lactation) and maintained a higher prevalence (32% compared to 8.5% for North Island herds). Transmission pathways associated with bulk tank BVD status were purchasing cows, neighbour's stock, and stock movements off-farm. The other factors associated with bulk tank BVD status were herd size, herd BVD vaccination, and herd ownership structure.

In 10 BTM PCR positive herds, all lactating cows (n=3,793) were tested for BVD antibody at the start of the seasonal breeding programme (planned start of mating; PSM), and again 125 days later, to identify cows that seroconverted during the observation period. Improved cutoff values were derived for the IDEXX milk antibody ELISA. There were few (3.8%) susceptible lactating cows at PSM in herds with a lactating persistently infected cow (PI), but most of these susceptible cows (82%) seroconverted. This required 4.6 contacts per PI each day. There were more susceptible (31%), and a smaller proportion of susceptible cows seroconverted (32%) in herds without a lactating PI. Seroconversion was associated with 13% longer PSM to conception (3.2 days), 4% lower pregnancy rate, 6% lower conception to AI, and \$11.97 (1.9 times) greater cost of clinical disease. The average cost per transient infection was \$91.08.

These results contributed to voluntary BVD control efforts in New Zealand and will be essential for developing a comprehensive cost-benefit model to estimate the average total cost of BVD, and assessing the benefit of various control strategies.

**Keywords**: Bovine Viral Diarrhoea; BVD; BVDV; virus; diarrhea; pestivirus; Flaviviridae; veterinary; epidemiology; New Zealand; dairy; prevalence; incidence; herd; cow; reproduction; disease; transient infection; immune suppression; PI; PCR; ELISA; antibody; milk; economic; cost; seasonal; pasture-based; observational study; longitudinal; cross-section; risk factor; risk; probability; proportion; rate; mastitis; lactation; seroconversion; regression; generalised estimating equation; GEE; Hurdle model; accelerated failure time; AFT; questionnaire; survey; sharemilker; cow behaviour; herd management; model.

## **Acknowledgements**

I would like to thank the farmers involved for their time, completing and returning questionnaires, giving permission to test their bulk milk, the use of their facilities and animals, and for following instructions and recording activities and events; New Zealand dairy farmers for funding the study through DairyNZ; The McKenzie legacy trust for additional funding contributions; Eltham District Veterinary Services for financial support, in-kind contributions (particularly Joan Hughes), and the patience and encouragement from other staff members; Livestock Improvement Corporation for in-kind contributions toward the testing costs, particularly Hinrich Voges who provided valuable feedback and advice about diagnostic testing; my supervisors Cord Heuer, Scott McDougall, and Mark Stevenson for their guidance, advice, and feedback. I would also like to acknowledge my family: my wife Kath and my children Sarah, Esther, and Daniel who sacrificed much for this PhD, and my parents Alistair and Heather Weir for their on-going support.

Approval for this research was obtained from the Massey University Animal Ethics Committee.

## **Table of Contents**

Abstract	ii
Acknowledgements	iii
Table of Contents	iv
List of Illustrations, Tables	v
Chapter 1: Introduction	1
Chapter 2: Prevalence and changes in bulk tank Bovine Viral Diarrhoea (BVD) virus status in a sample of New Zealand dairy herds.	5
Introduction	5
Materials and methods	6
Herds	6
Testing	7
Analysis	7
Results	9
Discussion	14
Chapter 3: Risk factor distribution and association with bulk tank Bovine Viral Diarrhoea (BVD) virus PCR and antibody status in New Zealand dairy herds	21
Introduction	21
Materials and methods	22
Herds	22
Testing	22
Questionnaires	23
Analysis	23
Results	24
Discussion	28
Chapter 4: Use of an Enzyme-Linked Immunosorbent Assay for detecting Bovine Viral Diarrhoea virus antibodies in individual cow milk samples	
Introduction	34
Materials and methods	35
Cows	35
Laboratory procedures	35
Statistical Analysis	35
Results	36
Discussion	39
Chapter 5: Impact of transient infection with Bovine Viral Diarrhoea (BVD) virus on reproductive performance and health of dairy cows	e
Introduction	42

