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EDITED BY

Elizabeth Anne Jessie Cook,
International Livestock Research
Institute (ILRI), Kenya

REVIEWED BY

Kebede Amenu,
International Livestock Research
Institute, Ethiopia
Gezahegn Alemayehu Ayalew,
International Livestock Research
Institute, Ethiopia
Judy Bettridge,
University of Greenwich,
United Kingdom
Wudu Temesgen Jemberu,
International Livestock Research
Institute, Ethiopia

*CORRESPONDENCE

Andrew R. Peters
andy.peters@ed.ac.uk

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Patterns of mortality in domesticated ruminants in Ethiopia

Giles T. Innocent¹, Ciara Vance², David A. Ewing¹,
Iain J. McKendrick¹, Solomon Hailemariam³,
Veronica R. Nwankpa⁴, Fiona K. Allan², Christian Schnier² and
Andrew R. Peters^{2*}

¹Biomathematics and Statistics Scotland (BioSS), Edinburgh, United Kingdom, ²Centre for Supporting Evidence Based Interventions-Livestock, Royal (Dick) School of Veterinary Studies, University of Edinburgh, Midlothian, United Kingdom, ³Private Consultant, Addis Ababa, Ethiopia, ⁴Department of Microbial, Cellular and Molecular Biology, Addis Ababa University, Addis Ababa, Ethiopia

Background: Premature death of livestock is a problem in all ruminant production systems. While the number of premature ruminant deaths in a country is a reasonable indicator for the nation's health, few data sources exist in a country like Ethiopia that can be used to generate valid estimates. The present study aimed to establish if three different data sets, each with imperfect information on ruminant mortality, including abortions, could be combined into improved estimates of nationwide mortality in Ethiopia.

Methods: We combined information from a recent survey of ruminant mortality with information from the Living Standards Measurement Study and the Disease Outbreak and Vaccination Reporting dataset. Generalized linear mixed and hurdle models were used for data analysis, with results summarized using predicted outcomes.

Results: Analyses indicated that most herds experienced zero mortality and reproductive losses, with rare occasions of larger losses. Diseases causing deaths varied greatly both geographically and over time. There was little agreement between the different datasets. While the models aid the understanding of patterns of mortality and reproductive losses, the degree of variation observed limited the predictive scope.

Conclusions: The models revealed some insight into why mortality rates are variable over time and are therefore less useful in measuring production or health status, and it is suggested that alternative measures of productivity, such as number of offspring raised to 1 year old per dam, would be more stable over time and likely more indicative.

KEYWORDS

Ethiopia, ruminant, cattle, sheep, goat, mortality, disease, abortion

Introduction

Livestock mortality is of great importance globally, and especially in low- and middle-income countries (LMICs) where livestock often have multiple roles. Livestock mortality has typically been considered one of the most important measures of population dynamics and productivity of livestock production systems (1), therefore high mortality risks can be considered a significant constraint in traditional livestock systems in sub-Saharan Africa (sSA) (2). Stock lost is a loss in nutrition, livelihood, and genetics, as well as lost wealth (3).

In Ethiopia, there is consistently low ruminant productivity and profitability (4). Between 2005 and 2015, average annual mortality in cattle and small ruminants was reported to be 7 and 20%, respectively (5, 6). Infectious diseases are frequently reported in household surveys as being major constraints on smallholder productivity (7) and annual disease-associated losses are estimated at US \$150 million (8). The rate of abortion in Ethiopian cattle is reported to range from 2.2 to 29%, varying across regions (9–13), where an abortion rate of 2–5% per annum is considered to be expected (14). In central Ethiopia, 57% of farmers have reported abortions in their flocks (15), with the reported mean annual abortion/stillbirth rate for Ethiopian flocks ranging from 3 to 36% over different ecosystems (16–18). Where the rate of abortion is between 2 and 5%, it is suggestive of underlying endemic disease, but when the rate exceeds this, it is important to investigate likely causes (19). Overall, information on small ruminant abortions in Ethiopia is considered lacking (20).

