

FULL PAPER

Public Health

Evaluation of control measures for bovine viral diarrhea implemented in Nemuro District, Hokkaido, Japan, using a scenario tree model

Norikazu ISODA^{1,2)}, Akihiro ASANO³⁾, Michiru ICHIJO³⁾, Shiho WAKAMORI⁴⁾, Hiroshi OHNO⁵⁾, Kazuhiko SATO⁶⁾, Hirokazu OKAMOTO⁷⁾, Shigeru NAKAO⁸⁾, Hajime KATO⁸⁾, Kazuma SAITO⁹⁾, Naoki ITO¹⁰⁾, Akira USUI⁵⁾, Hiroaki TAKAYAMA³⁾ and Yoshihiro SAKODA^{2,4)}*

¹⁾Unit of Risk Analysis and Management, Research Center for Zoonosis Control, Hokkaido University, Kita 20, Nishi 10, Kita-ku, Sapporo, Hokkaido 001-0020, Japan

²⁾Global Station for Zoonosis Control, Global Institute for Collaborative Research and Education (GI-CoRE), Hokkaido University, Kita 20, Nishi 10, Kita-ku, Sapporo, Hokkaido 001-0020, Japan

³⁾Hokkaido Nemuro Livestock Hygiene Service Center, 69, Betsukai-Midorimachi, Betsukai, Notsuke-gun, Hokkaido 086-0214, Japan

⁴⁾Laboratory of Microbiology, Graduate School of Veterinary Medicine, Hokkaido University, Kita 18, Nishi 9, Kita-ku, Sapporo, Hokkaido 060-0018, Japan

⁵⁾Hokkaido Veterinary Medical Association Nemuro Branch, 119, Betsukai-Midorimachi, Betsukai, Notsuke-gun, Hokkaido 086-0292, Japan

⁶⁾Western Nemuro Operation Center, Hokkaido Higashi Agricultural Mutual Aid Association, 109, Nishi-Syunbetsu, Betsukai, Notsuke-gun, Hokkaido 088-2576, Japan

⁷⁾Northern Nemuro Operation Center, Hokkaido Higashi Agricultural Mutual Aid Association, 37, Tawara-bashi 14, Naka-shibetsu, Sibetsu-gun, Hokkaido 086-1137, Japan

⁸⁾Southern Nemuro Operation Center, Hokkaido Higashi Agricultural Mutual Aid Association, 119, Betsukai-Midorimachi, Betsukai, Notsuke-gun, Hokkaido 086-0292, Japan

⁹⁾Betsukai Town Office, 280, Betsukai-Tokiwa, Betsukai, Notsuke-gun, Hokkaido 086-0205, Japan ¹⁰⁾The Federation of Agricultural Cooperatives in Nemuro, 2, Higashi 1, Minami 1, Nakashibetsu, Sibetsu-gun, Hokkaido 086-1006, Japan

ABSTRACT. A scenario tree model was developed to propose efficient bovine viral diarrhea (BVD) control measures. The model used field data in eastern Hokkaido where the risk of BVDV infection in cattle has been reduced by an eradication program including mass vaccination, individual tests prior to communal pasture grazing, herd screening tests using bulk milk, and outbreak investigations of newly infected herds. These four activities were then used as hypothesized control measures in the simulation. In each simulation, the numbers of cattle infected persistently and transiently with BVDV detected by clinical manifestations and diagnosis tests and of missed by all of the diagnosis tests were calculated, and the numbers were used as indicators to be compared for the efficacy of the control measures. The model outputs indicated that the adoption of mass vaccination decreased the number of missed BVD cattle, although it did not increase the number of detected BVD cattle. Under implementation of mass vaccination, the efficacy of individual tests on selected 20% of the young and adult cattle was equal to that of the herd screening test performed in all the herds. When the virus prevalence or the number of sensitive animals becomes low, the efficacy of herd screening test was superior to one of individual tests. Considering the model outputs together, the scenario tree model developed in the present study was useful to compare the efficacy of the control measures for BVD.

