

Modelling Animal Systems Research Paper

Cite this article: Lynch R, Kenny DA, Parr MH, Barrett D, Kelly AK, Crosson P (2018). Modelling the impact of bovine herpesvirus-1 seropositivity on the technical and economic performance of a pastoral-based suckler beef system. *The Journal of Agricultural Science* **156**, 557–564. <https://doi.org/10.1017/S0021859618000576>

Received: 23 January 2018
Revised: 28 June 2018
Accepted: 9 July 2018
First published online: 14 August 2018

Key words:

Bio-economic modelling; bovine herpesvirus 1; infectious bovine rhinotracheitis; whole-farm systems model

Author for correspondence:

R. Lynch, E-mail: richard.lynch@teagasc.ie

Modelling the impact of bovine herpesvirus-1 seropositivity on the technical and economic performance of a pastoral-based suckler beef system

R. Lynch^{1,2}, D. A. Kenny^{1,2}, M. H. Parr², D. Barrett³, A. K. Kelly¹ and P. Crosson²

¹School of Agriculture and Food Science, University College Dublin, Belfield, Dublin 4, Ireland; ²Animal & Grassland Research and Innovation Centre, Teagasc, Grange, Dunsany, County Meath, Ireland ³Department of Agriculture Food and Marine, Backweston, County Kildare, Ireland

Abstract

Bovine herpes virus 1 (BHV-1) manifests as a latent viral infection putatively affecting bovines. Understanding its effect on cattle herds is critical to maintaining sustainable beef and dairy production systems, as well as aiding in the development of herd health policies. The primary objective of the current study was, therefore, to use a whole-farm bio-economic model to evaluate the effect of herd seroprevalence to BHV-1 on the productive and economic performance of a spring calving beef cow herd. As part of a wider epidemiological study of herd pathogen status, a total of 4240 cows from 134 spring calving beef cow herds across the Republic of Ireland were blood sampled to measure the seroprevalence to BHV-1. Using data from a national breeding database, productive and reproductive performance indicators were used to parameterize a single year, static and deterministic whole-farm bio-economic model. A spring-calving, pasture-based suckler beef cow production system with an emphasis on calf-to-weanling production was simulated. The impact of BHV-1 seropositivity on whole-farm technical and economic performance was relatively small, with a marginal drop in the net margin of 4% relative to a baseline seronegative herd. Subsequent risk factors for increased pathogenicity were considered such as total herd size, percentage of intra-herd movements and vaccination status for BHV-1. In contrast to all others, scenarios representing herds that were either small in size or those which indicated an active vaccination policy for BHV-1 had no reduction in net margin against the baseline as a result of seropositivity to BHV-1.

Introduction

Bovine herpes virus 1 (BHV-1) is a latent viral infection of bovines (Nandi *et al.*, 2009). It is the causative agent of infectious bovine rhinotracheitis (IBR), which has been linked to infertility and production losses in cattle (Biuk-Rudan *et al.*, 1999; Raaperi *et al.*, 2012). Infection with BHV-1 prior to breeding can result in an animal undergoing irregular oestrus cycles (Givens, 2006), while abortifacient effects are a direct result of naïve contraction in mid-to-late gestation of cattle (Muyllkens *et al.*, 2007). The virus has also been shown to impact upon the production of viable offspring for sale as a result of immunosuppression and the onset of subsequent secondary bacterial infections (Fairbanks *et al.*, 2004; Sharon *et al.*, 2013).

Previous research has examined the prevalence of pathogens such as BHV-1 (Cowley *et al.*, 2011); however, none has investigated the implications of the pathogen on whole-farm economics. An understanding of the economic consequences of this pathogen is imperative for the development of herd health control programmes. Furthermore, government policy needs to be informed of the likely implications for national herd health. In this respect, many EU countries are currently engaged in, or have already implemented, national control programmes aimed at acquiring BHV-1-free status (Czech Republic, Germany, Italy, Denmark, Sweden, Norway, Finland, Austria, Switzerland; Cowley *et al.*, 2014). There is currently no such programme in Ireland; thus, the economic implications of exposure to BHV-1 in Irish cattle herds is of considerable interest.

Given the multifactorial nature of farm systems economics, an appropriately parameterized whole-farm model is required to establish the economic implications of an infectious disease outbreak. Mathematical models simulating farm dynamics are essential in examining how farm systems adapt to environmental changes such as the introduction of a pathogen. A number of epidemiological models have previously been developed to evaluate the economic implications of various control strategies for bovine viral diarrhoea virus (BVDv) and BHV-1 at both farm (van Schaik *et al.*, 2001) and national levels (Noordegraaf *et al.*, 1998; Gunn *et al.*, 2004). Although these models provide important information in the evaluation of a

BHV-1 infection, they lack the ability to examine its effect on whole-farm economics, due to the absence of a whole-farm modelling approach.

Furthermore, a paucity of performance data with respect to pasture-based suckler beef systems, to validate such models, may undermine their effectiveness. Previous studies such as those conducted by Noordegraaf *et al.* (1998) used a combination of expert opinion and experimental data to parameterize a state transition model for a BHV-1 infection. Where studies have used performance data, they focussed on the likely implications of a BHV-1 outbreak on dairy production systems (van Schaik *et al.*, 1999).