Smallholder production systems in Ethiopia are broadly classified into three categories: mixed crop-livestock, pastoral and agropastoral systems. Mixed crop-livestock farming is the predominant system in Ethiopia, combining crop cultivation and livestock production. There is an emphasis on crop cultivation, with low numbers of livestock per household. For example, in Central Ethiopia, the average number of livestock per household is 4.5 cattle, 1.1 sheep and 0.5 goats (21). Pastoral and agropastoral livestock production is the second most practiced system, predominantly in southern and eastern Ethiopia. There is no crop production in pastoral farming, and agropastoralism involves only limited crop and predominantly livestock production (22). Typically, pastoralists own an average of 5–29 cattle and 42–168 sheep and goats (23).

Research and development of animal health projects and interventions in Ethiopia focus on transboundary or zoonotic diseases and those which affect trade, with relatively little attention given to endemic diseases, despite their negative effects on production, food security and livelihoods (24). Indeed, there are limited quantitative data on the causes of ruminant losses in sSA which constrains the development and application of interventions to improve livestock health and productivity. The Center for Supporting Evidence Based

Interventions (SEBI), University of Edinburgh, was established to mobilize and apply evidence and data on livestock health and productivity from LMICs. One of their initial programmes was to establish mortality rates and major causes of mortality in cattle and small ruminants, in Ethiopia, Nigeria and Tanzania. Outcomes have been published for studies carried out in Nigeria (25) and Tanzania (26). This study aimed to establish if three different datasets, each with imperfect information on ruminant mortality, including abortions, could be combined into improved estimates of nationwide mortality. These estimates could then be used to monitor changes in the nation's herd health over time.

Materials and methods

Ethical approval

No animals were involved in this study. Verbal informed consent was obtained from each participant prior to commencement of the survey; all chose to participate. They were informed that their participation was voluntary, that their responses would be kept confidential and no names or identifying information linking them to the survey would be disclosed. In addition, they were assured that they were free to terminate their participation at any time. Surveys were read out to participants in the local language and responses translated into English by local translators.

At the outset of the study, ethical review was unintentionally omitted, however full details and supporting documentation of the study were submitted for retrospective ethical review on 21st February 2022 by the Human Ethical Review Committee, University of Edinburgh, the outcome of which stated that had the study been reviewed prior to commencement, ethical approval would likely have been granted.

Datasets

A total of three datasets were used in the study: a Farmer Survey (FS), the Living Standards Measurement Study (LSMS) from the World Bank, and the Disease Outbreak and Vaccination Reporting (DOVAR) dataset. Each dataset contained data on number of deaths, state, and species. Species included were cattle, sheep and goats or 'small ruminants'. Datasets also had varied additional information (Table 1), however, these were the only data that were consistent for all three datasets. The term 'herd' is used, hereafter, to refer to the varied reporting units, including 'households', 'holdings', 'herds' and 'flocks'.

For numerical reasons, we aggregated sheep and goats in a single small ruminant (SR) class. Due to data availability, we were only able to analyse abortions in the FS.

TABLE 1 Data available in each dataset.

Dataset	Parameters available
Farmer survey (FS)	State, species, disease, total number, number dead, production system, ecosystem, age, sex
Living Standards Measurement Study (LSMS)	Year, state, species, total number, number dead, rural/urban
Disease Outbreak and Vaccination Reporting (DOVAR)	Year/month, state, species, disease, number of outbreaks, number dead, number at risk

Data were sparse and there was a lack of independence of potential explanatory variables, such as Age, Sex, Disease, State, Ecosystem and Production System. Statistical models did not converge when 'Disease' was included as a variable, therefore disease variables were recoded and grouped as shown in Table 2.

Farmer survey

Farmers were recruited in purposively selected zones, districts and kebele in all nine regional states¹ and in Dire Dawa city administration, based on livestock population, and included different production systems and agro-ecological zones (AEZ). Production systems were mixed crop-livestock, pastoral and agropastoral, as already described, as well as semi-intensive, where livestock are enclosed during the day and housed at night (2). A total of 17 zones were selected (average of two zones per region, depending on the size of the region), and one district per zone ($n = 17$). Two kebele per district were selected, based on accessibility and ruminant production ($n = 34$). Approximately 10–12 households were selected per kebele, resulting in a total of 382 households for sampling. Households or producers (farmers, pastoralists and urban producers) were selected primarily based on their availability and accessibility to the study teams.