KEY WORDS: BVD, epidemiology, scenario tree analysis

Bovine viral diarrhea (BVD) is a chronic disease with a global distribution caused by infection of bovine viral diarrhea virus (BVDV) in cattle [15], and can result in economic losses within the livestock industry [5, 6, 11, 16, 17, 21, 23, 31]. Transient

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^{*}Correspondence to: Sakoda, Y., Laboratory of Microbiology, Graduate School of Veterinary Medicine, Hokkaido University, Kita 18, Nishi 9, Kita-ku, Sapporo, Hokkaido 060-0018, Japan. e-mail: sakoda@vetmed.hokudai.ac.jp

infection (TI) of BVDV in adult ruminants shows subclinical or non-severe symptoms [8, 25]; however, BVDV viruses can be transmitted to the fetus in pregnant cattle during the 40–120 days of gestation, thereby inducing fetus immune tolerance to BVDV and resulting in the delivery of persistently infected (PI) calves [3, 9–10, 17, 22, 26]. The PI calves shed BVDVs through oculonasal discharge and other secretions throughout their lives, without showing apparent clinical manifestations except for fatal mucosal disease [7, 12]. It is commonly believed that PI cattle are the primary source of BVDV infection in a herd [17, 27]. Therefore, it is essential for the control of BVD that PI cattle are promptly detected and culled [1].

BVDVs have been detected in Japan since 1969. After the designation of BVD as a notifiable disease in 1998, the numbers of BVD diagnosed cases stabilized between 100 and 200 animals per year [19]. According to the results of the latest survey on PI animals in dairy farms in Japan, the proportion of farms with PI animals and that of cattle diagnosed as PI animals were 7.6% (95% confidence limits [CI]: 3.1–16.4%) and 0.12% (95% CI: 0.05–0.25%), respectively [19]. The majority of dairy cattle in Japan are raised in Hokkaido Prefecture, which is an island located in the northern part of Japan. As dairy husbandry is one of the main industries in the eastern part of Hokkaido, some husbandry communities have implemented independent control measures for BVD virus infection in cattle. In Betsukai Town, Nemuro District, mass vaccination using a combination of inactivated and attenuated vaccines and screening tests using bulk tank milk followed by individual serum tests have been implemented since 2006 [18]. These independent but well-designed control measures reduced the numbers of PI detected by active surveillance from 21 in 2006 to 2 in 2012 [29]. As a result of these intensive control measures, the risk of BVDV infection in this area was reduced dramatically; however, the sustainability of these control measures should be a focus so as to prevent the BVDV endemic in this area.

As the ecology of BVDV in cattle is complicated and some of the interventions are only effective over specific limited conditions, there is no single effective intervention to control BVD that is commonly used worldwide. A scenario tree model is a useful risk assessment model to conduct quantitative assessment so as to identify an effective control measure or a combination of measures that could be used concurrently or in rotation. A scenario tree model of the surveillance program was developed to estimate the probability of freedom from BVDV infection in Norway, with the model results validated by the surveillance program undertaken with a high probability of disease freedom [28]. Another scenario tree model was developed to estimate the probability of the introduction of BVDV to Danish dairy herds by import or transfer or to identify effective interventions to minimize the risk of BVD introduction [14].

In the present study, a stochastic scenario tree model was developed using the field data relating to husbandry and epidemiology of BVD in Nemuro, Hokkaido, Japan, to evaluate each of the targeted interventions which had been implemented under the estimation of a BVD epidemic situation. The outputs obtained from the model indicated that implementation of mass vaccination should definitely be effective for reducing BVDV in the area, and efficacies of disease monitoring interventions would be influenced by the epidemiological situations, including vaccination coverages and BVDV prevalence.

MATERIALS AND METHODS

Model structure

To estimate the number of detected and missed cattle infected with BVDV as transient and persistent infection, we developed a stochastic scenario tree model composed of nodes representing cattle categorization, immunity, virus infection, and monitoring based on implementation in a certain area in Nemuro District, Hokkaido Prefecture, where the cattle infected with BVDV had been reduced by the interventions [29]. A scenario tree model for risk assessment is generally composed of three types of nodes, namely category nodes, infectious nodes, and detection nodes, so as to theoretically calculate the probabilities of infected animals being detected at a given level of prevalence. To generalize the output of this model, the number of initial cattle was set as tens of thousands instead of the actual number of cattle in Nemuro. The numbers of either persistent or transient infection with BVDV in each animal category were calculated as a product of all conditional probabilities describing the nodes of each specific categorization and infection pathway. The overall probabilities of detecting and missing cattle persistently or transiently infected with BVDV (BVD cattle) through the monitoring nodes as interventions for BVD control in a default or scenario settings were calculated as a product of the probabilities of infected and missed BVD cattle were calculated as an output of the default or scenarios and were compared to evaluate the efficacy of the interventions in the scenarios. The scenario tree was developed in an Excel spreadsheet (Microsoft Office Excel 2013), using the software @Risk 6 (Palisade Corporation, Ithaca, NY, U.S.A.), and was run using 10,000 iterations.

