Assessing the Mandatory Bovine Abortion Notification System in France Using Unilist Capture-Recapture Approach

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Abstract

The mandatory bovine abortion notification system in France aims to detect as soon as possible any resurgence of bovine brucellosis. However, under-reporting seems to be a major limitation of this system. We used a unilist capture-recapture approach to assess the sensitivity, i.e. the proportion of farmers who reported at least one abortion among those who detected such events, and representativeness of the system during 2006–2011. We implemented a zero-inflated Poisson model to estimate the proportion of farmers who detected at least one abortion, and among them, the proportion of farmers not reporting. We also applied a hurdle model to evaluate the effect of factors influencing the notification process. We found that the overall surveillance sensitivity was about 34%, and was higher in beef than dairy cattle farms. The observed increase in the number of notifying farmers from 2007 to 2009 followed an increase in the surveillance sensitivity in 2007/2008 and an increase in the proportion of farmers who detected at least one abortion in 2008/2009. These patterns suggest a raise in farmers’ awareness in 2007/2008 when the Bluetongue Virus (BTV) was detected in France, followed by an increase in the number of abortions in 2008/2009 as BTV spread across the country. Our study indicated a lack of sensitivity of the mandatory bovine abortion notification system, raising concerns about the ability to detect brucellosis outbreaks early. With the increasing need to survey the zoonotic Rift Valley Fever and Q fever diseases that may also cause bovine abortions, our approach is of primary interest for animal health stakeholders to develop information programs to increase abortion notifications. Our framework combining hurdle and ZIP models may also be applied to estimate the completeness of other clinical surveillance systems.

Introduction

In the context of increasing cross-border movements of people and growing international trade of animals and animal products, efficient and reliable surveillance systems are the basis to assess the actual disease situation in a country or region. Mandatory notification systems are essential in human and animal health as they allow health authorities to react promptly and control the spread of diseases. In France, clinical (or “passive”) surveillance covers 65 animal diseases that may cause important public health or economical impacts; among them, 60% are sporadic or currently exotic. Passive surveillance systems play an important role in early warning systems but their sensitivity in detecting cases has not been routinely evaluated [1].

The mandatory bovine abortion notification system, implemented in France in 1965, aims to detect as soon as possible any resurgence of bovine brucellosis [2]. This highly infectious zoonosis due to *Brucella abortus* (and less frequently to *B. melitensis* and *B. suis*) in cattle affects primarily the reproductive organs of infected animals. In particular, once introduced in a cattle herd, bovine abortion is the main clinical sign of the disease. Bovine brucellosis was endemic to French cattle population in the 1960’s but was then progressively eradicated by vaccination and test and cull management. France was declared as officially free from bovine brucellosis in 2005 [3]. Nevertheless, the recent outbreaks in Belgium and France (April 2012, [4]) reminded that the risk of introduction of bovine brucellosis still persists and emphasized the importance of an effective clinical surveillance system. In addition, the mandatory bovine abortion notification system might be useful for surveillance of other diseases causing bovine abortion. Thus, high abortion rate in ruminants may also be a signal of the introduction of Rift Valley Fever, an acute, zoonotic viral disease that currently circulates in numerous African countries [5]. Similarly, a clinical surveillance of Q fever has been recently implemented in ten departments (administrative units) in France to...
estimate the prevalence of this zoonosis in cattle and small ruminants herds [6].

In case of bovine abortion, European regulations require farmers to consult their veterinarian who report the abortion to veterinary services and sample the aborted cow for a serological analysis for Brucella [7,9]. Abortion is defined by national regulation as “the expulsion of the fetus or the calf, stillborn or dying less than 40 hours after being born” [7]. About 8 million reproductive cows distributed in 166,000 herds compose the French bovine stock [9]. For the last 5 years, 25 to 30,000 farmers participated in the national mandatory abortion notification system and 30 to 60,000 abortions were reported each year, which correspond to about 20% of the herds and 0.70 to 0.80% of the pregnant cows (considering a barrenness of about 5–10% [10]). However, a recent study of the delay between artificial insemination and calving in dairy cattle estimated that the rates of abortion occurring in mid-pregnancy and late pregnancy were about 6.4% and 5.1%, respectively [11]. “Usual” annual abortion risks have been cited elsewhere to vary from 3 to 5% once cows are about 6.4% and 5.1%, respectively [11]. “Usual” annual abortion risks have been cited elsewhere to vary from 3 to 5% once cows are beyond 42 days of pregnancy [12]. Although abortions are not always detected (it has been estimated that only 20 to 30% of abortions are detected visually [13]), the low proportion of cases reported to the system suggests that some abortions remain unreported.

Under-reporting of abortions in cattle may impede the early detection of incursions of bovine brucellosis or other infectious agents that may cause abortion. To our knowledge, no study has evaluated the completeness of the register. Estimating the proportion of under-reporting farmers, i.e. who detected at least one abortion but did not report any, and determining the factors that drive the decisions of farmers to report abortions are central goals for public health authorities to assess the performances and sensitivity of the surveillance system.