Materials and methods	43
Herds	43
Testing	43
Data	44
Analysis	44
Results	
Reproduction	
Disease	50
Discussion	53
Chapter 6: General discussion	57
Conclusions	
Bibliography	
Appendix 1: Questionnaire	
Appendix 2: Statement of contribution for chapter 2	
Appendix 3: Statement of contribution for chapter 3	
Appendix 4: Statement of contribution for chapter 4	
Appendix 5: Statement of contribution for chapter 5	
List of Illustrations, Tables	
Table 1: Enrolled herds by season and district	
Table 2: Count (and row percentage) of North Island PCR positive herds in season 1 and season 2	
Table 3: Count (and row percentage) of South Island PCR positive herds in season 1 and season 2 Table 4: Count of herds in each antibody category by season with row percentage in brackets	
Figure 1: Bulk tank antibody level change between seasons with vertical lines corresponding with fir	
bulk tank antibody categories (none, low, moderate, high, and very high from left to right)	
Figure 2: Antibody level and PCR status in all herd-season bulk tank samples (sorted by antibody level horizontal lines marking the cut-off values between "none", low, medium, high, and very high a categories.	vel) with antibody
Table 5: Count of PCR negative and positive herds by antibody category with row percentage in brace	
Table 6: Comparison of status change between seasons with positive status defined as antibody S/P > PCR positive.	= 1.0, or
Table 7: Summary of selected continuous variables	
Table 8: Generalised estimating equation logistic regression model of bulk tank BVD PCR status	
Table 9: Generalised estimating equation linear regression model of bulk tank BVD ELISA sample to	-
(S/P) ratioFigure 3: Plot of IDEXX BVD milk antibody sample to positive-control ratio (S/P ratio) by serum S/	
recommended cut-off values (dark horizontal and vertical lines at 0.2 and 0.3) and adjusted cut- (grey horizontal lines at 0.08 and 0.13), the equivalency line (diagonal black 1:1 – implied by u same cut-off values for the different sample types), and the best fit regression line (grey curve, and the best fit regression line).	off values sing the
= 0.89)  Table 10: Comparison of IDEXX BVD antibody ELISA sample to positive-control optical density racategory of paired serum and milk samples using previously recommended cut-off values	ıtio

Figure 4: Bland-Altman plot showing the relationship between paired milk and serum sample to positive-control
ratio values for raw data (a), and with serum values rescaled to the milk value scale using the equation
from a linear regression model to account for the systematic difference (b). Serum was converted to the
milk value scale because that was the method used to derive new milk cut-off values. The upper and lower
dotted lines are 1.96 standard deviations away from zero mean value difference, and the vertical lines
divide the mean values into negative, suspect, and positive status (left to right) based on the previously
recommended cut-off values (a), and the adjusted cut-off values (b). Plot (a) shows that milk values tend to
be lower than serum values greater than zero. Plot (b) shows that applying the conversion accounts for this
systematic difference and brings them in line with each other with few values falling outside the 95%
interval range
Table 11: Comparison of IDEXX BVD antibody ELISA sample to positive-control ratio category of paired
serum and milk samples using new, adjusted cut-off values for the milk samples
Table 12: Number of cows, number of animals confirmed as persistently infected BVD virus carriers (PI),
number and percentage of initially susceptible cows, number of cows defined as BVD virus immune,
number of naïve, and number of transiently infected (TI) cows with percent of susceptible and percent of
herd amongst 10 dairy herds that were test positive for the presence of BVD virus in bulk tank milk 47
Table 13: Number of cows, number of animals confirmed as BVD virus carriers (PI), number and percentage of
initially susceptible cows, number of cows defined as BVD virus immune, number of naïve, and number of
transiently infected (TI) cows with percent of susceptible and percent of herd by parity group48
Figure 5: Kaplan-Meier survival curves of the time from the planned start of mating to conception for transiently
infected cows (black), and other cows (grey).
Table 14: AFT model results for the planned start of mating (PSM) to conception interval accounting for cows
clustered in herds using a log-logistic distribution
Table 15: Final pregnancy status generalised estimating equation model accounting for herd clusters with
exchangeable correlation structure
Table 16: Summary of the effect of transient infection on reproductive outcomes from logistic generalised
estimating equation models accounting for parity and calving to the planned start of mating date <sup>1</sup> 51
Table 17: Results of the hurdle model for the risk of disease and, if it occurred, the cost of disease (treatment
cost plus discarded milk cost)
Table 18: Summary of the effect of transient infection on various disease measure regression models accounting
for parity and herd53