Therefore, a whole-farm bio-economic model incorporating the effects of BHV-1 within a beef cow herd and parameterized using novel data from a national-level animal disease study, is required. The objectives of the current paper were to:

- (1) Evaluate the impact of BHV-1 seropositivity within pasture-based suckler beef cow herds, on key economically important animal production traits.
- (2) Quantify the impact of a BHV-1 infection on whole-farm technical and economic performance using literature-sourced risk factors for this pathogen.

Materials and methods

Epidemiological study

A comprehensive epidemiological study (full title, *An integrated multi-disciplinary approach to improving the reproductive efficiency of seasonal calving beef cow herds in Ireland*; short title, *BeefCow*) was carried out to identify the key factors affecting the reproductive efficiency of commercial Irish beef cow herds, with particular emphasis on the prevalence and impact of infectious diseases (Barrett *et al.*, 2018). Information regarding seroprevalence to BHV-1, BVDv, leptospirosis and *Neospora caninum* were collected on a large cohort of breeding females. This consisted of 161 spring calving suckler beef herds, containing 6049 suckler cows. During the breeding season (May to July) in 2014 and 2015, calved cows from these farms were blood sampled to measure their seroprevalence (antibodies) to each of the pathogens using commercially available antibody test kits (BHV-1 gE, gB X3 antibody kit; Idexx Laboratories, Inc. One IDEXX Drive, Westbrook, Maine, USA). Additionally, trans-rectal uterine ultrasonography was carried out approximately 1 month after the end of the breeding season to obtain a pregnancy diagnosis. For the purposes of the current study, a sample set of beef cows from herds within the Republic of Ireland were extracted for analysis from this larger group. The sample set consisted of 134 spring calving suckler beef herds in the Republic of Ireland, containing 4240 suckler cows.

Animal-level performance data

To permit the investigation of the effects of BHV-1 seropositivity on animal performance, it was necessary to combine serology data with animal-level performance data. The animal-level performance data were retrieved from the database of the Irish Cattle Breeding Federation (ICBF), which collates data from all bovine animals in Ireland (Wickham *et al.*, 2012). Individual records were obtained for each cow serologically tested, together with performance and health-related records of their immediate progeny.

Mortality and live-weight performance traits

Mortality data from the ICBF database are categorized as mortality in the neonatal period (0–28 days of age) and mortality of older, pre-weaned calves (29–225 days of age). For the current study, these data were analysed based on the year the cow was blood tested to account for both the potential for placental–foetal transfer of the virus and potential of the calf suckling the dam contracting the virus (Fig. 1). Also, within the year of blood sampling, average daily live-weight gain was measured, up to a maximum of 225 days, which is consistent with the weaning protocol typical of Irish spring calving herds (McGee *et al.*, 2005). Live-weight gain data were obtained from livestock marts as well as on-farm weight recordings and were adjusted to account for gender and age.

Reproductive performance traits

Two parameters were used to examine the possible effects of BHV-1 on the reproductive output of beef cows (Fig. 1). Firstly, calving interval (CIV), defined as the interval in days between successive calvings, was used as an indicator of reproductive irregularities such as delayed oestrus or failure to conceive. Secondly, reappearance percentage was used to indicate the culling of non-pregnant cows due to mid- to late-term abortion and/or an extended CIV. Animals were assumed to have aborted and thus require replacement if they had a positive pregnancy diagnosis but did not reappear in the calving records of the ICBF database before 30 June of the following year. Thus, reappearance percentage was calculated as the number of cows which had a positive pregnancy scan and subsequently carried gestation successfully through to full term.

Statistical analysis

Statistical significances were obtained for the effects of seropositivity within the pre-determined risk factor groupings based on the thresholds in Table 1. Analysis was carried out using the software package Statistical Analysis Systems (SAS version 9.1.2 SAS Institute Inc., Cary NC, USA, 2004). PROC UNIVARIATE was used to confirm that all data adhered to a normal distribution, while PROC GLM was used to analyse the outcome variables of interest; CIV, reappearance percentage, live-weight gain, calf mortality and weanling mortality. Independent variables such as sero status for all other respective pathogens and body condition score of cows were included in the statistical model, along with controls for cow breed, number of cow movements, parity, herd, year, calf sire and calf sex. Significance levels were initially set at a level of $P \geq 0.20$ in order to eliminate non-significant effects using a stepwise approach; however, the final significance threshold was set at a level of $P \leq 0.05$.

Risk factors for BHV-1 pathogenicity

Statistical analysis was conducted for a range of scenarios with respect to the performance traits mentioned. Firstly, overall effects of seropositivity on performance traits were defined. Risk factors were then identified to quantify the effects on performance traits of herd seropositivity in the context of specific herd characteristics that were measurable in the present study. Van Wuijckhuise *et al.* (1998) identified herd size as an important risk factor for herd seropositivity to BHV-1 in Dutch dairy herds. The level of biosecurity has also shown to be an important management-related risk factor that could increase the pathogenicity of a BHV-1