As often advocated in public health surveillance, data collected by a surveillance system may be analyzed by capture-recapture methods to take into account the fact that some events of interest are not observed. Unlist capture-recapture approach is implemented when data are collected by a single source of observations with repeated entries [14–16]. In our case, only herds with reported abortion(s) were “observed” by the mandatory bovine abortion notification system [17,18]. These data were incomplete because some farmers may have detected abortion(s) but did not report any. Multi-response or mixture models, including zero-inflated Poisson (ZIP) and hurdle models, provide flexible approaches to model the heterogeneity among farmers in the notification process [19].

Our objectives were to assess the sensitivity (i.e. the proportion of farmers who reported at least one abortion among those who detected such events) and representativeness of the mandatory bovine abortion notification system in France during 2006–2011, and to study the factors influencing the notification process. We applied a ZIP model to estimate the proportion of farmers who detected at least one abortion and among them, the proportion of farmers not reporting. We then quantified the effect of the factors influencing the farmers’ probability to report at least one abortion and the number of abortions reported by notifying farmers using a hurdle model.

### Materials and Methods

#### Data Sources and Study Population

We extracted data related to abortion notification data from the French national animal health information database SIGAL, including the herd identification number, dates of veterinarian visits (or dates of abortion), and pregnancy stages of the aborted cows. For all farmers, we extracted information about cattle farm location (department), animals (identification number, birth date, sex and breed), and animal movements (herd identification number, date, reason for entry [birth], reason for exit [death]) from the French National Cattle Register.

We considered five reproductive seasons between 2006 and 2011: reproductive season X started on August 1st of year X and ended on July 31st of year X+1 to follow the seasonality of calving, peaking in September/October for dairy cows and in March for beef cows. The study focused on all departments that reported at least one abortion per reproductive season since 2005. A department is a French administrative unit with a mean area of 5,800 km². For each department, we considered bovine farmers who owned at least one reproductive cow per reproductive season. A reproductive cow was defined as a female aged two years and over at the start of a reproductive season; thus, we included 98% of the calving records for each reproductive season. For each reproductive season, a cattle herd was characterized by its department, size, production type and the reported number of abortion(s). Herd size was calculated as the mean number of reproductive cows held per day. Three production types were defined according to the breeds of the animals that composed the herd: beef, dairy, and mixed (combining beef and dairy) cattle herd.

#### Multi-response Models

The data consisted on counts of abortions $Y_j$ reported by each farmer $i=1,2,...,N$ within a given reproductive season $j$. $Y_j$ was a random variable, which could take values $0,1,2,...$ for each herd. Herds were “observed” (or “captured”) by the mandatory bovine abortion notification system only when farmers reported abortion(s) (i.e. $Y_j>0$). Thus, the observed proportion of notifying farmers ($\Pr(Y_j>0)$) was defined as the ratio of the observed number of farmers who reported at least one abortion to the total number of farmers. However, farmers with no observation (referred as “non-notifying farmers” hereafter) included farmers who detected no abortion whether an abortion occurred or not and farmers who detected abortion(s) but did not report any. Even if abortion notification implied abortion occurrence, detection, and notification itself, we focused only on the detection and notification processes.

Preliminary analyses considering $Y_j$ as a simple random Poisson variable did not provide a good fit to the data because it did not account for the excess of zeros [19,20], due to the presence of farmers who detected no abortion. Therefore, we used multi-response models to account for zero-inflation and to evaluate separately the various processes generating the observations [19].

The ZIP model described $Y_j$ as the result of two random variables: the binary $(1/0)$ realization of the farmers’ probability to detect at least one abortion (unobserved process) $w_j$ and the number of abortion(s) reported by farmers who detected at least one abortion $y_j$ such that $Y_j=w_j \times y_j$ [21] (Figure S1). The probability for a farmer to detect at least one abortion $p_w$ was expressed by a logistic regression (link logit): $w_j \sim Bernoulli(p_w)$. The number of abortion(s) reported by farmers who detected at least one abortion was modeled by a Poisson regression (link log): $y_j \sim Poisson(\lambda_y)$, with mean $\lambda_y$.