Data

A basic animal and epidemiological dataset was kindly provided by the Federation of Agricultural Cooperatives in Nemuro, the Hokkaido Nemuro Livestock Hygiene Service Center, and the Betsukai town office. Epidemiological, immunological, and virological data, as well as information relating to the efficacies of vaccination in cattle and sensitivities of diagnosis tests were sourced from peer-reviewed scientific papers. Some of the epidemiological parameters used in the present assessment were kindly advised by a total of 15 specialists of the Eastern Hokkaido Agricultural Mutual Aid Association and Hokkaido Nemuro Livestock Hygiene Service Center, since there were no data available from the literature. All of the data used in the present scenario tree model were shown in Table 1.

Table 1. Parameters used in the scenario tree analysis

Parameter	Name	Default	Source of information
N	Number of animals in a modeled area	10,000	Decided by the authors
R _a	Rate of each age group	$\begin{array}{l} 0-2 \ M \ old \ (R_{a0-2}): RiskPert \ (0.0346, \ 0.0372, \ 0.0398) \\ 3-8 \ M \ old \ (R_{a3-8}): RiskPert \ (0.1017, \ 0.1021, \ 0.1025) \\ 9-14 \ M \ old \ (R_{a9-14}): RiskPert \ (0.0825, \ 0.0903, \ 0.0981) \\ 15-18 \ M \ old \ (R_{a15-18}): RiskPert \ (0.0536, \ 0.0556, \ 0.0576) \\ > 19 \ M \ old \ (R_{a} > 19): \ 1-R_{a0-2}-R_{a3-8}-R_{a8-14}-R_{a15-18} \end{array}$	Federation of Agricultural Cooperatives in Nemuro
P _p	Probabilily of persistant infection	0.4%: RiskPert (0.003, 0.004, 0.006) 0.13%: RiskPert (0.0005, 0.0012, 0.0025) 0.04%: RiskPert (0.0003, 0.0004, 0.0005)	Damman et al., 2015 [13]
P _{vs}	Probability of vaccine administration under single-administration of vaccination program at each age group	0–2 M (P _{vs0-2}): 0 3–8 M (P _{vs3-8}): Riskuniform (0.95, 1.00) 9–14 M (P _{vs9-14}): Riskuniform (0.975, 1.00) 15–18 M (P _{vs15-18}): Riskuniform (0.975, 1.00) >19 M (P _{vs>19}): Riskuniform (0.975, 1.00)	Saino <i>et al.</i> , 2013 [29]
P _{vd}	Probability of vaccine administration under double-administrations of vaccination program at each age group	0–2 M (P _{vd0-2}): 0 3–8 M (P _{vd3-8}): Riskuniform (0.95, 1.00) 9–14 M (P _{vd9-14}): Riskuniform (0.95, 1.00) 15–18 M (P _{vd15-18}): Riskuniform (0.975, 1.00) >19 M (P _{vd >19}): Riskuniform (0.975, 1.00)	Saino <i>et al.</i> , 2013 [29]
P _{vt}	Probability of vaccine administration under triple-administrations of vaccination program at each age group	0–2 M (P _{vt0-2}): 0 3–8 M (P _{vt3-8}): Riskuniform (0.95, 1.00) 9–14 M (P _{vt9-14}): Riskuniform (0.95, 1.00) 15–18 M (P _{vt15-18}): Riskuniform (0.65, 0.75) >19 M (P _{vt15-19}): Riskuniform (0.975, 1.00)	Saino <i>et al.</i> , 2013 [29]
P _{il}	Probability of conferring immunity for protection against BVDV infection under single-administration of vaccination program at each age group	0–2 M: 0 3–8 M, 9–14 M, 15–18 M, >19 M (Pi1f): RiskBeta (70, 30)	Saino et al., 2013 [29]
P _{i2}	Probability of conferring immunity for protection against BVDV infection under double-administrations of vaccination program at each age group	0–2 M: 0 3–8 M (P _{i2t}): RiskBeta (70, 30) 9–14 M, 15–18 M, >19 M (P _{i2s}): RiskPert (0.85, 0.90, 0.95)	Saino et al., 2013 [29]
P _{i3}	Probability of conferring immunity for protection against BVDV infection under triple-administrations of vaccination program at each age group	0–2 M: 0 3–8 M (P _{i3f}): RiskBeta (70, 30) 9–14 M (P _{i3s}): RiskPert (0.85, 0.90, 0.95) 15–18 M, >19 M (P _{i3t}): RiskPert (0.975, 0.99, 1.00)	Saino et al., 2013 [29]
\mathbf{P}_{t}	Probability of Transient infection with BVDV	RiskUniform (0.0015, 0.0026)	Damman et al., 2015 [13], modified
P _{mp}	Probability that persistantly infected cattle is detected by clinical manifestation	0–2 M: RiskBeta (16,4) 3–8 M: RiskBeta (8,12) 9–14 M: RiskBeta (4,16) 15–18 M: 0.001 >19 M: RiskBeta (12,8)	Opinions from veterinarians in Nemuro District
P _{mt}	Probability that transiently infected cattle is detected by clinical manifestation	0–2 M, 3–8 M, 9–14 M, 15–18 M: 0 >19 M: 0.3	Opinions from veterinarians in Nemuro District
P _c	Probability of using communal summer pasture	Pasture (10%): 0.1 Pasture (20%): 0.2 Pasture (30%): 0.3 Pasture (40%): 0.4 Pasture (50%): 0.5	Set in the present study for scenario analysis
\mathbf{S}_{ec}	Test sensitivity of individual test prior to using communal pasture	RiskBeta (19,1)	Kim and Dubovi, 2003 [20]
S _{eb}	Test sensitivity of screening test using bulk tank milk sample	RiskBeta (19,1)	Kim and Dubovi, 2003 [20]
Po	Probability of performing outbreak investigation of newly infected herd	RiskPert (0.0015, 0.003, 0.0045)	Federation of Agricultural Cooperatives in Nemuro
S_{eo}	Test sensitivity of outbreak investigation	RiskBeta (19,1)	Kim and Dubovi, 2003 [20]
P _{ma}	Probability of maternal antibody being present in the 3–8 M cattle	Under single-administraion (P _{mi1}): RiskBeta (60,40) Under double-administrations (P _{mi2}): RiskBeta (33,7) Under triple-administrations (P _{mi3}): RiskBeta (18.5,1.5)	Opinions from veterinarians in Nemuro District



