



Title	Evaluation of bovine viral diarrhoea virus control strategies in dairy herds in Hokkaido, Japan, using stochastic modelling
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3

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26

27 **Summary**

28 Bovine viral diarrhoea virus (BVDV) infection in cattle can result in growth retardation,
29 reduced milk production, reproductive disorders, and death. Persistently infected animals are
30 the primary source of infection. In Hokkaido, Japan all cattle entering shared pastures in
31 summer are vaccinated before movement for disease control. Additionally, these cattle may be
32 tested for BVDV and culled if positive. However, the effectiveness of this control strategy
33 aiming to reduce the number of BVDV infected animals has not been assessed. The aim of this
34 study was to evaluate the effectiveness of various test and cull and/or vaccination strategies on
35 BVDV control in dairy farms in two districts of Hokkaido, Nemuro and Hiyama. A stochastic
36 model was developed to compare the different control strategies over a 10-year period. The
37 model was individual-based and simulated disease dynamics both within and between herds.
38 Parameters included in the model were obtained from the literature, the Hokkaido government
39 and the Japanese Ministry of Agriculture, Forestry and Fisheries. Nine different scenarios were
40 compared: no control, test and cull strategies based on antigen testing of either calves or only
41 cattle entering common pastures, vaccination of all adult cattle or only cattle entering shared
42 pastures, and combinations thereof. The results indicate that current strategies for BVDV
43 control in Hokkaido slightly reduced the number of BVDV infected animals; however,
44 alternative strategies such as testing all calves and culling any positives or vaccinating all
45 susceptible adult animals dramatically reduced those. To our knowledge, this is the first report
46 regarding the comparison of the effectiveness between the current strategies in Hokkaido and
47 the alternative strategies for BVDV control measures.

48

49 **Keywords:** Bovine viral diarrhoea virus, Common pasture, Control strategies, Culling,
50 Modelling, Vaccination

51

52

53 **Introduction**

54 Acute infection with Bovine viral diarrhea virus (BVDV) in cattle results in temporary
55 fever, respiratory symptoms and diarrhea (Bachofen et al., 2010). Rarely, acutely infected
56 animals may suffer from high fever and internal bleeding due to thrombocytopenia. More
57 importantly, infection during specific stages of pregnancy can result in the birth of
58 immunotolerant calves that are persistently infected (PI) with BVDV (McClurkin et al., 1984).
59 These PI animals, which continuously excrete virus into the environment, are accepted as the
60 main source of transmission regarding the short time span of low-intensity viral excretion by
61 transiently infected animals within a herd. In some cases, clinical signs in PI animals can be
62 differentiated pathogenetically into mucosal disease (Bachofen et al., 2010). Economic losses
63 in dairy and beef industries due to BVDV infection are substantial (Houe, 2003; Weldegebriel
64 et al., 2009). Thus, many countries have implemented BVDV control programs (Greiser-Wilke
65 et al., 2003).

66 Identification and removal of PI animals is an effective strategy for clearing infected
67 herds from BVDV infection, and a common approach to prevent new herds from becoming
68 infected. BVDV control strategies differ between countries (Lindberg and Houe, 2005; Campen,
69 2010). In most European countries, BVDV control is aimed at systematic eradication without
70 vaccination. Systematic programs have been implemented in Denmark, Finland, Norway,
71 Sweden, Ireland, Belgium, and Scotland (Bitsch et al., 2000; Lindberg et al., 2006; Graham et
72 al., 2014; Hanon et al., 2014). Based on Scandinavian strategies, which is test and culling only
73 PI cattle but no vaccination, a regional BVDV control program was launched in lower Austria
74 in 1996 and later extended to the entire country (Rossmanith et al., 2005). A national
75 compulsory eradication program has been in place in Switzerland since 2008 (Presi and Heim
76 et al., 2010; Presi et al., 2011). This program is based on identification and culling of PI animals
77 through antigen testing of newborn calves using ear notch samples, a ban on vaccination, and
78 movement restrictions. In contrast, in the United States (US) several voluntary BVDV control

79 programs that include vaccination have been implemented. It has been estimated that
80 approximately 80% of US cattle are vaccinated with either inactivated or modified live virus
81 vaccines to prevent both fetal infections and acute disease (Campen, 2010).

82 In Japan, BVDV was first reported in 1967 and is currently endemic (Nagai et al., 2008;
83 Kadohira and Tajima, 2010; Abe et al., 2015). According to a pilot survey on PI animals in dairy
84 farms in Japan, the prevalence of PI animals at the herd level and animal level were 7.59% and
85 0.12%, respectively, in some regions in 2014 (Kameyama et al., 2016). In Hokkaido, which is
86 located in northern Japan between 41°31' and 45°30' N latitude and between 139°20' and
87 148°53' E longitude, BVDV has already spread and become an endemic disease in dairy herds;
88 thus, some regional voluntary BVDV control programs have already been implemented
89 (Yasutomi et al., 2004; Kadohira et al., 2007). The current control activities targets pastured
90 animals (Saino et al., 2013). In Japan, a part of cattle are released in a common pasture shared
91 by the multiple herds. All cattle that graze on common pastures during the summer must be
92 vaccinated against BVDV before their movement to common pasture. Both modified live
93 vaccines and killed vaccines are used with the aim to prevent fetal infections and acute disease
94 in most prefectures in Japan including Hokkaido. These cattle are also tested by RT-PCR or
95 virus isolation before entering common pastures, and any positive PI animals are culled. These
96 tests are conducted at Livestock Hygiene Service Center in Hokkaido. Grazing on shared
97 pastures is a major risk factor for BVDV spread between herds (Bitsch et al., 2000; Rossmannith
98 et al., 2005; Valle et al., 1999). However, it is not known whether the current BVDV control
99 strategies are effective for reducing the prevalence of PI animals at the regional level. Therefore,
100 the aim of this study was to evaluate the BVDV control programs in Hokkaido using stochastic
101 modelling. Epidemiological models of BVDV infection are well established (Courcoul and
102 Ezanno, 2010; Gates et al., 2013; Gates et al., 2014; Tinsley et al., 2012). However, common
103 pastures are not included and control strategies concerning common pastures were not tested in
104 previous studies. Here, we developed an individual-based model to describe the dynamics of

105 BVDV transmission within and between herds and common pastures. Nine different scenarios
106 were assessed with this model: no control, a test-and-cull strategy based on antigen testing of
107 calves or only cattle entering common pastures, vaccination of either adult cattle or only cattle
108 entering common pastures, and combinations thereof. Additionally, we compared the
109 effectiveness of voluntary and compulsory programs.

110

111 **Materials and methods**

112 **Model**

113 To assess the effectiveness of interventions, we constructed a mathematical model describing
114 the transmission process of BVDV at two levels simultaneously, within a dairy herd and
115 between dairy herds. To capture the stochasticity of transmission events, we employed an
116 individual-based stochastic model. Our model is a compartmental SIR-like model; the transition
117 of infection status was previously described (Ezanno et al., 2007). The hosts were classified by
118 the following infection states: *M*: maternal antibodies, *S*: susceptible, *TI*: transiently infected,
119 *PI*: persistently infected, *CP*: recovered cows still carrying *PI* animals, *R*: other recovered
120 animals, or *V*: vaccinated animals (Fig. 1).