For farms where no abortion was detected ($w_j=0$), the observed number of reported abortions was zero ($Y_j=0$). On the other hand, for farms where at least one abortion was detected ($w_j=1$), the number of reported abortion(s) was $y_j=0$ for farmers not reporting or $y_j>0$ for notifying farmers. Thus, among non-
notifying farmers \((Y_j = 0)\), the ZIP model differentiated farmers who did not detect any abortion \((w_j = 0)\) from under-reporting farmers, who detected \((w_j = 1)\) but did not report any abortion. Therefore, the probability for a farmer to report 0 abortion was calculated as \(P(Y_j = 0) = (1 - p_y) + p_y e^{-\lambda_y}\). The probability for a farmer to report \(k\) abortions \((k = 1, 2, \ldots)\) was

\[
P(Y_j = k) = p_y \frac{\lambda_y^k e^{-\lambda_y}}{k!}
\]

For each reproductive season \(j\), the proportion of farmers who detected abortion(s) \((Pr \text{op}_{Dj})\) was calculated as the ratio of the number of farmers who detected at least one abortion to the total number of farmers: \(Pr \text{op}_{Dj} = \frac{\sum p_y}{N}\). The proportion of under-reporting farmers \((Pr \text{op}_{URj})\) was calculated as the ratio of the number of farmers who detected but did not report abortion(s) to the total number of farmers who detected at least one abortion, \(Pr \text{op}_{URj} = \frac{\sum p_y e^{-\lambda_y}}{\sum p_y}\).

**Hurdle model.** The hurdle model described \(Y_j\) as the result of two random variables: the binary \((1/0)\) realization of the farmers’ probability to report at least one abortion \(w_j\) and the number of abortion(s) reported by notifying farmers \(y_j\), such that \(Y_j = w_j y_j\) \([21]\) (Figure S1). The probability for a farmer to report at least one abortion \(p_y\) was expressed by a logistic regression (link logit): \(w_j \sim \text{Bernoulli}(p_y)\). The number of abortion(s) reported by notifying farmers, i.e. farmers who had decided to “cross the hurdle” of the notification process, was modeled by a zero-truncated Poisson regression (link log): \(y_j \sim \text{Poisson}(\lambda_y)\), with mean \(\lambda_y\).

Once the farmer had decided to report at least one abortion \((w_j = 1)\), the number of reported abortion(s) \(y_j\) was positive \((y_j > 0)\). Thus, this model analyzed separately the “zero count” (i.e. non-notifying farmers, \(w_j = 0\)) and counts of one or more reported abortions (i.e. notifying farmers, \(w_j = 1\)) \([19, 22]\). With the hurdle model, the probability for a farmer to report 0 abortion was: \(P(Y_j = 0) = 1 - p_y\), and the probability for a farmer to report \(k\) abortions \((k = 1, 2, \ldots)\) was: \(P(Y_j = k) = \frac{p_y \lambda_y^k e^{-\lambda_y}}{1 - e^{-\lambda_y}} \frac{\lambda_y^k e^{-\lambda_y}}{k!}\).

**Inference on the Number of Bovine Abortion Notification(s) by Farmers**

**Covariates.** For each model, the Bernoulli and \((zero-truncated)\) Poisson responses could be predicted by a set of explanatory variables \([23, 24]\). We considered the production type of herd (dairy, beef, mixed), herd size (with categories based on quartiles) and reproductive season as categorical variables in the logistic and \((zero-truncated)\) Poisson regressions. In the \((zero-truncated)\) Poisson regression, the herd size was also included as an offset in order to take into account the suspected linear relationship between the number of abortion notifications and herd size (mechanistic effect). The location of the herd (department) and the farmer were studied as random effects. The R MCMCglm function package used to run Bayesian simulations \([25]\) included automatically an additive over-dispersion parameter \(e\) (or “residuals”) in both linear predictors as a random effect, for which a residual variance was estimated \([26, 27]\).

We ran the ZIP model for each reproductive season \(j\) separately in order to estimate for season-specific \(Pr \text{op}_{Dj}\) and \(Pr \text{op}_{URj}\). The ZIP models included covariates in the logistic and Poisson regressions, respectively, as

\[
\text{Logit}(p_y) = \beta_0 + \beta_1 \times \text{Department}_i + \beta_2 \times \text{Production Type}_i
\]

and

\[
\text{Log}(\lambda_y) = \beta_0 + \beta_1 \times \text{Department}_i + \beta_2 \times \text{Production type}_i + \log(\text{Size}_i) + \beta_3 \times \text{Size}_i + \epsilon_y,
\]

where \(\beta_0\) was the intercept, \(\beta\) were the covariate coefficients, \(b\) was the random effect, and \(\epsilon\) corresponded to the residuals.

We ran a single hurdle model including reproductive season as a fixed effect in order to study a potential between-seasons variation in the probability for a farmer to report at least one abortion and in the number of abortion(s) reported by notifying farmers. The hurdle model included covariates in the logistic and truncated-Poisson regressions, respectively, as

\[
\text{Logit}(p_y) = \beta_0 + \beta_1 \times \text{Farmer}_i + \beta_2 \times \text{Production type}_i + \beta_3 \times \text{Reproductive season}_i + \epsilon_y
\]

and

\[
\text{Log}(\lambda_y) = \beta_0 + \beta_1 \times \text{Farmer}_i + \beta_2 \times \text{Production type}_i + \beta_3 \times \text{Size}_i + \beta_4 \times \text{Reproductive season}_i + \epsilon_y
\]

For the logistic regression, we calculated the odds ratio (OR) of each covariate as \(OR = \exp(\beta)\). Similarly, for the \((zero-truncated)\) Poisson regressions, we calculated the relative risk (RR) for each covariate, as \(RR = \exp(\beta)\) \([28]\).