Structured questionnaire surveys (Supplementary material) were designed by SEBI and conducted with farmers between October and December 2017. Five study teams conducted the surveys with farmers and pastoralists, collecting information on demographics, causes of mortality and reproductive performance losses in cattle and small ruminants.

Herd data collection categorized cattle and small ruminants by age and sex. Cattle ages were classified as young stock (≤ 2 years), juveniles ($> 2-3$ years) and adults ($4+$ years). Small ruminants were grouped as < 3 months, 3 months - 1 year, and > 1 year. Information captured for the previous year (2016)

¹ Since the surveys were conducted, a tenth regional state has been created.

TABLE 2 Grouping of disease variables.

Original variable	Recoded variable
Anthrax	Infectious
Babesiosis ²	
Blackleg ²	
Cystic echinococcosis ¹	
Cowdriosis ²	
Foot and mouth disease (FMD)	
Peste des petits ruminants (PPR) ¹	
Lumpy skin disease (LSD) ²	
Streptothricosis (STRE) ²	
Trypanosomiasis (TRYPS)	
Infectious ²	
Contagious bovine pleuropneumonia (CBPP) ²	Infectious
Contagious caprine pleuropneumonia (CCPP) ¹	(respiratory)
Other infectious respiratory disease	
Other respiratory disease	
Pasteurellosis	
Endoparasites	Gastrointestinal
Nutrition	(G.I.)
Neonatal Calf Diarrhea (NCD) ²	
Other digestive problem	
Footrot ¹	Other
Injuries	
Sudden death	
Other Reproductive Problem	
Unspecific	

¹Disease only recorded in small ruminants. ²Disease only recorded in cattle.

on causes of mortality and reproductive performance losses was collected from farmers and pastoralists. The incidence of stillbirths was not raised and not specifically included; producers rather reported pregnancy terminations as abortions. The study involved 10,291 cattle (1,266 cross breeds) and 16,587 small ruminants (sheep and goats).

Living standards measurement study (LSMS)

The LSMS-Integrated Surveys on Agriculture (LSMS-ISA) are household survey projects that collaborate with the national statistics offices in partner countries, of which Ethiopia is one. The LSMS-ISA project supports the design and implementation of the Ethiopia Socioeconomic Survey and is implemented every two years (27).

For our study, we used data collected from rural households ($n = 4,000$) in 2011/2012 (wave 1) and the same households revisited and expanded to include 1,500 urban households ($n = 5,500$) in 2013/2014 (wave 2). Sampling followed a stratified random sampling scheme intended to select households

according to population size. For information on sampling design, refer to The World Bank (27). Data were collected face to face *via* a livestock questionnaire interview, administered where at least one member of the household was involved in rearing livestock. Information was collected for the previous 12 months about animal holdings and costs, and production, cost and sales of livestock byproducts. Data were collected for deaths but not abortions. Completed questionnaires were sent to the Central Statistical Agency in Addis Ababa. Questionnaire design for both waves of data collection and details for field work implementation are described previously (27, 28).

In this dataset there was a total of 11,650 non-duplicated cattle records for 2011 and 2013. Of these, 7,081 records were deemed valid animal holdings, of which 6,182 recorded zero deaths (87%). Cause of death (disease) was not recorded. There was a total of 23,250 non-duplicated small ruminant records for 2011 and 2013. Of these, 6,457 records were deemed valid animal holdings, of which 5,305 recorded zero deaths (82%). As for cattle, cause of death (disease) was not recorded.