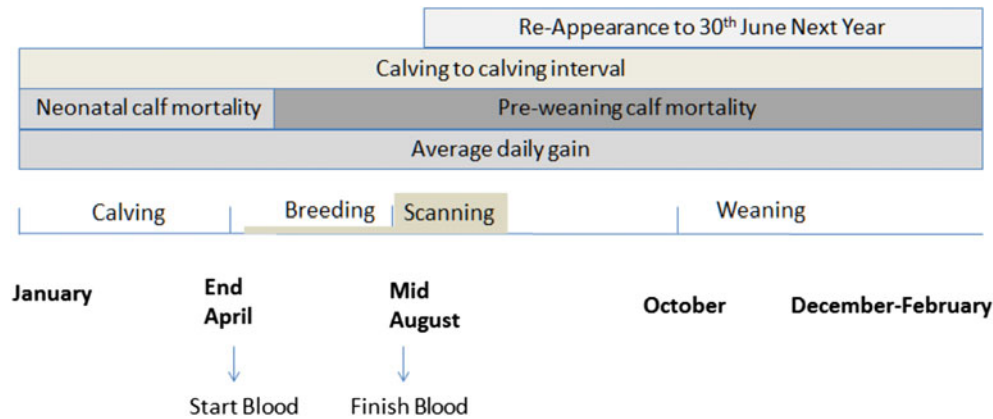


Fig. 1. Schematic representation illustrating the chronology of the BHV-1 epidemiological study and duration of measurement of each of the key performance traits. Reappearance is indicative of pregnant cows at scanning that successfully calve and thus reappear on national records, before 30 June the following year.

infection (van Schaik *et al.*, 1998; O'Grady *et al.*, 2008). In this respect, percentage of inter-herd movements was used as a measure of biosecurity within the herds sampled in the present study. Vaccination was also included as a risk factor in the current study, given its effect on the pathogenicity of the virus (Ackermann and Engels, 2006). Scenarios for each risk factor were derived from thresholds representative of the study data itself (Table 1 and Supplementary Table 1, available online at <https://www.cambridge.org/core/journals/journal-of-agricultural-science>).

Bio-economic model

General framework

The Grange Beef Systems Model (Crosson *et al.*, 2006) is a whole-farm budgetary simulation model of Irish suckler cow-based production systems. It is a single-year, static model with a monthly time step. The model is configured by specifying the farm area, proportion of the cow herd calving in each month, breeding policy (natural mating or artificial insemination (AI)), replacement rate, cattle trading strategy (month/age at sale and feed management practices) and feeding system, along with various price variables.

The feed management criteria for animal groups are based on a combination of grazing, grass silage, concentrate feeds and alternative forages (e.g. maize or whole crop cereal silage). All feeding activities are specified on a monthly basis to incorporate the seasonal variation in animal diets during the year. Forage production (herbage produced monthly expressed as kg dry matter (DM)/ha) is calculated based on conservation strategy and rate of fertilizer application. Animal feed requirements are determined according to the net energy system (Jarrige, 1989) which was modified for Irish conditions by O'Mara *et al.* (1997) and Crowley *et al.* (2002).

Costings for animal, forage and fixed items are formulated and specified within output reports. This allows for a detailed evaluation of economic performance of the farm system. Technical performance such as average animal numbers, stocking rates, live weight at key periods (e.g. weaning) and feed consumption is outlined in a summary report. Financial performance in the form of net margin includes all revenues accrued, direct and overhead costs but does not account for non-market-based subsidies (e.g. support payments made according to the European Union Common Agricultural Policy) or costs associated with the farmers own family labour or land ownership.

Parameterization of the Grange Beef Systems Model

The approach taken in the present study was to incorporate the epidemiological data from the BeefCow project into the Grange Beef Systems Model. The model was configured to represent a 40 ha spring calving suckler herd selling weanlings in the autumn. In the case of the herd size risk factor scenario, farm size was maintained at 40 ha to remove the potential confounding caused by economics of scale. Although there was no charge for owned land, where a change in farm system required additional land, this land was rented using a prevailing rental charge.

The baseline herd was assumed to be seronegative and non-vaccinating for BHV-1 and to meet industry targets for reproductive (Diskin and Kenny, 2014) and live-weight (Drennan and McGee, 2009) performance. Stocking rate was set at 2.2 live-stock units (LU) per ha; accordingly, economic performance was commensurate with the top third of Irish suckler beef herds (Teagasc, 2016). Default animal management assumptions were as follows: the calving profile modelled was 0.3, 0.4 and 0.3 of the herd calving in February, March and April, respectively, with a mean calving date of 15 March. Suckler progeny were assumed to be weaned at 225 days of age and sold as weanlings at 235 days of age. Heifers calved for the first time at 24 months of age with all females bred using AI.

The values of the performance traits in the baseline scenario were: CIV, 365 days; replacement rate, 18%; average daily live-weight gain pre-weaning for male calves, 1200 g/day and female calves 1100 g/day; neonatal mortality, 5%; and pre-weaning (excluding the neonatal period) mortality, 1%.

The effect of seropositivity within each scenario was observed as the cumulative difference between seronegative and seropositive animals for each of the performance traits, as presented in Table 2. The impact of seropositivity was thus modelled as a change in the corresponding performance traits within the baseline scenario. All animals within modelled seropositive herds were assumed to be seropositive and therefore impacted by BHV-1 seropositivity.

CIV changes were modelled as changes to the calving profile such that each additional day increase in CIV moved the calving season to later in the spring. Since reappearance percentage due to BHV-1 infection is a component of the overall herd replacement rate, its effect was modelled as an increase to the baseline replacement rate of 18%. In the case of the live-weight gain and mortality performance traits, these were modelled as increases or decreases to the values used in the baseline scenario.