121 The transmission of BVDV occurs by horizontal transmission (from *PI* and *TI* animals
122 to *S* animals) and vertical transmission (*TI* or *PI* animals carry *PI* offspring). The transmission
123 probability of horizontal transmission differs according to management groups. The animals
124 were stratified by five groups based on the breeding place in a herd: calves, young heifers,
125 pregnant heifers, dry cows, and lactating cows. Age structure of cows is parameterized based
126 on the data of the duration of age group provided by the Hokkaido government. We modeled
127 the heterogeneity of horizontal transmission probability due to the breeding place in herds
128 stratified by age group of cows. The probability of horizontal transmission per a susceptible

129 host, so-called the force of infection at specific time point t , $\lambda(t)$, is given by the following
 130 formulae:

131

$$\begin{aligned}
 \lambda_{calves,x}(t) &= \beta_{PI} \frac{PI_{calves,x}(t)}{N_{calves,x}(t)} + \beta_{TI} \frac{TI_{calves,x}(t)}{N_{calves,x}(t)} + \beta_{PI,diff} \left(\frac{PI_{heifers,x}(t)}{N_{calves,x}(t)N_{heifers,x}(t)} + \frac{PI_{adults,x}(t)}{N_{calves,x}(t)N_{adults,x}(t)} \right), \\
 \lambda_{heifers,x}(t) &= \beta_{PI} \frac{PI_{heifers,x}(t)}{N_{heifers,x}(t)} + \beta_{TI} \frac{TI_{heifers,x}(t)}{N_{heifers,x}(t)} + \beta_{PI,diff} \left(\frac{PI_{calves,x}(t)}{N_{heifers,x}(t)N_{calves,x}(t)} + \frac{PI_{adults,x}(t)}{N_{heifers,x}(t)N_{adults,x}(t)} \right), \\
 \lambda_{adults,x}(t) &= \beta_{PI} \frac{PI_{adults,x}(t)}{N_{adults,x}(t)} + \beta_{TI} \frac{TI_{adults,x}(t)}{N_{adults,x}(t)} + \beta_{PI,diff} \left(\frac{PI_{calves,x}(t)}{N_{adults,x}(t)N_{calves,x}(t)} + \frac{PI_{heifers,x}(t)}{N_{adults,x}(t)N_{heifers,x}(t)} \right),
 \end{aligned} \tag{1}$$

132

133 where $\lambda_{calves,x}$, $\lambda_{heifers,x}$, and $\lambda_{adults,x}$ denote transmission probability for calves (calves and young
 134 heifers), heifers (pregnant heifers), and adults (dry cows and lactating cows) in the x -th herd,
 135 respectively. The stratification of transmission probability, calves, young heifers, dry cows and
 136 lactating cows, is based on the place of management in herds. β_{PI} and β_{TI} denote the
 137 transmission coefficient of transmission from PI and TI animals within the same age group and
 138 same herd, respectively. $\beta_{PI,diff}$ denotes the transmission coefficient of transmission from PI
 139 animals between different age groups in the same herd. We set $\beta_{PI} = 0.5$, $\beta_{TI} = 0.03$, and $\beta_{PI,diff}$
 140 $= 0.1$ per 2 weeks in our simulation runs (Ezanno et al., 2007). PI , TI , and N denote the number
 141 of PI , TI , and total animals, respectively; e.g., $PI_{heifers,x}$ indicates the number of heifers whose
 142 infection status is PI in the x -th herd. For vertical transmission, only TI or PI pregnant heifers
 143 and lactating cows can carry PI offspring in our model. The offspring from PI animals are
 144 always PI . TI cows carry PI offspring (TI becomes CP) when the infection timing is early to
 145 mid-pregnancy (weeks 7–22) (Brownlie et al., 1987), otherwise TI carries M . TI cows obtain
 146 life-long immunity against BVDV (Brownlie et al., 1987; Young et al., 2006; Liebler-Tenorio
 147 et al., 2004), we assumed no re-infection for TI animals in our model. The infectious period is
 148 assumed to be 2 weeks for TI and life-long for PI animals, similar to the setting in the previous
 149 study (Ezanno et al., 2007). We assumed that there was an endemic equilibrium and the BVDV
 150 transmission dynamics have reached an endemic equilibrium at the initial time point of the
 151 simulations. To this end, we started the simulation runs with the frequency of transiently

152 infected cows = 0.001, the frequency of *PI* cows = 0.02 and other cows denoted as *R* for all
153 herds, and discarded the first four years to obtain the initial condition. We set 2 weeks as the
154 unit time. The development of the model and all subsequent analyses were performed using the
155 statistical software R 3.1.3 (R Development Core Team, 2015).

156 **Movement of animals**

157 Parameter values describing the movement patterns of cows were determined based on the
158 current situation in Hiyama and Nemuro, sub-regions of Hokkaido (Table 1 and 2), using data
159 from each sub-region provided by the Hokkaido government. We modeled two movement
160 patterns: i) movement of animals between herds (the common pasture is not included), and ii)
161 movement of animals between the herd and the common pasture. We parameterized the
162 movement of animals between herds based on field data as shown in Table 2. Animals that
163 moved between herds and destination herds were randomly selected. For movement between
164 herds and the common pasture, animals move to the common pasture at the beginning of May
165 and return to their herd at the end of October. The herds using common pastures are fixed over
166 time; constant herds used common pastures. The proportions of herds using common pastures
167 were 0.31 for Nemuro and 0.21 for Hiyama. Based on the situation in Japan, the animals moved
168 to common pastures in the model includes only young heifers. Young heifers moving to the
169 common pasture are determined randomly with probabilities of 0.19 for Nemuro and 0.32 for
170 Hiyama. Both movement patterns were independent of the infection status of animals.

171 **Herd demographics**

172 In addition to the classification according to infection status and herds, cattle were classified
173 into five groups according to age and reproductive status to describe the herd demographics.
174 The duration of each class and the mortality rate in each class are summarized in Table 3. We
175 modeled the birth of calves deterministically; cows over 15 months old (pregnant heifers or dry
176 cows) become pregnant once per year. At birth, newborn calves are *PI* or *M*. *M* describes
177 animals protected against BVDV infection by maternal antibodies. The length of protection by

178 maternal antibodies is parameterized as 84 days by the field data (Palfi et al., 1993). After this
179 period, M animals become S animals and can be infected with BVDV. At the same time,
180 newborn calves become heifers, and the duration before breeding is 280 days. After
181 insemination, all young heifers moved to the pregnant first-calf heifer group for 280 days.
182 Thereafter, a calf was born and the dam moved to the lactating cow group for 304 days and then
183 to the dry cow group for 60 days. At the end of the dry period, a new calf was born and the dam
184 moved to the lactating cow group again. The maximum cow lifespan in this model was assumed
185 to be 7 years because dairy cattle are usually slaughtered at approximately 7 years of age in
186 Japan. We assumed that breeding occurred constantly throughout the year based on the situation
187 in Japan. The mortality rate of PI and non-PI animals were based on the field data of mean
188 mortality rate of the entire cattle population in Japan during five years, which was obtained
189 from the Japanese Ministry of Agriculture, Forestry and Fisheries (Table 3). We included only
190 female cows in the model; males (50% of newborn animals) were assumed to be sold or culled
191 within 14 days after birth.