**Random sampling.** Due to the high number of herds included in the study population \((n = 94,640)\), models were implemented using a stratified randomly selected sample of herds continuously monitored from 2006 to 2011 with department as strata. We selected 10% \((n = 9,462)\) of the entire population of herds for the ZIP model and 5% \((n = 4,732)\) for the hurdle model, to maintain precision while reducing computing time. We checked the robustness of the results by performing a second round of estimations for the selected models with another sample.

**Priors distribution.** We used proper uninformative distributions for hyperparameters, except for the offset (Table 1) \([27, 29]\). As proposed by Hadfield \([24]\), the variance of the over-dispersion parameter integrated in the logistic part of the ZIP and hurdle models was set to 1 and the fixed effects and variances were rescaled. To investigate the sensitivity of the posterior inference to the prior information, we fitted the selected model using different prior distributions. We assessed the relative changes in the posterior distributions of the parameters by multiplying by 2 the scale of the half-Cauchy prior distributions and by 10 the parameter of the inverse-gamma prior distribution.

**Models implementation.** A total of 800,000 samples were run including a burn-in period of 500,000 and we used a thinning sample of 100. Two simulation chains were obtained and convergence was assessed using the Gelman-Rubin convergence criterion \([30]\). Models were selected considering the statistical
significance of covariates and their convergence. We checked model fit by studying the discrepancy between the observed and predicted numbers of farmers who reported \( k \) abortion(s) and assessing the posterior mean of the standardized residuals [31].

Posterior distributions of \( \text{PropD} \) and \( \text{PropUR} \) were obtained from each ZIP model. Differences in posterior means among reproductive seasons and types of production were evaluated with a z-test using the significance level \( \alpha = 0.05 \). We approximated standard errors by dividing the 95% credible intervals by 3.92 (2 \( \times \sigma_{0.05} \)). P-values were adjusted using the Bonferroni method [32].

Results

Population Characteristics
The study population included 94,640 herds from 37 departments (which represented 78.7% of the 120,213 herds registered during the 2006–2011 period), among which 36.5 to 40.0% were beef cattle, 21.1 to 25.3% dairy cattle, and 38.2 to 39.1% mixed cattle depending on the reproductive season. During the study period, 25% of the herds held less than 18 reproductive cows on average, 50% less than 38, and 75% less than 60. On average, there were 37 ± 3 (sd), 39 ± 24, and 52 ± 38 reproductive cows in beef, dairy and mixed cattle, respectively. Between 20.0% (in the 2010/2011 reproductive season) and 26.1% (in 2007/2008) of farmers reported at least one abortion per reproductive season (Figure 1). Among notifying farmers, an average of 60.5% (min-max: 58.3–62.0%) reported one abortion and 21.9% (min-max: 21.3–22.7%) two abortions depending on the reproductive season. 76.5% of reported abortions occurred after 6 months of pregnancy (among the 92.4% of reported abortions for which the information was available). Characteristics of the herds included in the two subsamples used to run the ZIP and hurdle models did not differ from the study population characteristics (Table S1).

Inferences from the ZIP Model
We found a good fit between the observed number of farmers who reported \( k \) abortion(s) and the ZIP-model predictions for all reproductive seasons (Figure S2). Model predictions were robust to a change in prior distributions and when analyzing another subsample (results not shown).

\( \text{PropD} \) varied from 60.6% to 80.1% depending on the reproductive season and \( \text{PropUR} \) from 59.5% to 70.0% (Figure 1). For all reproductive seasons, \( \text{PropD} \) in beef cattle (about 58%) was lower than in dairy cattle (about 84%) but similar to mixed cattle (Tables 2 and 3). In contrast, \( \text{PropUR} \) was lower for dairy (about 59%) than beef cattle farmers (about 79%; Table 3). Overall, \( \text{PropD} \) was significantly higher during the 2008/2009 reproductive season compared to other seasons and \( \text{PropUR} \) was significantly lower in 2007/2008 than during other seasons.

Table 1. Prior distributions of the hyperparameters for the ZIP and hurdle models.

| Hyperparameter | Submodel | Prior distribution |
|----------------|----------|--------------------|
| Fixed effects (except the offset) | Logistic regression | \( N(0, 10^{10}) \) |
| Offset | (truncated-)Poisson regression | \( N(0, 10^{10}) \) |
| Standard deviation of the “Department” and Herd effects | Logistic regression | Half-Cauchy(0, 5) |
| Variance “Over-dispersion parameter” | Logistic regression | Fixed to 1 |
| | (truncated-)Poisson regression | Inv(0.001, 0.001) |

\( N \) refers to a normal distribution and \( \text{Inv} \) to an inverse gamma distribution. Poisson regression refers to the ZIP model and truncated-Poisson regression to the hurdle model.