Table 1. Threshold levels used to define scenarios for a whole-farm bio-economic model of BHV-1 pathogenicity^a

Risk factors	Threshold level	Category
Herd size ^b	>139 head of cattle	Large
	<72 head of cattle	Small
Movement status	>18% annual purchases	High move
	<6% annual purchases	Low move
Vaccination status	≥1 animal	Vacc
	<1 animal	Non-Vacc

^aSee supplementary material for threshold definitions.

^bBased on entire herd.

Table 2. Implications of BHV-1 seropositivity on mean values, s.e.m. (±) and *P*-values of key performance traits in pasture-based suckler beef cow herds

Variables	Serostatus		
	Sero (-)	Sero (+)	<i>P</i> -Value
Calving interval (days)	372 ± 1.2	372 ± 1.4	0.637
Live-weight gain 0–225 days (g/day)	1166 ± 12	1163 ± 14	0.850
Replacement rate (%)	12 ± 1.3	14 ± 1.5	0.126
Neonatal calf mortality <28 days (%)	4 ± 0.5	4 ± 0.6	0.829
Pre-weaning calf mortality 28–225 days (%)	1 ± 0.3	2 ± 0.4	0.321

Results

Effects of seropositivity

BHV-1 seropositivity analysis

A grand scenario was developed comparing seronegative and seropositive herd scenarios. When compared with animals which were seronegative to BHV-1, there was no significant effect of seropositivity for any of the key performance variables measured (Table 2). The greatest numerical differences were for replacement rate and pre-weaning calf mortality, which showed differences of 1.9 and 0.5%, respectively.

Technical and economic performance

When modelled at whole-farm level, the effects of seropositivity to BHV-1 were modest (Table 3). Land use and feed budgets remained broadly similar across both scenarios with no change in the quantities of grazed grass, grass silage or concentrates. Differences in replacement rates resulted in different mature and primiparous cow numbers, with corresponding effects observed in carcass and live-weight output.

Similarly, only marginal effects were found in farm financial performance, with net margin for the seronegative scenario being 4% greater than the seropositive scenario.

Effects of seropositivity within risk factor scenarios

Risk factor analysis

Further statistical analysis evaluated the effect of a change due to seropositivity in key performance traits within each of the chosen

Table 3. Impact of herd seropositivity to BHV-1 against the baseline (high performing herd, seronegative to BHV-1) on the technical and economic performance of a suckler cow herd, as modelled using the Grange Beef Systems Model (Crosson *et al.*, 2006)

	Baseline ^a	BHV-1 herd ^a
<i>Land use and feeding</i>		
Grazing area (proportion of total land area)	0.44	0.44
Grass silage area (proportion of total land area) ^b	0.56	0.56
Length of grazing season (days)	244	245
Grazed grass (tDM)	248.6	249.3
Grass silage (tDM)	92.5	92.7
Concentrates (tDM)	13.9	14.0
Inorganic N applied (kg N/ha)	139	140
<i>Livestock numbers and output</i>		
Multiparous cows at weaning	53.9	51.4
Primiparous cows at weaning	14.2	15.4
Stocking rate (LU/ha)	2.27	2.27
Live-weight output (kg/ha)	495	480
Carcass output (kg/ha)	85	94
<i>Farm economic performance (€/ha)^c</i>		
Gross output	59 800	59 600
Total variable costs	25 360	25 480
Gross margin	34 480	34 080
Net margin	14 160	13 600

^aBaseline and BHV-1 herds used the respective performance traits as follows: CIV 365, 365; ADG 1.18, 1.18; replacement rate 18.0, 19.9%; neonatal calf mortality 5.0, 4.9%; pre-weaning mortality 1.0, 1.5%.

^bGrass silage area is also available for early spring grazing and aftermath grazing following silage harvest.

^cPrices used were as follows: weaning price, €2.50/kg; beef carcass, €3.35/kg; concentrate feedstuffs, €299/tDM, inorganic fertilizer; urea, €360/t; CAN, €320/t.

risk factors (Supplementary Table S1). When compared with animals seronegative for BHV-1, seropositivity resulted in a small increase in CIV under 'Vacc', 'large' and 'high move' scenarios; however, a decline in CIV was observed for all others, most noticeably in the 'low move' scenario (Table 4). Average daily live-weight gain was not impacted by seropositivity. In contrast, seropositivity had a large effect on replacement rate for all scenarios, with the effect particularly noticeable, and opposite in its impact, for the 'large' and 'small' scenarios. The 'low move' scenario had the greatest increase in neonatal calf mortality, while the 'Vacc' and 'high move' scenarios displayed the greatest decreases. Pre-weaning mortality was reduced to a large degree in the 'small' herd scenario.