192 **Intervention**

193 In this study, we assessed the effectiveness of two types of interventions, vaccination and testing
194 and culling. We assumed that all cows in the herd were vaccinated if the herd used the
195 vaccination strategy, and revaccination was repeatedly conducted to maintain protection by the
196 vaccine. Vaccination scenarios assumed that all herds used vaccination. The vaccine efficacy
197 was assumed to be 80% (Newcomer et al., 2015), and the infection probability λ among
198 vaccinated animals was consequently reduced by 80%. For culling, we assumed that animals
199 were tested by RT-PCR for adults or by antigen Enzyme-Linked Immunosorbent Assay
200 (ELISA) testing using ear notch samples for calves, and that positive PI animals were culled.
201 Both tests are set to have the same performance. The sensitivity and specificity of RT-PCR and
202 ELISA were considered to be 99% and 100% for detecting PI animals, respectively (Presi et al.,
203 2010). The timing of culling was dependent on the strategy, as described below.

204 **BVDV control strategies**

205 In Hiyama and Nemuro, all animals that graze on common pastures must be vaccinated.
206 Furthermore, all animals are tested for BVDV before entering common pastures in those regions.
207 RT-PCR and/or virus isolation of BVDV from the blood are considered suitable diagnostic
208 techniques for BVDV infection in Japan, and positive animals cannot enter common pastures.
209 To assess the strategies including the current strategy used in Hiyama and Nemuro, we
210 constructed nine scenarios (Table 4) for simulation using our model. The baseline scenario
211 consisted of no control program (S1). In S2, all animals that moved to common pastures were
212 tested by RT-PCR using blood samples for adult animals before moving to common pastures,
213 and positive animals were culled. In S3, all animals were vaccinated before entering common
214 pastures. In S4, all animals that moved to common pastures were tested before moving to
215 common pastures and positive animals were culled, and all animals were vaccinated before
216 entering common pastures. In S5, all calves were tested by ELISA at the time of birth and
217 positive animals were culled. In S6, all animals over 6 months of age (young heifers, pregnant
218 heifers, lactating cows, and dry cows) were vaccinated with killed vaccine once per year. S7 is
219 the combination of ELISA testing of all calves and vaccinating all animals over 6 months of
220 age. S8 is the combination of antigen ELISA testing of all calves and vaccinating all animals
221 moving to common pastures. In S9, animals moving to common pastures were tested by RT-
222 PCR and animals over 6 months of age were vaccinated.

223 The effectiveness of interventions was measured by the prevalence of PI animals at the
224 end of the 10-year simulation period. To compare scenarios, 1,000 iterations in each scenario
225 were compared by pairwise Wilcoxon rank sum tests. P values <0.05 were considered
226 statistically significant.

227

228 **Results**

229 The results of the different BVDV control scenarios were similar in both Hiyama and Nemuro

230 (Fig. 2). The prevalence of *PI* animals with no control intervention (S1) fluctuated seasonally
231 based on summer pasture use for 10 years in this model (Fig. 2 and 3). In simulation modelling
232 of the current strategies in Hokkaido, the prevalence of *PI* animals gradually decreased by
233 testing and culling (S2), vaccination (S3), or the combination of testing and culling all animals
234 moving to common pastures and vaccinating all animals moving to common pastures (S4),
235 compared to the no control scenario (S1) (Fig. 2A and 2B). However, testing and culling of
236 positive calves (S5) resulted in a dramatic decrease in the prevalence of *PI* animals to almost
237 eradicated levels (Fig. 2C and 2D). Vaccinating all susceptible adult animals (S6) also reduced
238 the number of *PI* animals (Figure 2C and 2D). The combination strategies, i.e., testing and
239 culling all positive calves and vaccinating all susceptible adult animals (S7) and testing and
240 culling all positive calves and vaccinating all animals moving to common pastures (S8), yielded
241 similar results to those obtained from the single strategy of testing and culling all positive calves
242 (Fig. 2E and 2F). The combination strategy of testing all animals moving to common pastures
243 and vaccinating all susceptible adult animals (S9) also showed similar results to the single
244 strategy of vaccinating all susceptible adult animals (Fig. 2E and 2F). At the 10-year time point
245 an important and significant decrease in the prevalence of *PI* animals in comparison with no
246 control was demonstrated for the following strategies: testing and culling all positive calves
247 (S5), vaccinating all susceptible adult animals (S6), and combinations thereof (S7–S9) (Fig. 4).
248 Although a statistically significant decrease was found also for S2 and S4 after 10 years, the
249 prevalence of *PI* animals was still at sufficiently high levels to discard these as relevant options
250 for control. We then assessed the sensitivity of the proportion of herds participating in the
251 interventions S5 and S6 in Nemuro. Changes in the proportions of herds participating in S5 and
252 S6 control programs influenced the prevalence of *TI* and *PI* animals at the herd level (Fig. 5).
253 In both scenarios, the prevalence of *TI* and *PI* animals decreased as the proportion of
254 participants increased. When 100% of herds participated, S5 and S6 strategies markedly
255 reduced *TI* and *PI* prevalence at the herd level (Fig. 5A and 5B). We also assessed the

256 effectiveness of the intervention "test and culling all calves" with the varied sensitivity of
257 ELISA (Fig. 6). If the sensitivity of ELISA is not high enough, the intervention cannot eradicate
258 the BVDV epidemic within ten years from the beginning of intervention.

259

260 **Discussion**

261 Epidemiological models allow investigation of projected dynamics of virus spread and
262 various control strategies (Ezanno et al., 2007; Ezanno et al., 2008; Innocent et al., 1997;
263 Sorensen et al., 1995; Viet et al., 2007; Gunn et al., 2004; Viet et al., 2004). In the current study,
264 we established a stochastic model of within- and between-dairy herd BVDV infection dynamics,
265 including the effects of common pasture use, based on general and epidemiological data from
266 Hokkaido. In general, movement of animals is of most importance for the spread of BVDV
267 infection between herds on a global scale. The use of common pastures is also one of the risk
268 factors for the spread in areas where common pastures are used frequently (Bitsch et al., 2000;
269 Rossmanith et al., 2005; Valle et al., 1999); thus, it is important to consider the use of common
270 pastures in modelling.