Disease outbreak and vaccination reporting (DOVAR)

DOVAR collated data from six regional diagnostic laboratories within Ethiopia (Mekele, Asosa, Bahardar, Shola, Dire Dawa, and Soddo) as well as the federal National Animal Health Diagnostic and Investigation Centre (NAHDIC) in Sebeta, and from the Department of Disease Outbreak Investigation and Vaccination in Addis Ababa. The purpose of DOVAR is to conduct national passive surveillance through the collection of animal disease outbreak and vaccination data on a monthly basis.

The data were collected by the laboratories monthly between 2009 and 2015, and annually in 2016 and 2017. The laboratories collected samples from sick animals during outbreaks for analysis as well as conducting post-mortems. Data in our study were taken from the laboratory registers, between December 2017 and early 2018.

These data represent monthly returns of number of outbreaks, cases and deaths from notifiable diseases, at state level. If there were two outbreaks ongoing, it was possible to have two separate records for a single disease in a state in the same month, therefore data were aggregated by State, Disease and Year in order to be more in line with the other datasets analyzed, with values for all States, Diseases and Years fully calculated or inferred i.e. where no entry occurred in the data for a specific State/Disease/Year combination a zero was recorded, since no deaths were reported. For cattle, a total of 1,386 records resulted, of which 1,060 were zero deaths, 79 were >100 deaths and 5 were >1,000 deaths (maximum value 2,471). For small ruminants, as described for the cattle data, these data were aggregated by State, Disease and Year. A total of 900 records resulted, of which

677 were zero, 80 were >100 and 6 were >1,000 (maximum value 4,838).

Data analysis

In each analysis, data as analyzed consist of the number of animals that died or aborted due to a specific disease, on a specific geographical unit (farm or governmental region) during a specific time period (year). These data include zeros where no animals died from a specific disease. They are used as the outcome, or dependent, variable in all analyses, with associated covariates, including the specific disease which caused the deaths or abortions. These data can be considered independent from one another, except in the unfortunate case where the majority of a flock or herd are lost to a disease. Even in this case, restocking could lead to further, unrelated deaths or abortions.

Initially, where the total number of animals was recorded, a binomial model was proposed for these data, that is, the number of deaths or abortions relative to the numbers of animal kept. However, a large number of records indicated more deaths or abortions than animals kept. This uncertainty about numbers may represent where a number of animals were kept at the beginning of the year, some animals died, and the remaining animals are reported as the number kept. Therefore, the binomial model approach was abandoned, in favor of using the number of animals as a covariate in the model as a more robust approach. This choice of model means, however, that even where the number of animals per herd is specified, it is possible for the model to predict that a greater number of deaths will occur.

Therefore, all datasets were analyzed using a generalized linear mixed model, (GLMM), specifically a Poisson model with overdispersion fitted as a random intercept associated with herd and zero-inflation, if it improved the model fit. Use of a mixed model allowed for the variation between herds and the random effect represents the accumulation of many small herd-level effects that were not measured, such as training, cleanliness etc. Where possible, an offset term was included to account for the differing numbers of animals in each herd. Where herd size was available, predictions were standardized, and this is stated where relevant. Where herd size was not available either a Poisson or Negative Binomial model was used, whichever had the better fit.

In practice, an initial model was proposed for each dataset, with a Poisson error structure with a random effect based on herd ID, where this was available. The model was compared to a similar model allowing for an increased number of zero observations using `glmmTMB` (version 1.1.4) (29) in R (version 4.1.2) (30), and where there was support for the latter model, an initial logistic regression model was considered, using `glm`, and the optimum model for zero-non-zero observations, including explanatory variables, was identified. A conditional (non-zero

observation) model was then fitted using glmmTMB. The non-zero observations were modeled as a zero-truncated Poisson distribution with herd fitted as a random intercept, and the optimum set of explanatory variables identified. Formally this can be described as a “hurdle” model, reflecting the two stages of the fitting process.