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![Figure 1. Proportion of farmers who detected, reported or failed to report abortions during the 2006–2011 period.](https://doi.org/10.1371/journal.pone.0063246.g001)
reproductive seasons (Figure 1). The number of abortions reported by farmers who detected at least one abortion was influenced by production type and herd size. Among farmers who detected at least one abortion, farmers with dairy cattle reported on average twice more abortions that those with beef cattle (Table 4). When adjusting for the proportional increase in the number of abortions with herd size (using an offset in the model), farmers with 18 to 60 reproductive cows tended to report more abortions than those with less than 10 reproductive cows; in contrast, farmers with more than 60 reproductive cows reported as many abortions as those with less than 18 reproductive cows. The residual variance of the over-dispersion parameter was higher than the variance of the department (Table 4).

Discussion

Inferences from the Hurdle Model

The hurdle model provided a good fit to the observed number of farmers who reported k abortion(s) (Figure S2). Model predictions were robust to a change in prior distributions and when analyzing another subsample (results not shown).

The probability for a farmer to report at least one abortion was higher for dairy than for beef cattle herds (OR = 3.94 [3.40–4.54]), and increased with the number of reproductive cows (Table 5); it was higher in the 2007/2008 reproductive season than in other seasons (Table 5). The number of abortion notifications was 1.77 [1.55–2.00] higher for notifying farmers with dairy than beef cattle herds (Table 6). When adjusting for the proportional increase in the number of abortions with herd size (using an offset in the model), the ability to report abortions was equal among farmers whatever the size of their herds (Table 6). The number of abortion notifications by notifying farmers was higher in 2007/2008 than in other reproductive seasons (Table 6).

Table 2. Odds ratios (OR) for the probability of detecting at least one abortion in cattle.

| Reproductive season | Production type (OR) | Variance “Department” |
|---------------------|----------------------|-----------------------|
|                     | Dairy                | Mixed                 |
| 2006/2007           | 4.9 [1.52–30.89]     | 1.3 [0.78–2.21]       | 0.15 [0.0–0.39] |
| 2007/2008           | 2.49 [1.49–4.04]     | 1.32 [0.9–1.9]        | 0.31 [0.07–0.64] |
| 2008/2009           | 8.88 [1.5–146.06]    | 1.17 [0.54–3.02]      | 0.95 [0.05–2.29] |
| 2009/2010           | 3.53 [1.57–8.09]     | 1.44 [0.69–2.52]      | 0.2 [0.0–0.51]   |
| 2010/2011           | 5.75 [2.46–32.4]     | 1.84 [1.09–2.85]      | 0.35 [0.02–0.81] |

The probability for a farmer to detect at least one abortion (p) was modeled by the logistic regression of the ZIP model. 95% credible intervals are mentioned in square brackets and bolded values indicate significant differences (i.e. credible interval not including 1.00). Beef cattle herds were considered as the reference for each reproductive season. Variance corresponded to the amount of variation among departments.

Unilist Capture-recapture Assumptions

Unilist capture-recapture approach assumes capture homogeneity and independence between capture and recapture [18]. In our study, validating the homogeneity assumption required that the number of detected abortions and the probability of reporting them were the same between herds presenting the same set of covariates. As bovine abortions are generally sporadic events, we expected only few variations in the number of detected abortions between herds. But because the probability of detecting abortion(s) in a farm certainly depends on the frequency of contacts between farmers and their herds, we included the production type as a covariate in both models. In addition, we considered potential variation in the probability of reporting abortion(s) among farmers by including an over-dispersion parameter in the ZIP model and a random effect “farmer” in the hurdle model. Secondly, in the capture-recapture approach the reporting probabilities of successive abortions within a cattle herd are assumed to be independent. This assumption was admissible as most of abortions are sporadic events. However, the dependence between captures in herds exposed to epizootic abortions, such as observed during neosporosis outbreaks, may cause an underestimation of the proportion of under-reporting farmers [18,33].

Factors Driving Farmers’ Decision to Report Abortions

Our analysis predicted that, during each reproductive season and within the 37 departments included in the study, 19.9 to 39.4% of farmers did not detect any abortion in their herd and among farmers who did detect abortions, 59.5 to 70.0% did not report any. This high proportion of under-reporting farmers is surprising as the veterinarian visit in case of abortion is financed by veterinary services in France. We suspect that the absence of known brucellosis outbreak during 2006–2011 did not encourage farmers to report clinical suspected cases. In addition, farmers may not get worried about the economical impact of abortions as long as these are sporadic: farmers may contact their vet for a diagnosis only if the number of abortions goes over a “threshold” that may vary from one farmer to another. However, our results indicate some variations among departments in the probability of abortion notification likely because the mandatory bovine abortion notification system is implemented at the department level and some local animal health associations encourage farmers to diagnose other diseases (in addition to brucellosis) that may have an economic impact.