Fig. 1. Overview of the scenario tree developed in the present study. A stochastic scenario tree is composed of nodes representing cattle categorization, vaccination program, virus infection, and monitoring based on implementation in Nemuro District. This tree represents the disease detection dynamics of over 19-months old cattle. Nodes with superscripts of a, b and c indicate that node is included only on the tree of 3–8 months group, nodes are removed on the tree of 0–2 months group, and nodes are included only on the tree of >19 months group, respectively. All of the parameters in along with branches are referred from the Table 1.

Structure of the scenario tree model

We developed a scenario tree to represent the distribution of animals, persistent or transient infection with BVDV in cattle, immunity to BVDV infection, and detection of BVD cattle (Fig. 1).

Animals: All of the 10,000 cattle were divided into five age groups: 1) 0–2 months (0–2 M); 2) 3–8 months (3–8 M); 3) 9–14 months (9–14 M); 4) 15–18 months (15–18 M); and 5) >19 months (>19 M). Cattle of 0–2 M were assumed to potentially be PI with BVDV but not TI due to possessing maternal antibodies, and were assumed not to use the communal summer pastures. On the other hand, some of the 3–8 M cattle were considered to potentially be TI with BVDV due to a decline of the maternal antibodies to a level that is insufficient for protection. As field data to reveal the duration of maternal antibodies in the calves are scarce, the probability of maternal antibodies being present in the 3–8 M cattle (P_{ma}) was defined according to the opinions of local veterinarians in Nemuro. In parallel, 3–8 M cattle were set to be injected with the first administration of the vaccine under the adoption of mass vaccination programs, and could use the communal summer pastures. Cattle of 9–14 M were assumed to no longer possess maternal antibodies, and were assumed to be injected with the second or third administration of the vaccine under the adoption of mass vaccination programs. Cattle of 15–18 M were assumed to be injected with the second or third administration of the vaccine under the adoption of mass vaccination programs. Cattle of 15–18 M were assumed to be injected with the second or third administration of the vaccine under the adoption of mass vaccination programs. Rates of each of the five age distributions (R_a) were determined by the data of cattle age distribution in Nemuro.

Persistent Infection: The probability of cattle being PI with BVDV was set as P_p , and the default was determined as 0.4% according to the field data in Hokkaido [2].

Immunity by vaccination: In the default setting, none of the cattle were set to be injected with vaccines and to possess sufficient maternal antibodies for protection against BVDV infection. On the other hand, mass vaccination programs in the scenario settings were set as control measure options by referring to the current vaccination program implemented in Nemuro [29]. The probabilities of possessing insufficient immunity for protection against BVDV infection, even after receiving the vaccine injection in each of four age groups, were set as in Table 2.