271 If animals are in early pregnancy during pasture and become infected, then their
272 offspring become *PI* calves. Newborn *PI* calves can then spread the virus within their own herd
273 at the end of the summer season. *PI* calves born in the grazing period in common pastures are
274 also important. Trojan *PI* animals are an issue regardless of common pasture use. Movement of
275 animals between herds is another risk factor for BVDV spread. Introduction of infected animals
276 into an uninfected herd results in the spread of BVDV to the previously uninfected herd. In
277 endemic areas for BVDV, the prevalence of *PI* animals in the standing cattle population has
278 been reported to remain approximately 0.5–2% (Houe, 1999). This is considered to be due to
279 the balance between new persistent infections and removal of *PI* animals. The prevalence of *PI*
280 animals in Japan was 0.12% in the pilot survey performed in other regions in 2014 (Kameyama
281 et al., 2016). The low prevalence of *PI* has been observed in previous studies, but that currently

282 there are no good explanations for this. Further studies might help to understand BVDV
283 situation in Japan.

284 We evaluated the BVDV control strategies in two sub-regions of Hokkaido, Hiyama and
285 Nemuro, using stochastic modelling. Dairy farming is the main industry in Nemuro, and there
286 are more herds in this region than in Hiyama. However, the results of our simulations were
287 similar between Hiyama and Nemuro. These results indicate that a common BVDV control
288 strategy could be used in both areas. The current strategy of targeting animals moving to pasture
289 with the combination of testing and culling and vaccination (S4) significantly reduced the
290 number of BVDV infected animals in our model, but not too a sufficiently low level to achieve
291 control or eradication. Likewise, the other current strategies of targeting animals moving to
292 pasture, i.e., testing and culling (S2) and vaccination (S3) also significantly, but not sufficiently,
293 reduced the number of BVDV infected animals compared with no control. This is likely because
294 the numbers of tested or vaccinated animals were insufficient to control BVDV transmission.
295 The proportions of animals moving to common pastures were very low in both areas and fewer
296 animals moving to common pastures resulted in fewer animals being tested or vaccinated.
297 Hence, the effect of control strategies targeting only pastured animals on the prevalence of PI
298 animals in Hokkaido is only limited. Certainly the use of common pastures is a risk factor for
299 spreading BVDV between herds; however, farm management situations differ between each
300 farm, area, or country. Therefore, it is important to control risk factors according to individual
301 herd situations. Our results suggest that testing all calves and culling the positive ones (S5) is
302 an effective control method. Elimination of *PI* animals is a rapidly effective means of
303 controlling BVDV within a herd. In Switzerland, a national BVDV eradication program
304 launched in 2008 (Presi and Heim, 2010). All calves must be ear notched and the ear tissue
305 tested using the antigen ELISA test for BVDV within 5 days of birth, and all positive calves are
306 culled. This compulsory eradication program has resulted in a reduction in the prevalence of *PI*
307 animals in Switzerland (Presi et al., 2011). However, combining S5 with the strategies in other

308 scenarios is unlikely to produce a synergistic or additive effect because the test sensitivity is
309 very high. The effectiveness of S5 depends primarily on the participation rate of the herds in
310 this BVDV control program. The Swedish program was voluntary for almost 10 years before it
311 became compulsory (Hult and Lindberg, 2005; Lindberg et al., 2006; Lindberg and Alenius
312 1999). Compulsory control approaches are biosecurity based and aim to prevent introduction
313 of the infection into uninfected herds, eliminate *PI* animals from infected herds, and rapidly
314 detect new infections (Lindberg et al., 2006). These approaches have been successful in BVDV
315 control (Hult and Lindberg, 2005); however, compulsory testing and culling of all positive
316 calves is costly and requires significant manpower. Therefore, bulk milk testing should be used
317 to establish herd status before individual testing. Then, in infected herds, all animals should be
318 tested using ear notch or blood samples. The results of our model showed that vaccination of
319 all adult animals (S6) could also effectively lead to BVDV control even without culling *PI*
320 animals. Though *PI* animals are not actively removed, they leave the herd by natural means.
321 This suggests that the role of vaccination is to prevent new infection, while testing and culling
322 calves directly decreases the number of *PI* animals. Although the prevalence of *PI* animals
323 decreased slowly in S6 compared with S5, it was sufficient for BVDV control through
324 vaccination.

325 The optimal vaccine strategy has been greatly debated. A meta-analysis on the efficacy
326 of BVDV vaccination to prevent reproductive disease was conducted, which showed that fetal
327 infection was decreased by over 80% with killed vaccine (Newcomer et al., 2015). Live vaccine
328 was more efficient in fetal infection than killed vaccine. We similarly would not expect a
329 synergistic or additive effect if the control strategy in S6 were combined with other strategies
330 targeting pastured animals (as in S9). Though *PI* animals are not actively removed in S6, 100%
331 of herds participated in the vaccination program, drastically reducing *TI* and *PI* prevalences.
332 This would be effective to control BVDV, which suggests that herd immunity contributes to the
333 effectiveness of this strategy as a BVDV control measure. If a sufficient proportion of the

334 population is vaccinated and immunized, then the potential for contact between infected and
335 susceptible animals decreases and the epidemic fails to spread (Garnett, 2005). However,
336 eradication of BVDV has never been achieved in practice in spite of many decades of
337 vaccinations. The vaccination effectiveness could be reduced by low vaccine coverage, low
338 quality of vaccine and improper use of vaccine. Taking into account these issues is required to
339 evaluate the vaccination effectiveness in practical settings. Because each control strategy has
340 advantages and disadvantages, it is important to find the right balance between testing and
341 culling and vaccination to control or eradicate BVDV. Rapid reduction of *PI* animals may not
342 necessarily be the correct strategy when considering the cost of *PI* animal testing, *PI* animal
343 removal, vaccination, and manpower needed for these strategies. Measures should be selected
344 according to the situation. Furthermore, a compulsory program is necessary for BVDV
345 control/eradication programs to ensure their effectiveness.

346 Our study has several limitations. In this study, we focused the evaluation of control
347 program on the dairy farms with common pasture since those programs have been implemented
348 on the dairy farms. If non-dairy farms were involved in the programs, it would need to be
349 considered. In addition, other “hot spots” for BVDV transmission, e.g., livestock markets and
350 exhibitions, are also important for the BVDV transmission modelling. The time trend of
351 demographic changes, e.g., the number of dairy herds decrease but the average herd size
352 increases, is also important for the quantitative assessment of BVDV interventions. Our model
353 did not take into account the heterogeneity of the daily herd network with respect to
354 geographical distance, social preferences between herds, and seasonality of animal movement
355 due to the difficulty in obtaining the data. These heterogeneities may affect to our results.

356 In conclusion, stochastic modelling allows us to predict the extent and duration of
357 infection and evaluate the efficacy of control strategies. The results indicate that the current
358 strategies for BVDV control in Hokkaido slightly reduced the number of BVDV infected
359 animals; however, alternative strategies such as testing all calves and culling any positives or

360 vaccinating all susceptible adult animals drastically reduced those in this region. The proportion
361 of herds participating in the intervention is also a major driver of success for BVDV control.
362 These findings give us the opportunity to reconsider BVDV control strategies and highlight the
363 importance of using control measures that have been proven effective.