Our analysis indicated that the probability for a farmer to report at least one abortion was lower for beef that for dairy cattle herds: the proportion of farmers who detected at least one abortion was lower for beef than dairy cattle herds, and farmers with beef cattle...
Variations among Reproductive Seasons

Our results indicated that the proportion of non-notifying farmers was lower in 2007/2008 compared to other reproductive seasons. Although the predicted proportion of farmers who detected at least one abortion did not change, we found that farmers were more prone to report detected abortions during that reproductive season. However, in 2008/2009, the proportion of non-notifying farmers had increased again, in spite of a higher number of farmers who detected at least one abortion. Several events during these years may explain these findings. First, the implementation in 2005 of an annual official vet visit to inform farmers about the abortion issue may have increased farmers’ awareness of notification in 2007/2008. Besides, following the introduction of Bluetongue Virus (BTV) in 2006, the number of cases started to rise dramatically with 7,607 outbreaks reported in the 2007/2008. The coverage by national medias about the suspected role of BTV in bovine abortions may have also encouraged abortion notification [34,35]. In 2008/2009, the proportion of farmers who detected at least one abortion increased while the number of BTV outbreaks peaked (12,243 reported outbreaks) [36], suggesting a direct impact of the virus on abortion occurrence [11]. The concurrent increase of the proportion of under-reporting farmers (Figure 1) may be explained by the fact that the role of vaccination against BTV was debated at that time, and tensions between farmers and vets may have discouraged farmers to notify. Afterwards, the decline in BTV cases to 13 cases in 2009/2010 and none in 2010/2011 may explain the lower proportion of farmers who detected at least one abortion those years.

Advantages and Limits of the Study

Combining hurdle and ZIP models provides a comprehensive framework to examine the sensitivity of the bovine abortion notification system and the factors influencing the farmers’ decisions to report these events. These two models complement

| Reproductive season | Proportion | Beef | Dairy | Mixed |
|---------------------|------------|------|-------|-------|
| 2006/2007 detection | 59.3 [49.6–69.5] | 86.6 [78.0–99.5] | 66.3 [59.3–74.1] |
| 2007/2008 detection | 78.1 [74.9–81.5] | 59.9 [55.8–64.4] | 60.0 [56.1–63.9] |
| 2008/2009 detection | 77.3 [70.5–76.6] | 52.3 [48.8–55.9] | 53.1 [50.5–56.4] |
| 2009/2010 detection | 74.0 [63.2–87.5] | 93.8 [85.5–99.9] | 78.1 [69.2–88.4] |
| 2010/2011 detection | 82.6 [79.8–85.2] | 62.1 [58.8–65.2] | 63.7 [59.7–67.6] |

The proportion of farmers who detected at least one abortion (\(\text{PropAs}\)) was the ratio of the number of farmers who detected at least one abortion to the total number of farmers. The proportion of under-reporting farmers (\(\text{PropUR}\)) was the ratio of the number of farmers who detected but did not report abortion to the number of farmers who detected at least one abortion. Both proportions were estimated from the ZIP model. 95% credible intervals are mentioned in square brackets.

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| Variables | 2006/2007 | 2007/2008 | 2008/2009 | 2009/2010 | 2010/2011 |
|-----------|-----------|-----------|-----------|-----------|-----------|
| Production type (RR) Dairy | 1.83 [1.43–2.32] | 1.85 [1.53–2.22] | 2.17 [1.79–2.67] | 2.08 [1.60–2.84] | 2.04 [1.61–2.68] |
| Mixed | 1.51 [1.20–1.87] | 1.5 [1.27–1.77] | 1.69 [1.38–2.07] | 1.59 [1.19–2.14] | 1.58 [1.24–2.09] |
| Herd size (RR) (18–38) | 1.18 [0.97–1.46] | 1.24 [1.03–1.52] | 1.57 [1.25–1.95] | 1.36 [1.06–1.74] | 1.34 [1.05–1.70] |
| (38–60) | 1.09 [0.90–1.35] | 1.25 [1.04–1.5] | 1.44 [1.15–1.78] | 1.38 [1.10–1.76] | 1.24 [0.98–1.57] |
| > 60 | 0.94 [0.76–1.15] | 1.02 [0.84–1.23] | 1.23 [0.99–1.54] | 1.12 [0.89–1.43] | 1.05 [0.83–1.31] |
| Variance “Department” | 0.11 [0.04–0.19] | 0.07 [0.03–0.13] | 0.17 [0.06–0.31] | 0.1 [0.03–0.17] | 0.11 [0.04–0.20] |
| Variance “Overdispersion parameter” | 0.62 [0.48–0.78] | 0.49 [0.38–0.59] | 0.71 [0.53–0.85] | 0.66 [0.50–0.83] | 0.7 [0.50–0.95] |

The number of abortion notification(s) by farmers who detected at least one abortion (\(i\)) was modeled by the Poisson regression of the ZIP model. 95% credible intervals are mentioned in square brackets and bolded values indicate significant differences (i.e. credible interval not including 1.00). Beef cattle farmers with less than 18 reproductive cows during the 2007/2008 reproductive season were considered as the reference. Variance corresponded to the amount of variation associated with the corresponding random variable.