Transient infection: Cattle which did not receive any vaccine administrations or those that had insufficient immunity for protection against BVDV infection, even after receiving vaccine administrations, were considered as potential targets to become

Vaccine	Probability of insufficient immunity enough to protect in each of age group					
programme	0–2 M	3–8 M	9–14 M	15–18 M	>19 M	
Single-shot	0	$P_{v1 (3-8)}*(1-P_{i1f})*(1-P_{mi1})$	$P_{v1 (9-14)} * (1-P_{i1f})$	$P_{v1 (15-18)} * (1-P_{i1f})$	$P_{v1(19-)}*(1-P_{i1f})$	
Double-shots	0	$P_{v2(3-8)}*(1-P_{i2f})*(1-P_{mi2})$	$\begin{array}{c} P_{v2\;(9-14)}*(1{-}P_{i2s}){+}\\ (1{-}P_{v2\;(9-14)})*P_{v2\;(3-8)}*(1{-}P_{i2f})\end{array}$	$P_{v2(15-18)}*(1-P_{i2s})$	$P_{v2(19-)}*(1-P_{i2s})$	
Triple-shots	0	$P_{v3(3-8)}*(1-P_{i3f})*(1-P_{mi3})$	$\begin{array}{c} P_{v3}_{(9-14)}*(1-P_{i3s})+\\ (1-P_{v3}_{(9-14)})*P_{v3}_{(3-8)}*(1-P_{i3f})\end{array}$	$\begin{array}{c} P_{v3\;(15-18)}*(1-P_{i3t})+\\ (1-P_{v3\;(15-18)})*P_{v3\;(9-14)}*(1-P_{i3s})\end{array}$	$P_{v3(19-)}*(1-P_{i3t})$	

Table 2. Probability of non-inducement of enough immunity to protect with vaccine injection

Each of the abbreviations are referred in Table 1.

TI cattle. In non-vaccinated cattle and insufficiently immunized cattle, TI cattle would be produced with a probability of transient infection with BVDV (P_t). As there were few field data to verify P_t , we referred to the simulated results of probability of transient infection by Damman *et al.* [13]. P_t was considered to be common in all of the age groups, but would be affected by the P_p , which was shown in the simulation analysis in Damman *et al.* P_t was set as 25% of P_p ; thus, the default P_t was defined as 0.1%.

Clinical manifestations: All of the probabilities of showing clinical manifestations of BVD were defined according to the opinions of field veterinarians in Nemuro. PI cattle of 0–2 M were assumed to be detected by apparent clinical manifestations of insufficient growth. In the present model, the probability that 0–2 M PI cattle being detected by clinical manifestations, $P_{mp (0-2)}$, was set to approximately 80%. In parallel, a BVD cow that delivered its calf diagnosed as PI should be detected. The probability that >19 M PI cattle would be detected by clinical manifestations, $P_{mp (>19)}$, was set to approximately 80%, with a hypothesis that 83% of the >19 M cattle deliver calves in a year. Although most of the PI calves were assumed to be identified during 0–2 M, some of the PI calves would be detected at a later phase, such as 3–8 or 9–14 M as insufficiently grown cattle. The probabilities that 3–8 and 9–14 M PI cattle would be detected by clinical manifestations, $P_{mp (3-8)}$ and $P_{mp (9-14)}$, respectively, were set to approximately 40% and 20%, respectively. On the other hand, PI cattle of 15–18 M were assumed to be rarely detected by clinical manifestations since the older the PI cattle, the fewer signs they are expected to show, besides reproductive disorder. Transient infection of BVDV in cattle is mainly subclinical or clinical over a short period. In the present model, TI cattle from the >19 M group would only be detected by reproductive disorder with a probability of 30%, which was derived from the rates of calving in delivered cattle, and delivery of the PI calf as an immature fetus would be caused by BVDV infection over 33% of the gestation period. We did not assume the presence of other reproductive disorders, including conception failure and abortion, as these are also caused by other infectious diseases, or environmental factors.