Technical performance

Whole-farm modelling showed that land use was similar across all scenarios, with a modest change in land usage seen within the 'small' herd scenario, where a larger grazing area was needed as a result of a decrease in the CIV (Table 5). The effect of a shorter CIV is to advance the mean calving date. Since cows are assumed

Table 4. Effect of seropositivity to BHV-1 on key performance traits for spring calving suckler beef herds within each of the risk factors for BHV-1 pathogenicity

Key performance traits	Baseline	Herd size		Movement status		Vaccination status	
		Large	Small	High move	Low move	Vacc ^a	Non-Vacc ^b
Calving interval (days)	365	+2.13	-1.44	+1.70	-5.27	+2.88	-1.48
Live-weight gain 0–225 days (g/day)	1180	-6	-31	-7	-44	+18	-16
Replacement rate (%)	18	+3.7	-3.6	+3.6	-0.9	-0.9	+2.6
Neonatal calf mortality <28 days (%)	5	-0.3	+0.6	-1.0	+0.9	-1.7	+0.4
Prewaning mortality 28–225 days (%)	1	+0.4	-1.1	-0.7	+0.7	+0.7	+0.5

^aVaccinating herd.^bNon-vaccinating herd.**Table 5.** Effect of seropositivity to BHV-1 on the technical performance of suckler cow systems for herds differing in size, movement status and vaccination status when compared with the baseline herd (high performing, seronegative to BHV-1)

Scenario	Baseline	Herd size		Movement status		Vaccination status	
		Large	Small	High move	Low move	Vacc	Non-Vacc
<i>Land use and feeding</i>							
Grazing area (proportion of total land area)	0.44	0.43	0.45	0.43	0.45	0.43	0.44
Grass silage area (proportion of total land area) ^a	0.56	0.57	0.55	0.57	0.55	0.57	0.56
Length of grazing season (days)	244	245	244	245	246	243	246
Grazed grass (tDM)	248.6	247.4	246.8	247.8	248.2	247.6	248.7
Grass silage (tDM)	92.5	94.7	91.5	94.2	91.0	94.0	92.7
Concentrates (tDM)	13.9	14.3	14.0	14.4	14.4	13.9	14.3
Organic N applied (kg N ha) ^b	170	170	170	170	170	170	170
Inorganic N applied (kg N ha)	139	141	136	141	137	140	140
<i>Livestock numbers and output</i>							
Multiparous cows at weaning	53.9	49.0	58.9	49.0	55.0	55.1	50.5
Primiparous cows at weaning	14.2	16.6	11.7	16.5	13.7	13.5	15.9
Stocking rate (LU/ha) ^b	2.27	2.27	2.28	2.26	2.27	2.27	2.27
Live-weight output (kg/ha)	495	466	516	475	485	513	468
Carcass output (kg/ha)	85	103	66	102	81	80	98

^aGrass silage area is also available for early spring grazing and aftermath grazing following silage harvest.^bBoth stocking rate and organic N/ha were kept at approximate equilibrium throughout all scenarios.

to be turned out to pasture post-calving, the effect of this was to increase and decrease grazed grass and grass silage demand, respectively, in the composition of the entire diet. The length of the grazing season was similar across all scenarios except vaccination status, wherein 'Vacc' scenarios were 3 days shorter when compared with 'non-Vacc'. Again, this was mostly attributable to the increase in the CIV variable seen for vaccinating herds within the study data.

There were differences in the ratio of multiparous *v.* primiparous cows across all scenarios. This was due to differences in replacement rates between scenarios. Increased replacement rate had the effect of reducing and increasing the number of multiparous and primiparous cows, respectively, at weaning. Scenarios with the highest amount of weanling sales, hence the greatest live-weight outputs, were observed for the 'small' and 'Vacc' herd scenarios at 516 and 513 kg/ha, respectively.

Economic performance

The effect of seropositivity to BHV-1 in each scenario against the baseline differed according to differences in beef price (Fig. 2). Overall, effects were minor across all the price variables (<5% deviation from the baseline), with the exception of 'low move' and 'non-Vacc' scenarios.

The lowest net margin was recorded in the 'non-Vacc' scenario with seropositivity resulting in a *c.* 10% reduction in net margin from the baseline scenario (€12 800 *v.* €14 160 per farm), when taken at a weanling price of €2.50 per kg live weight. All other scenarios, with the exception of 'small' and 'Vacc' herds, showed reduced profit due to herd seropositivity across all of the price ranges. The two highest net margins were seen in the 'small' and 'Vacc' herd scenarios at €23 588 and €24 040 per farm, respectively, when taken at a weanling price of €3.00 per kg live weight. This represented a €588 and €1040, respectively, farm

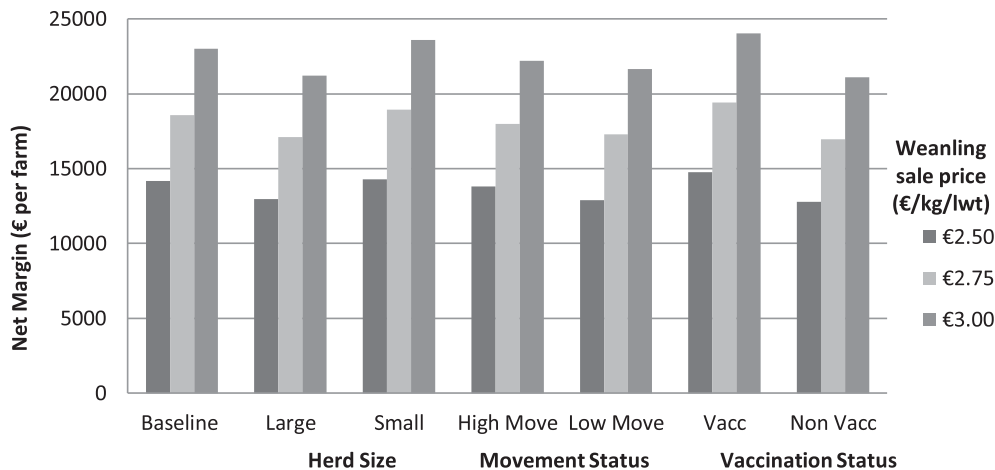


Fig. 2. Effect of weanling sale value on farm net margin for scenarios investigating the impact of seropositivity to BHV-1 within herds categorized according to the following risk factors: herd size (large and small), movement status (high move and low move) and vaccination status (Vacc and non-vacc). The baseline scenario represents a herd that is a seronegative to BHV-1 and is meeting industry targets for reproductive and live-weight performance.

net margin differential when compared with the baseline scenarios performance at a similar weanling price.