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each other: by accounting for the latent heterogeneity in detection, the ZIP model provided an unbiased estimate of the proportion of under-reporting farmers; on the other side, the hurdle model allowed to investigate the decision-making process of abortion notification.

We were not able to distinguish the relative influence of the probabilities of abortion occurrence, detection and notification. Knowing the frequency of abortion occurrence in cattle is challenging because the absence of registered calf for a reproductive cow during a given reproductive season may result from a fertility problem, or failure of the artificial insemination (AI) or natural breeding. Further analyses combining information from additional sources such as the dates of AI, mating period or intervals between calving would help evaluating the frequency of abortions in cattle. Moreover, our analysis does not enable to estimate the proportion of herds where abortions occurred: our study period. Difficulties in keeping actors participating in a surveillance system have already been observed in public health notification systems: under-reporting of notifiable diseases was found to be associated with the corresponding surveillance system compared to dairy cattle farmers. Given the high proportion of under-reporting farmers, will the surveillance system detect outbreaks of brucellosis or other critical diseases? Our results suggest that the early detection of any introduction or resurgence of brucellosis may be difficult. In particular, our findings concern all farms in activity since 2005, and we may expect a higher proportion of under-reporting farmers among those who started or stopped their activity during the study period. Difficulties in keeping actors participating in a notification system have already been observed in public health surveillance systems: under-reporting of notifiable diseases was estimated to reach 90% in some cases [39] and the knowledge, attitudes and beliefs of various actors were the main drivers of the decision to report or not [39,40]. Nevertheless, in contrast to active surveillance which is restricted to a certain time frame and is resource-consuming, passive surveillance presents the advantages of being ongoing, cost-saving and prompt to detect an outbreak. In 2012, one of the two brucellosis outbreaks reported in France was detected thanks to clinical surveillance. In Great Britain, a simulation study of brucellosis spread in cattle under several testing regimes found that suppressing abortion notifications would have a major effect on the rate of spread of infection before detecting a brucellosis outbreak [41]. Beside early detection, clinical surveillance has been suggested to be one of the main way to identify secondary outbreaks once the disease was introduced because disease awareness of animal health actors increases the report of suspect cases [35].

| Variable                  | Estimation     |
|---------------------------|----------------|
| Production type (OR)      | Dairy 3.94 [3.40–4.54] |
|                          | Mixed 2.27 [2.01–2.59] |
| Size (OR)                 | [18–38] 6.49 [5.35–7.79] |
|                          | [38–60] 12.16 [10.02–14.70] |
|                          | ≥60 16.13 [13.33–19.70] |
| Reproductive season (OR)  | 2006/2007 0.79 [0.71–0.88] |
|                          | 2008/2009 0.81 [0.72–0.90] |
|                          | 2009/2010 0.61 [0.55–0.69] |
|                          | 2010/2011 0.61 [0.54–0.68] |
| Variance “Farmers”        | 1.98 [2.19–1.77] |

The probability for a farmer to report at least one abortion (\( p \)) was modeled by the logistic regression of the hurdle model. 95% credible intervals are mentioned in square brackets and bolded values indicate significant differences (i.e. credible interval not including 1.00). Beef cattle farmers with less than 18 reproductive cows during the 2007/2008 reproductive season were considered as the reference. Variance corresponds to the amount of variation among farmers.

| Variable                  | Estimation     |
|---------------------------|----------------|
| Production type (RR)      | Dairy 1.77 [1.55–2.00] |
|                          | Mixed 1.45 [1.28–1.63] |
| Size (RR)                 | [18–38] 1.06 [0.73–1.51] |
|                          | [38–60] 0.93 [0.64–1.34] |
|                          | ≥60 0.72 [0.49–1.02] |
| Reproductive season (RR)  | 2006/2007 0.85 [0.76–0.94] |
|                          | 2008/2009 0.87 [0.79–0.97] |
|                          | 2009/2010 0.83 [0.74–0.92] |
|                          | 2010/2011 0.79 [0.71–0.89] |
| Variance “Farmers”        | 0.27 [0.21–0.33] |
| Variance “Overdispersion parameter” | 0.26 [0.19–0.32] |

The number of abortion notification(s) by farmers who reported at least one abortion (\( L_i \)) was modeled by the zero-truncated Poisson regression of the hurdle model. 95% credible intervals are mentioned in square brackets and bolded values indicate significant differences (i.e. credible interval not including 1.00). Beef cattle farmers with less than 18 reproductive cows during the 2007/2008 reproductive season were considered as the reference. Variance corresponded to the amount of variation associated with the corresponding random variable.