364

365 **Abbreviations**

366 BVDV: bovine viral diarrhea virus

367 PI: persistently infected

368 TI: transiently infected

369 RT-PCR: reverse transcription polymerase chain reaction

370 ELISA: enzyme-linked immunosorbent assay

371

372 **Declarations**

373 Competing interests

374 The authors declare that they have no competing interests.

375

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379

380 Authors' contributions

381 SS, PP, NI, and YS designed the simulation study. PP and RO designed the mathematical model.

382 SS and RO performed the simulations and drafted the manuscript. KS, MS, YY, TU, HN, and

383 YF participated in epidemiological analyses and critically reviewed the paper. All authors read

384 and approved the final manuscript.

385

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500

501 **Figure legends**

502 **Figure 1 Schematic representation of BVDV transmission dynamics within a dairy herd.**

503 Cattle were divided into one of five age categories (calves, heifers, pregnant first-calf heifers,
504 lactating cows, and dry cows). Cattle were also classified into one of six BVDV-related health
505 status groups (*M*: maternal antibody protection; *S*: susceptible animals; *TI*: transiently infected
506 animals; *PI*: persistently infected animals; *CP*: recovered cows still carrying *PI* animals; *R*:
507 other recovered animals; and *V*: vaccinated animals). Dashed line indicates the birth of cows.

508

509 **Figure 2 Changes in the prevalence of PI animals over a 10-year period.** Nine different

510 BVDV control scenarios in Hiyama (left column) and Nemuro (right column) were evaluated.
511 S1: no control. S2: testing all animals moving to common pasture and culling all positives. S3:
512 vaccinating all animals moving to pasture. S4: a combination of testing all animals moving to
513 pasture and culling all positives and vaccinating all animals moving to pasture. S5: testing all
514 calves and culling all positives. S6: vaccinating all adult animals. S7: a combination of testing
515 all calves and culling all positives and vaccinating all adult animals. S8: a combination of testing
516 all calves and culling all positives and vaccinating all animals moving to pasture. S9: a
517 combination of testing all animals moving to pasture and culling all positives and vaccinating
518 all adult animals.

519

520 **Figure 3 Change in the average prevalence over 10 years of PI and TI animals in Nemuro.**

521 The common pasture enhances the prevalence of TI and PI animals during its season when no
522 intervention is conducted.

523

524 **Figure 4 Boxplot of the prevalence of PI animals at the 10-year time point.** Nine different

525 BVDV control scenarios in Hiyama (A) and Nemuro (B) were tested. S1: no control. S2: testing
526 all animals moving to common pasture and culling all positives. S3: vaccinating all animals

527 moving to pasture. S4: a combination of testing all animals moving to pasture and culling all
528 positives and vaccinating all animals moving to pasture. S5: testing all calves and culling all
529 positives. S6: vaccinating all adult animals. S7: a combination of testing all calves and culling
530 all positives and vaccinating all adult animals. S8: a combination of testing all calves and culling
531 all positives and vaccinating all animals moving to pasture. S9: a combination of testing all
532 animals moving to pasture and culling all positives and vaccinating all adult animals.

533

534 **Figure 5 Changes in the prevalence of PI animals at the herd level in Nemuro.** The
535 proportion of herds participating in S5 and S6 changed. A: Prevalence for 0% (black), 25%
536 (green), 50% (brown), 75% (blue), and 100% (gray) of herds participating in a program that
537 tests and culls all calves. B: Prevalence for 0% (black), 25% (green), 50% (brown), 75% (blue),
538 and 100% (gray) of herds participating in a program that vaccinates all adult animals.

539

540 **Figure 6: The effectiveness of "test and culling all calves" with varying the sensitivity of**
541 **ELISA.** The effectiveness was measured by the prevalence of TI and PI animals in the
542 Nemuro setting. Prevalence for 60% (black), 70% (green), 80% (brown), 90% (blue), and
543 99% (gray) of sensitivity in a program that tests and culls all calves.

544

545 **Table 1 Farm structure in Nemuro and Hiyama.** The herd size and the number of animals
 546 in each age group per herd indicates its average in Nemuro and Hiyama. All values are
 547 derived from statistics data of Hokkaido government.

	Nemuro	Hiyama	Reference
Total herds	1500	62	
Total animals (heads)	189000	3472	
Average herd size (animals/farm)	126	56	
Number of animals in each age group per herd (heads)	Calves	23	5 Hokkaido
	Young heifers	11	3 government
	Pregnant heifers	8	8
	Dry cows	28	8
	Lactating cows	56	32

548
 549

550 **Table 2 Parameters regarding the movement of cows in Nemuro and Hiyama.** The
 551 proportion of animals moving between herds and those in each category per year indicate
 552 average in Hokkaido. Only young heifers are sent to common pasture for 6 months in
 553 Japanese farming system. All values are derived from statistics data of Hokkaido government.

	Nemuro	Hiyama	Reference
The proportion of animals moving between herds per year	0.2	0.2	
The proportion of animals moving between herds in each age category per year			
Calves	0.19	0.19	
Young heifers	0.08	0.08	
Pregnant heifers	0.16	0.16	
Lactating and dry cows	0.57	0.57	
Period of time for the common pasture per year (months)	6	6	Hokkaido government
Number of common pasture	11	1	
Number of farms using one pasture (herds/pasture)	42	13	
The proportions of herds using common pastures	0.31	0.21	
Proportion of animals going to pasture per farm	0.19	0.32	
Age group of the animals in common pasture	Young heifers	Young heifers	