Discussion

Within pastoral-based suckler beef cow herds, cow fertility is critical to herd economic performance (Diskin and Kenny, 2014) since (1) it underpins the level of output attained with respect to the progeny produced per breeding female, and (2) it determines the capacity of the system to take advantage of grazed grass by synchronizing calving with the onset of grass availability. Further to this, healthy progeny with good live weight for day of age is essential to the economic sustainability of a calf to weanling production system. Any factor that impacts upon animal health and subsequent productivity is likely to have considerable economic implications. The primary objective of the current paper was to evaluate the effects of BHV-1 seropositivity, an infection known to have both reproductive and live-weight effects, within a pasture-based suckler beef farm. A further aim was to quantify the impact of risk factors associated with the pathogenicity of a BHV-1 infection on the technical and economic performance of a spring calving beef cow herd.

Effects of seropositivity

Within the current study, the effects of seropositivity to BHV-1 on beef cow reproduction were considered using CIV (to represent delays in rebreeding) and replacement rate (to represent potential abortifacient effects). Progeny effects considered were mortality from 0 to 28 days (neonatal) and up to 225 days (pre-weaning), in addition to live-weight gain. Overall, there was little effect of seropositivity to BHV-1 on these variables, which resulted in a negligible difference in net margin. Only a minor change was noted in CIV; however, there was an increase, albeit statistically non-significant, of *c.* 2% on replacement rate within seropositive herds. As replacement rate was determined from the rate of cow reappearance after successful conception, this is assumed to be attributed to abortion in mid-to-late gestation. Indeed, Lassen *et al.* (2012) observed similar effects, whereby herd seroprevalence of BHV-1 increased the odds ratio of abortion and still births.

No economically significant reduction in calf live-weight gain or neonatal mortality rate was associated with seropositivity. However, an increase in pre-weaning mortality further reduced profitability to an aggregated difference in farm net margin of €560 between seropositive and seronegative herds. This concurs with studies by Sharon *et al.* (2013) and Yates *et al.* (1983) that showed the ability of the virus to cause increased mortality in older animals. Such a response may be indicative of the level of exposure to environmental stressors that older stock may have endured, which generally serves to reactivate a latent virus.

Risk factor analysis

Three risk factors, which were previously identified in the scientific literature as influencing the pathogenicity of BHV-1 for seropositive herds, herd size, proportion of inter-herd movements and the herd vaccination status, were evaluated.

Herd size and herd movements

In the previous Irish work, O'Grady *et al.* (2008) indicated that herd size was a significant factor in BHV-1 seroprevalence. In a separate large-scale Estonian study by Lassen *et al.* (2012), the authors came to a similar conclusion. The current study adds more precedence by indicating a possible link between animals which had previous exposure to BHV-1 within larger herds, and the amplification of its detrimental effects on the economics of pasture-based suckler beef farms. This would tend to support the hypothesis that more horizontal spread occurred through animal-to-animal interactions within larger enterprises and therefore served to amplify the negative implications of BHV-1 seropositivity. Subsequently, this translated into a reduction in farm profitability in larger herds.

Most other related studies have focussed on the level of herd movement as an indicator for a change in herd seroprevalence (van Wuijckhuise *et al.*, 1998; Van Schaik *et al.*, 2002). The current study did not examine the change in seroprevalence directly, but rather focussed on the magnitude of the effects of the BHV-1 virus on key performance traits of interest within herds with two different levels of animal movement. 'Low move' herds had a larger reduction in net margin in comparison with 'high move'

herds, mainly due to the increase in both neonatal and pre-weaning mortality.

Vaccination status

In a review by Ackermann and Engels (2006), the authors indicated how vaccination can prevent the economic losses attributable to IBR. The availability of sufficient cohorts of both vaccinating and non-vaccinating herds within the current study allowed for the testing of a scenario based on the efficacy of vaccination at reducing the pathogenicity of BHV-1.

Net margin was not reduced within a seropositive herd with an active vaccination programme. This indicates that seropositivity for BHV-1 within herds with an active vaccination programme had less of an impact on the key performance traits, and subsequent net margin, than seropositivity within herds not practicing vaccination. The current study found there was an almost 16% difference in net margin in favour of vaccinating herds compared with non-vaccinating, after accounting for the cost associated with vaccine administration. This is largely due to the combination of a lower replacement rate and lower neonatal calf mortality and indicates clearly that herds practising vaccination observed a greater positive impact on the offspring rather than on the cow itself.