Assessing the Mandatory Bovine Abortion Notification System

The sensitivity of the mandatory bovine abortion notification system varied from 30.0 to 40.5% depending on the reproductive season [37]. This surveillance system is not fully representative as it does not accurately describe the proportion of farmers detecting bovine abortions over time and its distribution in the population by production type [38]: the proportion of beef cattle farmers is under-represented in the mandatory bovine abortion notification system compared to dairy cattle farmers. Given the high proportion of under-reporting farmers, will the surveillance system detect outbreaks of brucellosis or other critical diseases? Our results suggest that the early detection of any introduction or resurgence of brucellosis may be difficult. In particular, our findings concern all farms in activity since 2005, and we may expect a higher proportion of under-reporting farmers among those who started or stopped their activity during the study period. Difficulties in keeping actors participating in a notification system have already been observed in public health surveillance systems: under-reporting of notifiable diseases was estimated to reach 90% in some cases [39] and the knowledge, attitudes and beliefs of various actors were the main drivers of the decision to report or not [39,40]. Nevertheless, in contrast to active surveillance which is restricted to a certain time frame and is resource-consuming, passive surveillance presents the advantages of being ongoing, cost-saving and prompt to detect an outbreak. In 2012, one of the two brucellosis outbreaks reported in France was detected thanks to clinical surveillance. In Great Britain, a simulation study of brucellosis spread in cattle under several testing regimes found that suppressing abortion notifications would have a major effect on the rate of spread of infection before detecting a brucellosis outbreak [41]. Beside early detection, clinical surveillance has been suggested to be one of the main way to identify secondary outbreaks once the disease was introduced because disease awareness of animal health actors increases the report of suspect cases [35].
To our knowledge, this study is the first to quantify the proportion of under-reporting farmers and evaluate the factors that influence the notification process of a specific health-related event. As underlined by the Centre for Disease Control, sensitivity and representativeness of public health surveillance systems are the major attributes of the system to assess when a surveillance system is evaluated [31]. Our approach may be applied to other clinical surveillance systems to help identify the ways to increase actors’ awareness and improve the notification of suspected cases.

Supporting Information

Figure S1 Graphical representation of the two-processes modeled by the ZIP and hurdle multi-response models. The ZIP model analyzed notification data (i.e., counts of abortions \( Y_{ij} \) reported by each farmer \( i \) within a given reproductive season \( j \)) as the result of two processes: 1) an unobserved probability for farmers to detect at least one abortion (Bernoulli regression) and 2) the number of notifications by farmers who detected at least one abortion (Poisson regression). We estimated the proportion of under-reporting farmers (\( Pr(P_{UR}) \)) as the ratio of the number of farmers who detected but did not report abortions to the number of farmers who detected at least one abortion. The hurdle model considered 1) the probability for farmers to report at least one abortion (Bernoulli regression) and 2) the number of notifications by notifying farmers (zero-truncated Poisson regression).

Figure S2 Posterior realization of the discrepancy statistic function \( D_K \) under the ZIP and hurdle models.
The predictions \( T_{ij}(y_{pred}) \) from the ZIP and hurdle models were calculated for \( m = 1,2,\ldots,3000 \) simulated samples. The discrepancy \( D_{K(m)} \) between the observed \( T_{ij}(y) \) and the predicted number of farmers who reported \( k \) abortion(s) \( T_{ij}(y_{pred}) \) was calculated as

\[
P_{ij}(m) = \frac{T_{ij}(y) - T_{ij}(y_{pred})}{n}, \quad n \text{ the number of farms per reproductive season [31].}
\]

The model underestimated the number of farmers if \( D_{K(m)} \) was positive and overestimated it if \( D_{K(m)} \) was negative. Figures present the posterior realization of \( D_K \) under the ZIP model for the reproductive season 2007/2008 (plots were similar for other reproductive seasons) and under the hurdle model.

(DOCX)

Table S1 Comparisons of the observed probability of reporting abortion(s) and herd characteristics between the samples used for the ZIP and hurdle models analyses (\( n = 9,462 \)) and the original data set (\( n = 94,610 \)). Chi square tests comparing distributions between each subsample and the original data set were not significant.

(XLSX)

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Author Contributions

Conceived and designed the experiments: AB EG VH DC EM PH. Performed the experiments: AB. Analyzed the data: AB. Contributed reagents/materials/analysis tools: TV JLV. Wrote the paper: AB.

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