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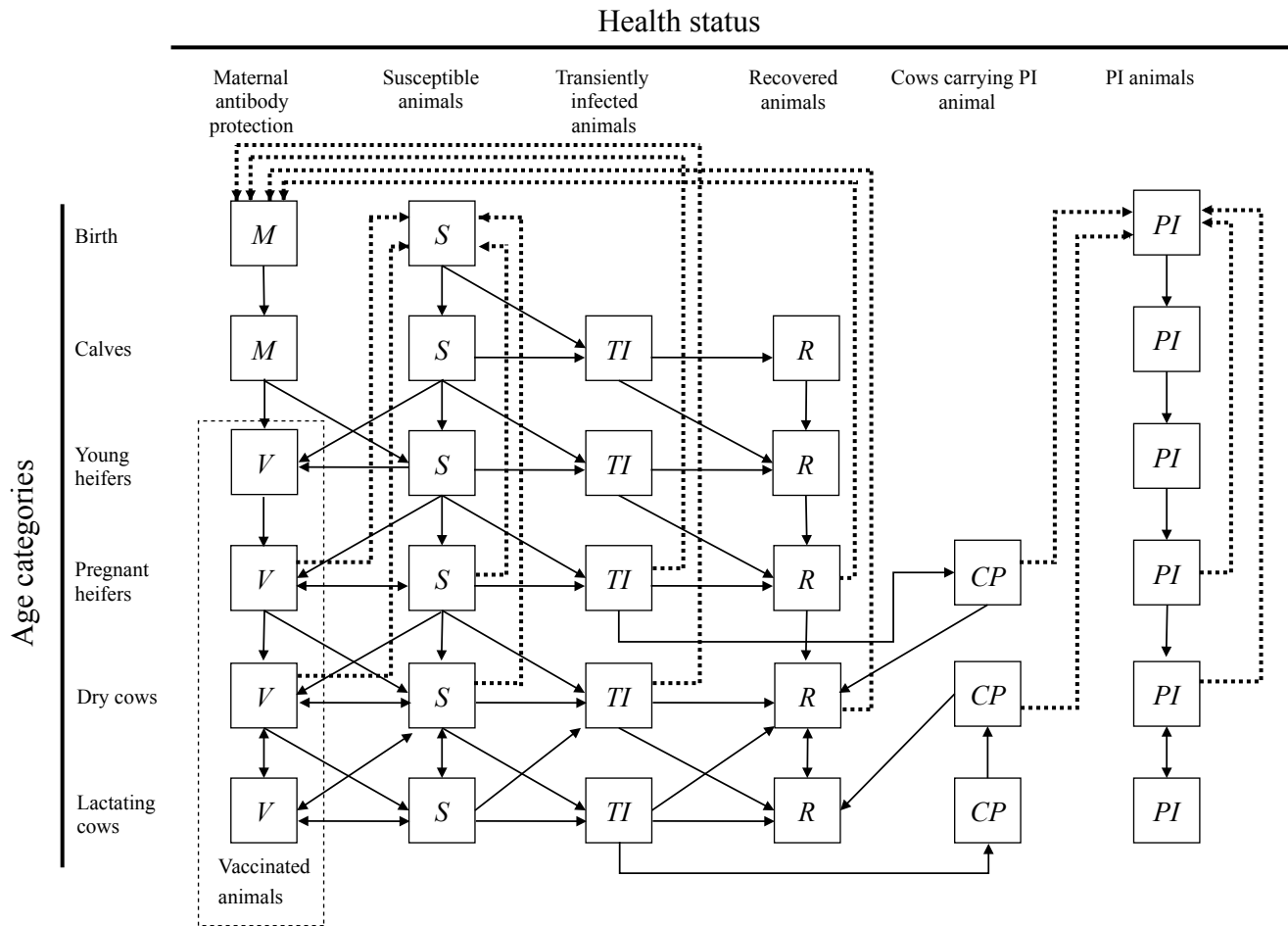
559 **Table 3 Mortality rate (including slaughter per 2 weeks) and duration in each class of**
 560 **cows.** The mortality rate of non-PI or PI per 2 weeks are derived from statistics data of
 561 Japanese Ministry of Agriculture, Forestry and Fisheries.
 562

Class	Duration	Mortality rate of non-PI per 2 weeks	Mortality rate of PI per 2 weeks	Reference
Birth	14 days	0.0012	0.0273	Japanese Ministry of Agriculture, Forestry and Fisheries
Calves	182 days	0.012	0.0273	
Young heifers	280 days	0.0057	0.0273	
Pregnant heifers	280 days	0.002	0.0273	
Dry cows	60 days	0.004	0.0273	
Lactating cows	304 days	0.004	0.0273	

563

564 **Table 4 BVDV control scenarios.** Nine different scenarios were evaluated. +: selected; -: not
565 selected. S1: no control; S2: all animals that moved to common pastures were tested by RT-
566 PCR before moving to common pastures, and positive animals were culled; S3: all animals
567 were vaccinated before entering common pastures; S4: all animals that moved to common
568 pastures were tested before moving to common pastures and positive animals were culled, and
569 all animals were vaccinated before entering common pastures; S5: all calves were tested by
570 ELISA at the time of birth and positive animals were culled; S6: all animals over 6 months of
571 age were vaccinated with killed vaccine once per year; S7: the combination of ELISA testing
572 of all calves and vaccinating all animals over 6 months of age; S8: the combination of antigen
573 ELISA testing of all calves and vaccinating all animals moving to common pastures; S9:
574 animals moving to common pastures were tested by RT-PCR and animals over 6 months of
575 age were vaccinated.
576
577

Scenario	Test & culling		Vaccination	
	All calves	All animals pasturing	All adults	All animals pasturing
S1	-	-	-	-
S2	-	+	-	-
S3	-	-	-	+
S4	-	+	-	+
S5	+	-	-	-
S6	-	-	+	-
S7	+	-	+	-
S8	+	-	-	+
S9	-	+	+	-