It is worth noting that most of the calf performance trait results indicate low levels of mortality and good live-weight gains, albeit with no statistical difference in most instances. However, the numerical differences seen may explain the positive net margins in small and vaccinating herds when compared with the baseline, as these data were incorporated into the model. There is a possibility that the decrease seen in 'large' and 'high move' herds with respect to calf mortality may be due to the fact that large herds or those which purchase a high number of animals annually may have a more robust overall herd health programme in place on their farm. This was further reinforced by the fact that there was a difference, by way of a reduction in calf mortality, between vaccinating and non-vaccinating herds in the current study. Another major point to note is that calf mortality pre-weaning may not have allowed for the expected effect of BHV-1 contraction to become manifest, as this excludes possible post-weaning mortality.

Strengths and limitations

The model coefficients for the current study were derived from a robust data set which originated from a comprehensive national epidemiological study carried out on BHV-1 seroprevalence within Irish suckler beef farms. The matching of these results with a range of definitive on-farm metrics from a well-established national breeding database (ICBF) allowed for a novel approach to investigating this issue.

Earlier studies have developed mathematical models to simulate the epidemiology of the onset of an endemic infection in cattle populations (Hage *et al.*, 1998; Keeling, 2005). Few studies, however, have incorporated an economic framework around such infections. Where they have (Noordegraaf *et al.*, 1998; van Schaik *et al.*, 1999, 2001), studies extended their analysis to farm-level effects such as milk yield, or risk factors for pathogenicity, independently. Furthermore, there has been no previous study of the economic effects of a BHV-1 infection in pasture-based suckler beef cow systems, using a whole-farm model. The current findings, therefore, are the first to focus on the effects of BHV-1

infection and synthesize these into a whole-farm economic modelling framework.

The major strength of this whole-farm bio-economic modelling approach is that it can account for a change in a multitude of variables on farm output simultaneously. Within the model used in the current study, adjustments to farm system variables such as calving date impact upon the feed resource allocation, while also altering the replacement rate. Subsequently, any effect on replacement rate alters the number of heifer progeny available for sale and hence farm live-weight output. Conversely, culling rates alter carcass output as was seen within 'large', 'high move' and 'Vacc' scenarios in the current study. Calf growth and mortality also impact upon the system as a whole by both decreasing the live-weight output and increasing the costs per cow calf unit, which ultimately reduces net margin.

The counterintuitive outcomes which resulted from the risk factor analysis complicated the farm-level assessment of the risk factors for BHV-1 pathogenicity. The objective of the whole-farm modelling analysis was to replicate the farm-level data as faithfully as possible – thus, the data pertaining to each scenario were used to parameterize the Grange Beef Systems Model. The very modest differences in many of the outcome variables were reflected in the negligible differences in net farm margin between scenarios.

This model provides additional information to aid the design and implementation of BHV-1 control programmes. Suckler beef herd owners will be more informed with respect to the cost-effectiveness of BHV-1 vaccination programmes based on the risk factors identified. Overall, BHV-1 has little impact on key performance traits specific to pasture-based suckler beef cow herds; however, within large herds, producers may need to be more vigilant regarding biosecurity measures on farm, one of which may be vaccination for BHV-1.

Supplementary material. The supplementary material for this article can be found at <https://doi.org/10.1017/S0021859618000576>.

Financial support. The lead author gratefully acknowledges the financial support of the Department of Agriculture, Food and the Marine under the Research Stimulus Fund (Project 13/S/515; short title, 'BeefCow').

Conflict of interest. None.

Ethical standards. Not applicable.

References

- Ackermann M and Engels M (2006) Pro and contra IBR-eradication. *Veterinary Microbiology* **113**, 293–302.
- Barrett D, Parr M, Fagan J, Johnson A, Tratalos J, Lively F, Diskin M and Kenny D (2018) Prevalence of bovine viral diarrhoea virus (BVDV), bovine herpes virus 1 (BHV 1), leptospirosis and neosporosis, and associated risk factors in 161 Irish beef herds. *BMC Veterinary Research* **14**, 8.
- Biuk-Rudan N, Cvetnić S, Madić J and Rudan D (1999) Prevalence of antibodies to IBR and BVD viruses in dairy cows with reproductive disorders. *Theriogenology* **51**, 875–881.
- Cowley DJB, Clegg TA, Doherty ML and More SJ (2011) Aspects of bovine herpesvirus-1 infection in dairy and beef herds in the Republic of Ireland. *Acta Veterinaria Scandinavica* **53**, 40.
- Cowley DJB, Graham DA, Guelbenzu M, Doherty ML and More SJ (2014) Aspects of bovine herpes virus-1 and bovine viral diarrhoea virus herd-level seroprevalence and vaccination in dairy and beef herds in Northern Ireland. *Irish Veterinary Journal* **67**, 18.
- Crosson P, O'Kiely P, O'Mara FP and Wallace M (2006) The development of a mathematical model to investigate Irish beef production systems. *Agricultural Systems* **89**, 349–370.