BVD cattle diagnoses: Diagnosis systems for cattle infected with BVDV in the present model were referred to the control measures implemented in Nemuro, including individual tests prior to communal pasture grazing, the herd screening test using bulk milk, and the outbreak investigation of newly infected herds. In Nemuro, all of the cattle kept at the summer communal pasture are mandatorily investigated for BVDV infection by RT-PCR method using serum samples prior to communal pasture grazing [29]. In the present model, the cattle older than three months were assumed to possibly use the summer communal pastures with a certain probability (P_c), which was changed from 10 to 50% in the scenario models. In Nemuro, independent of the use of summer communal pasture, all of the lactating cows are mandatorily screened for BVDV infection by RT-PCR method using bulk tank milk samples. As this screening test is usually performed over six month intervals during every spring and autumn, cows that were not lactating as dry cows at the time of the initial screening test would be investigated during the latter screening test scheme; therefore, theoretically, all of the delivered cattle would be screened no less than once per year. In the present model, PI or TI cattle in the >19 M group were estimated to be screened with a certain test sensitivity (S_{eb}) using bulk tank milk once a year. In the default setting, bulk milk tests were not performed (Seb was zero), and this was changed to approximately 95% in the scenario models. Under Japanese legislation, when any cattle in a herd are diagnosed with BVD, all of the cattle in the herd are mandatorily tested for BVDV infection either by RT-PCR or virus isolation method using serum samples as an outbreak investigation of a newly infected herd. In the present model, all of the cattle except those in which BVD was detected by any clinical manifestation, by the individual compulsory test, and by the screening test, were assumed to be tested within the outbreak investigation with a probability of 0.3% (P_o), which was calculated by the number of infected herds to that of all the herds in Nemuro. Test sensitivities of BVDV detection from serum and bulk milk samples were as according to the calculations by Kim and Dubovi [20].

Evaluation of interventions in the scenarios

To evaluate the efficacy of adoption of mass vaccination programs, the scenario models with single-, double-, and tripleadministrations of vaccination program were run, and the number of detected BVD cattle either by clinical manifestations or outbreak investigation, and the number of missed BVD cattle were calculated.

To evaluate the efficacy of diagnosis schemes by individual mandatory tests and the herd screening tests, the scenario models were run with the use of communal pasture at a rate of 10–50%, and/or with implementation of bulk milk tests with a test sensitivity of S_{eb} .

Phase one scenario

To evaluate the efficacy of the diagnosis scheme by the individual mandatory test and the herd screening test along with

A



Fig. 2. Evaluation of the control measures. The numbers of detected and missed cattle with or without control measures were estimated by the scenario tree model. A: Efficacies of vaccination with single-, double-, or triple-administrations were compared. B: Efficacies of the common pasture test with coverages from 10 to 50%, and the bulk milk test were compared. For each box-whisker plot, the width of the box represents the interquartile range; the black line indicates the mean; the ends of the whiskers represent the 5th and 95th percentile, respectively; the dots represent the outliers.

implementation of mass vaccination programs, the scenario models with adoption of mass vaccination program, use of communal pasture with a rate between 10 to 50%, and/or implementation of a bulk milk test were run.

Phase two scenario

With an assumption that risk of BVD in cattle is reduced by the implementation of effective control measures, the P_p was changed from 0.4% as a default setting to either 0.13, or 0.04%. Following the change of P_p , P_t was changed from 0.1% to either 0.03, or 0.01%. Under the low BVDV prevalence, the scenario models with adoption of a triple-administrations of vaccination program, use of communal pasture by 30% of cattle, and/or implementation of bulk milk tests were run.

RESULTS

Influence of each parameter on the model output

The numbers of detected or missed BVD cattle in the output of the simulation with each vaccination program were compared. Interestingly, the numbers of detected cattle were not apparently different among single-, double-, and triple-administrations of vaccination programs. On the other hand, the numbers of missed BVD cattle increased with decreases in the number of vaccine administrations (Fig. 2A).

The numbers of detected and missed BVD cattle in the outputs of the simulations with each rate of communal pasture use were compared. The rates of cattle kept at communal pasture in the total population ranged from 10 to 50%. The numbers of detected BVD cattle increased and the numbers of missed BVD cattle decreased with increases in rates of use of communal pasture (Fig. 2B). The numbers of BVD cattle detected and missed in the output of the bulk tank milk test were approximately equal to those with 40 and 20% of communal pasture use, respectively (Fig. 2B).

Given the comparison among vaccination program and individual test in measures of communal pasture, as the numbers of missed BVD cattle apparently decreased, it should be desirable under no interventions for the mass vaccination of cattle to be adopted as this is a more effective control for BVD control to decrease the number of missed animals.



Fig. 3. Evaluation of the control measures under three vaccination programs. The numbers of detected and missed cattle with or without control measures under three vaccination programs were estimated by the scenario tree model. Efficacies of common pasture tests with coverages from 10 to 50%, and bulk milk tests under vaccination with a single administration (A), double administration (B), and triple administrations (C) were compared. For each box-whisker plot, the width of the box represents the interquartile range; the black line indicates the mean; the ends of the whiskers represent the 5th and 95th percentile, respectively; the dots represent the outliers.