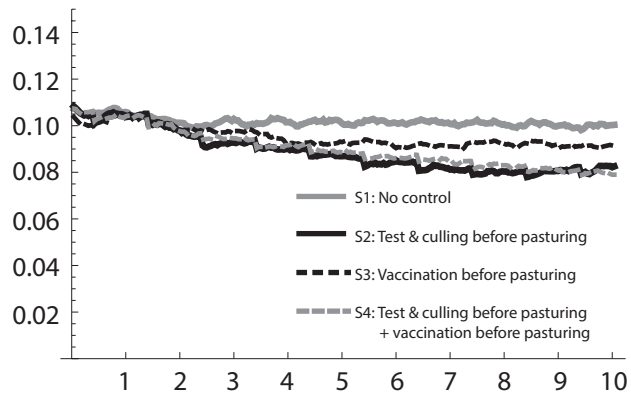


Sekiguchi et al. Figure 1

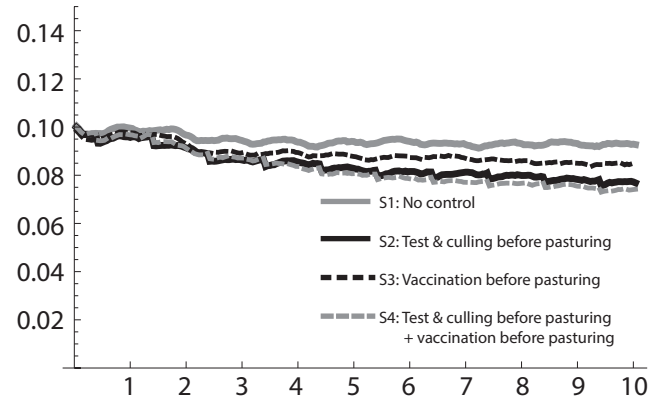
Hiyama

Nemuro

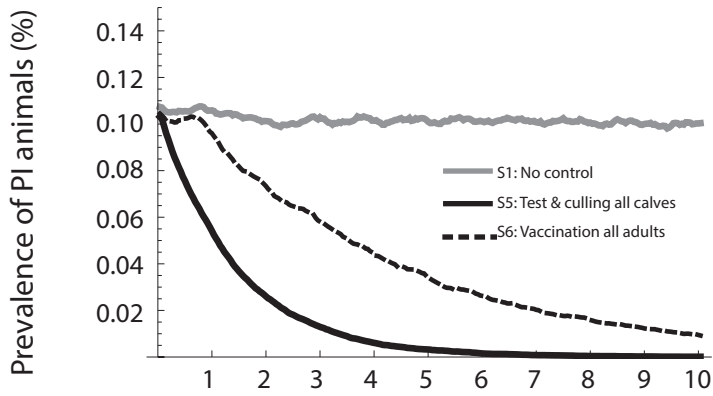
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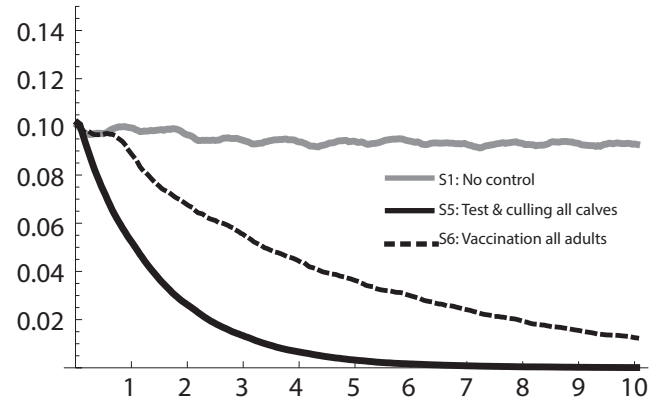
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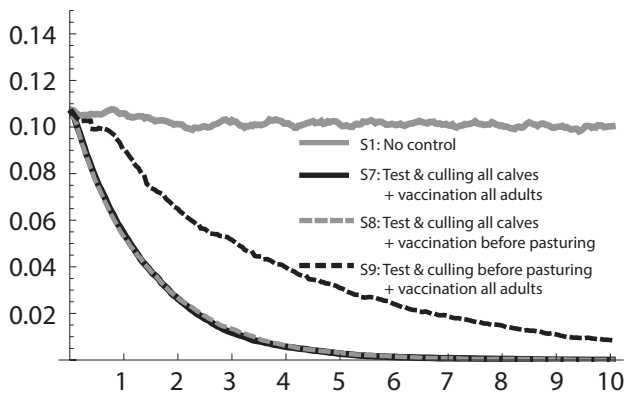
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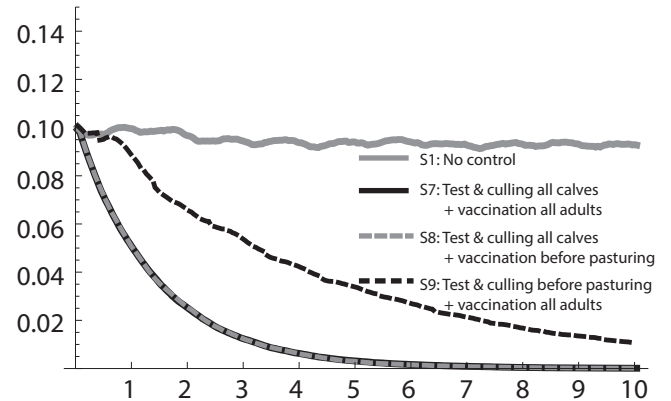
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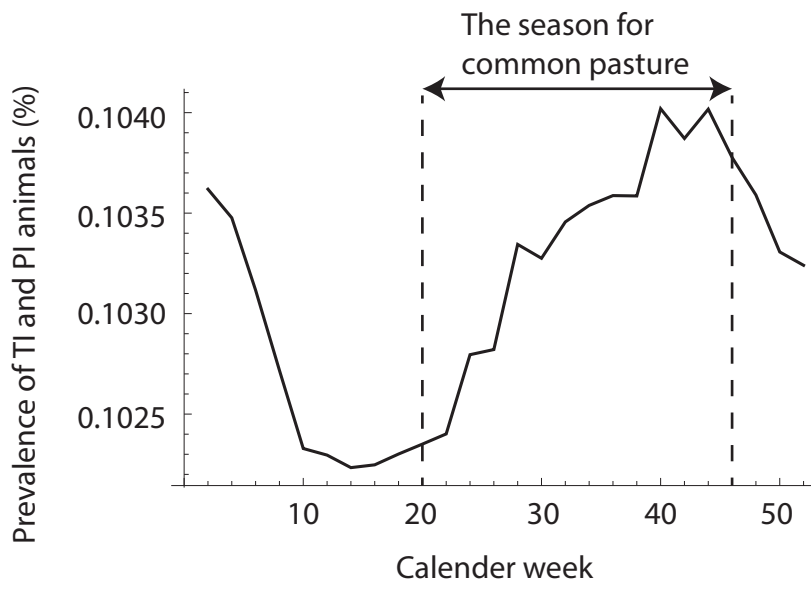
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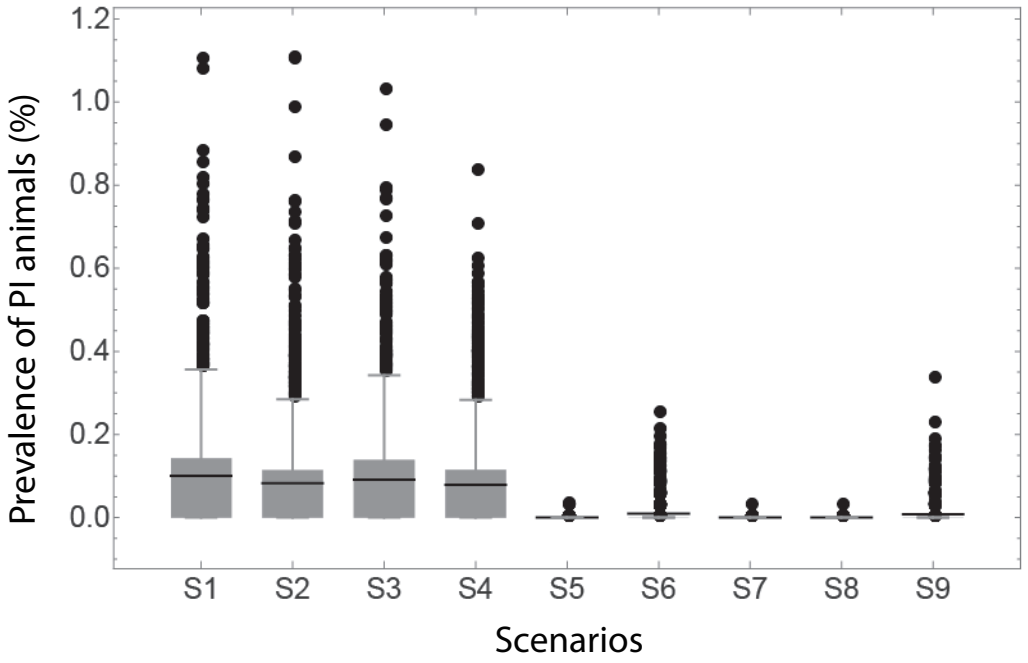
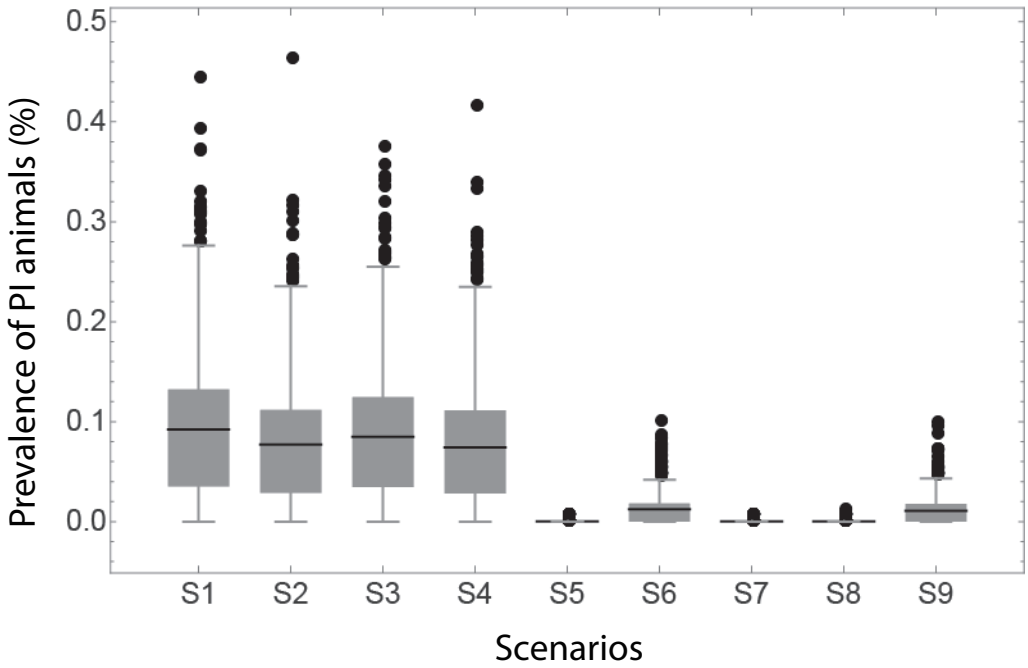
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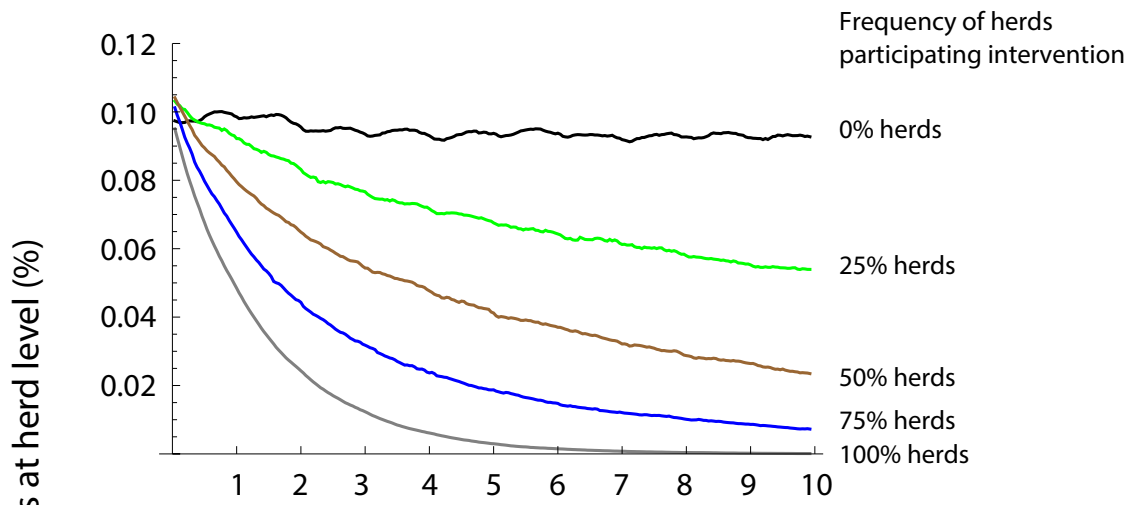
Time post infection (years)