- Crowley AM, Keane MG, Agabriel J and O'Mara FP (2002) Prediction of net energy requirements of beef cattle. *Proceedings of Agricultural Research Forum* **11**, 19.
- Diskin MG and Kenny DA (2014) Optimising reproductive performance of beef cows and replacement heifers. *Animal* **8**, 27–39.
- Drennan MJ and McGee M (2009) Performance of spring-calving beef suckler cows and their progeny to slaughter on intensive and extensive grassland management systems. *Livestock Science* **120**, 1–12.
- Fairbanks KF, Campbell J and Chase CCL (2004) Rapid onset of protection against infectious bovine rhinotracheitis with a modified-live virus multivalent vaccine. *Veterinary Therapeutics* **5**, 17–25.
- Givens MD (2006) A clinical, evidence-based approach to infectious causes of infertility in beef cattle. *Theriogenology* **66**, 648–654.
- Gunn GJ, Stott AW and Humphry RW (2004) Modelling and costing BVD outbreaks in beef herds. *Veterinary Journal* **167**, 143–149.
- Hage JJ, Schukken YH, Dijkstra TH, Barkema HW, van Valkengoed PHR and Wentink GH (1998) Milk production and reproduction during a sub-clinical bovine herpesvirus 1 infection on a dairy farm. *Preventive Veterinary Medicine* **34**, 97–106.
- Jarrige R (1989) *Ruminant Nutrition. Recommended Allowances and Feed Tables*. London, UK: John Libbey Eurotext.
- Keeling MJ (2005) Models of foot and mouth disease. *Proceedings of the Royal Society B: Biological Sciences* **272**, 1195–1202.
- Lassen B, Orro T, Aleksejev A, Raaperi K, Järvis T and Viltrop A (2012) *Neospora caninum* in Estonian dairy herds in relation to herd size, reproduction parameters, bovine virus diarrhoea virus, and bovine herpes virus 1. *Veterinary Parasitology* **190**, 43–50.
- McGee M, Drennan MJ and Caffrey PJ (2005) Effect of suckler cow genotype on energy requirements and performance in winter and subsequently at pasture. *Irish Journal of Agricultural and Food Research* **44**, 157–171.
- Muykens B, Thiry J, Kirten P, Schynts F and Thiry E (2007) Bovine herpesvirus 1 infection and infectious bovine rhinotracheitis. *Veterinary Research* **38**, 181–209.
- Nandi S, Kumar M, Manohar M and Chauhan R (2009) Bovine herpes virus infections in cattle. *Animal Health Research Reviews* **10**, 85–98.
- Noordegraaf AV, Buijtel JAAM, Dijkhuizen AA, Franken P, Stegeman JA and Verhoeff J (1998) An epidemiological and economic simulation model to evaluate the spread and control of infectious bovine rhinotracheitis in the Netherlands. *Preventive Veterinary Medicine* **36**, 219–238.
- O'Grady L, O'Neill R, Collins DM, Clegg TA and More SJ (2008) Herd and within-herd BoHV-1 prevalence among Irish beef herds submitting bulls for entry to a performance testing station. *Irish Veterinary Journal* **61**, 809–815.
- O'Mara FP, Caffrey PJ and Drennan MJ (1997) Net energy values of grass silage determined from comparative feeding trials. *Irish Journal of Agricultural and Food Research* **36**, 110.
- Raaperi K, Bougeard S, Aleksejev A, Orro T and Viltrop A (2012) Association of herd BHV-1 seroprevalence with respiratory disease in young stock in Estonian dairy cattle. *Research in Veterinary Science* **93**, 641–648.
- Sharon KP, Duff GC, Paterson JA, Dailey JW, Carroll JA and Marceau EA (2013) *Case study: effects of timing of a modified-live respiratory viral vaccination on performance, feed intake, antibody titer response, and febrile response of beef heifers. The Professional Animal Scientist* **29**, 307–312.
- Teagasc (2016) *e-Profit Monitor Analysis: Drystock Farms 2015*. Dunsany, Ireland: Teagasc. Available at <https://www.teagasc.ie/media/website/publications/2016/eProfit-Book.pdf> (Accessed 28 June 2018).
- van Schaik G, Dijkhuizen AA, Huirne RB, Schukken YH, Nielen M and Hage HJ (1998) Risk factors for existence of bovine herpes virus 1 antibodies on nonvaccinating Dutch dairy farms. *Preventive Veterinary Medicine* **34**, 125–136.
- van Schaik G, Shoukri M, Martin SW, Schukken YH, Nielen M, Hage JJ and Dijkhuizen AA (1999) Modeling the effect of an outbreak of bovine herpesvirus type 1 on herd-level milk production of Dutch dairy farms. *Journal of Dairy Science* **82**, 944–952.
- van Schaik G, Nielen M and Dijkhuizen AA (2001) An economic model for on-farm decision support of management to prevent infectious disease introduction into dairy farms. *Preventive Veterinary Medicine* **51**, 289–305.
- van Schaik G, Schukken YH, Nielen M, Dijkhuizen AA, Barkema HW and Benedictus G (2002) Probability of and risk factors for introduction of infectious diseases into Dutch SPF dairy farms: a cohort study. *Preventive Veterinary Medicine* **54**, 279–289.
- van Wuijckhuise L, Bosch J, Franken P, Frankena K and Elbers AR (1998) Epidemiological characteristics of bovine herpesvirus 1 infections determined by bulk milk testing of all Dutch dairy herds. *The Veterinary Record* **142**, 181–184.
- Wickham BW, Amer PR, Berry DP, Burke M, Coughlan S, Cromie A, Kearney JF, McHugh N, McParland S and O'Connell K (2012) Industrial perspective: capturing the benefits of genomics to Irish cattle breeding. *Animal Production Science* **52**, 172–179.
- Yates WD, Babiuk LA and Jericho KW (1983) Viral-bacterial pneumonia in calves: duration of the interaction between bovine herpesvirus 1 and *Pasteurella haemolytica*. *Canadian Journal of Comparative Medicine* **47**, 257–264.