Phase one scenario simulation

The efficacies of the disease detection program under the implementation of vaccination were compared (Fig. 3). When compared among the same monitoring test for detection, the numbers of detected and missed BVD cattle decreased with increases in the number of vaccine administrations, although the degree of decrease among vaccination programs found was small. When compared in the same vaccination program, the differences in the number of detected and missed BVD cattle between 10 and 50% were smaller under the vaccine programs incorporating a greater number of vaccine administrations. We compared the efficacies of the individual test prior to communal pasture grazing and bulk milk tests under the vaccination strategy so as to estimate which rates of use in communal pasture tests would be equally effective to the screening test using bulk tank milk. The numbers of detected cattle by the bulk tank milk tests under single-, double- and triple-administrations were almost equal to those of BVD cattle in 20, 30 and 30% use rates of communal pasture tests, respectively. In parallel, the numbers of missed cattle by the bulk tank milk tests respectively. Considering these results together, during the early phase of the vaccination program, it was indicated that the efficacy of the communal pasture test with low use rates would be equal to that of the bulk milk tests, whereas at the later phase of the vaccination program, the efficacy of the bulk milk test would not be inferior to that of the



Fig. 4. Evaluation of the control measures in low bovine viral diarrhea virus (BVDV) prevalence. The numbers of detected and missed cattle with control measures in low BVDV prevalence were estimated by the scenario tree model. Efficacies of bulk milk tests, common pasture test with coverage of 30%, and a mix of bulk milk test and common pasture test with 30% coverage in BVDV prevalence of 0.4% (A), 0.13% (B) and 0.04% (C) were compared. For each box-whisker plot, the width of the box represents the interquartile range; the black line indicates the mean; the ends of the whiskers represent the 5th and 95th percentile, respectively; the dots represent the outliers.

communal pasture test with low coverage rates. We should conclude that the communal pasture test is more effective to control BVD during the early phase of a vaccination program than a screening test using bulk tank milk.

Phase two scenario simulation

To understand the efficacy of the interventions for BVD control in lower BVDV prevalence, the model run was conducted under several scenarios of BVDV prevalence. The numbers of detected and missed BVD cattle by the individual test prior to 30% communal pasture use and/or the bulk tank milk test were calculated in each of the scenarios with 0.4% of BVDV prevalence (Fig. 4A), 0.13% of BVDV prevalence (Fig. 4B), and 0.04% of BVDV prevalence (Fig. 4C).

Generally, the numbers of detected BVD cattle in any detection tests decreased with a decrease of BVDV prevalence; however, in each of the three scenarios, the numbers of detected BVD cattle were not drastically different among the three detection tests. On the other hand, the numbers of missed BVD cattle in any detection tests were not drastically different among the three scenarios. However, in each of the three detection tests, the numbers of missed BVD cattle were not drastically different among the three scenarios of BVDV prevalence. There were differences in the numbers of missed BVD cattle under higher BVDV prevalence between the screening and individual tests; however, these differences became smaller under lower BVDV prevalence. Notably, under the high or middle level BVDV prevalence, not only a smaller number of missed BVD cattle, but also a smaller interval range of missed BVD cattle were found in the results of the screening test compared with the results of the individual test. Considered together, it was indicated that under lower BVDV prevalence resulting from the implementation of appropriate measures mentioned above, the efficacy of the bulk tank milk test would be more effective for control of BVD than the communal pasture test to steadily decrease the number of missed BVD cattle.

DISCUSSION

We developed a stochastic scenario tree model to evaluate the control measures of BVD using the field data in Nemuro, Hokkaido. Although most of the data were referred from the scientific literature or used the field data in Nemuro, some of the parameters in the present model were set according to the opinions of veterinarians in there. In order to confirm the confidence of the developed scenario tree model, the outputs of the model using the actual numbers of cattle, virus prevalence estimated in Saino *et al.*, and probability of using communal summer pastures, were compared with the numbers of reported cases of BVD in the last 5 years [29]. The results of model validation could support that the developed model would be reflecting the real situation with high confidence (Data not shown).