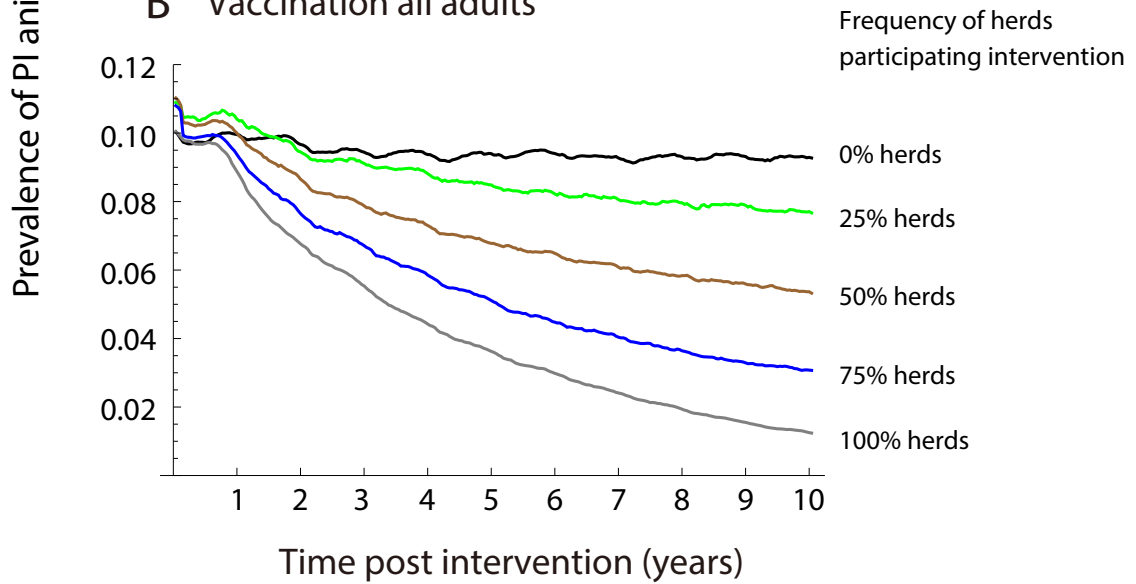
Sekiguchi et al. Figure 3

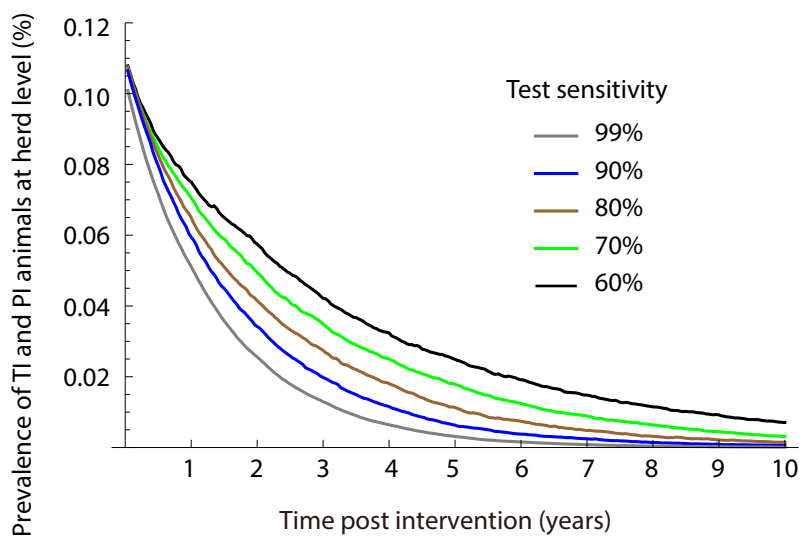
A**B**

A Test and culling all calves



B Vaccination all adults





Sekiguchi et al. Figure 6