In Scandinavian countries including Denmark, Sweden, Norway, and Finland, BVD had been eradicated due to implementation of an eradication program without vaccination [24, 30]. Moreover, the Norwegian government banned the use of the vaccination for pestiviruses as the vaccination could prevent the manifestation of severe clinical manifestations, but did not prevent the spread of the virus [24]. In these countries, cattle with antibodies against BVDV as potential risk of BVD spread is easily monitored but it can also cull no-virus-shedding cattle which was transiently infected in past. In Japan, on the other hand, we are monitoring virusshedding cattle and adopt the vaccination so as to prevent the further virus spread. According to the results of the present study, it was revealed that the adoption of mass vaccination decreased the number of missed BVD cattle, although it did not increase the number of detected BVD cattle. This may indicate that, in an area where no interventions for BVD control have been implemented, the adoption of the mass vaccination campaign would be the foremost intervention for BVD control to reduce the number of cattle infected with BVDV. In the present model, herd immunity conferred by mass vaccination did not affect the number of PI cattle, but did affect the number of TI cattle due to a decrease of the sensitive population. Under system of no BVD monitoring, PI cattle would not be detected without showing obvious clinical manifestations, such as reproductive disorder. Given these considerations, the missed BVD cattle under the mass vaccination program would be the TI cattle of any age or PI heifers. As these diseased cattle usually do not show obvious clinical manifestations, it is likely that undetected cattle would be left for a period, thereby allowing the shedding of infectious viruses. Even TI cattle would be a source of BVDV spread, resulting in horizontal and vertical infection over a limited duration.

It is estimated that immediately after initiating the mass vaccination program, the prevalence of BVDV is not low. Therefore, monitoring of infected cattle in parallel with the implementation of mass vaccination should be an effective intervention to reduce the risk of BVD in the population. In Nemuro, an individual test using a serum sample is mandatory applied to all cattle that use communal pastures [29]. In Japan, the use of communal pasture for summer grazing is common in any age of dairy cattle, including pregnant cows; therefore, the spread of BVDV infection horizontally or vertically in communal pastures has been suspected. An individual test prior to communal pasture grazing can detect young PI heifers that normally rarely show clinical manifestations; however, there are practical difficulties associated with adopting this test for a larger number of targeted cattle. On the other hand, the bulk tank milk test can be applied for a large number of animals, and some of the reports have already evaluated the efficacy of the bulk tank milk test [4, 18, 21]. As the target of the bulk tank milk test is lactating cows, this test is unlikely to detect the young PI heifers. Given that adoption of mass vaccination cannot contribute to a decrease of PI animals in the population, the individual test rather than the bulk tank milk test is more effective to reduce PI cattle immediately after conducting the mass vaccination. This consideration may support the results showing that the efficacy of the individual test with a lower coverage was equal to that of the bulk tank milk test in the present study. Comparison of efficacies between the individual test and screening test appears to be affected by some factors, including test sensitivity, virus prevalence, and distribution of targeted animals. Input of suitable data to the disease model as well as collection of accurate data in the field is necessary to reach a confidential output so as to contribute to the selection of a more effective intervention.

Implementation of appropriate interventions such as a mixture of mass vaccination and the individual test resulted in the decrease of numbers of PI animals in all the age groups. As the remaining PI cattle in the herd are known to be a source of new infection of BVDV in the herd, and may produce PI calves, we should aim to detect as many PI and TI cattle of any age as possible for the BVD eradication in the area by performing screening rather than individual tests. However, if animal-level prevalence of BVD becomes lower, it would be likely that the cost-effectiveness of the screening test using bulk tank milk would be higher due to lower between-herd prevalence. This would be because implementation of outbreak investigation could eradicate the BVD cattle in premises, thereby decrease of BVD cattle would affect to between-herd prevalence but not to within-herd prevalence in infected premises. In the situation of a higher number of herds being free from BVD under a very low prevalence of BVD, we could save resources by investigation once the infected herd is detected by the screening test. The decision to switch between implementation of the individual and screening test is critical so as to achieve risk management with high cost-effectiveness. In further analysis, the scenario model with more detailed conditions should be run to compare the outputs both at hygiene level and cost-effectiveness.

We developed the scenario tree model to evaluate the efficacies of interventions for BVD control using the archived data of Nemuro, Hokkaido where the cattle infected with BVDV have been reduced by the independent eradication program. The model output indicated that the adoption of mass vaccination apparently decreased the number of missed BVD cattle, although it did not increase the number of detected BVD cattle. Under implementation of mass vaccination, the efficacy of an individual test with lower coverage was equal to that of the herd screening test using bulk tank milk. Moreover, the switch from the individual to herd screening test under a situation of very low BVDV prevalence should be critical to achieve BVD eradication more efficiently. The scenario tree model developed was useful to simulate the BVD eradication program in the endemic area.

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