Models containing various combinations of explanatory variables were compared using Akaike’s Information Criterion (AIC), with the model with the lowest AIC selected as the best fitting model, subject to model assumptions being satisfied. Explanatory variables in the final model are presented on a case-by-case basis as the variables available for inclusion were not consistent across species and outcomes. Where models would not converge, a one-way non-parametric test equivalent to a parametric analysis of variance, a Kruskal-Wallis test, was used. Model results were summarized by predicted values based on the best fitting model and presented as box and whisker plots, where each plot represents a summary of the number of ruminants predicted to die in each of the sampled herds, and where every point is a herd based on the model predictions. For the DOVAR data the box and whisker plots are plots of the original data. For all other analyses the plots are predicted mean values for random farms in the appropriate area, management, disease etc. using all covariates from the final fitted model.

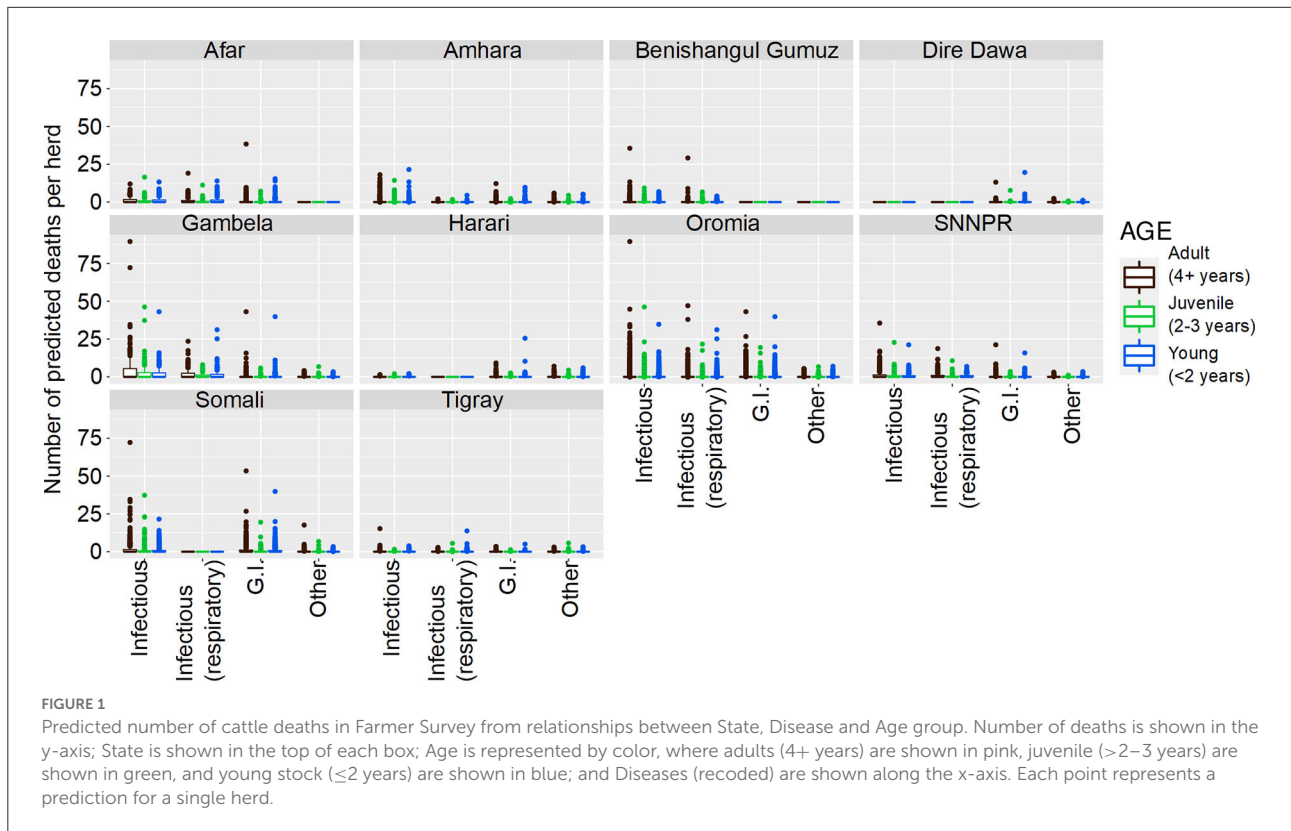
Results

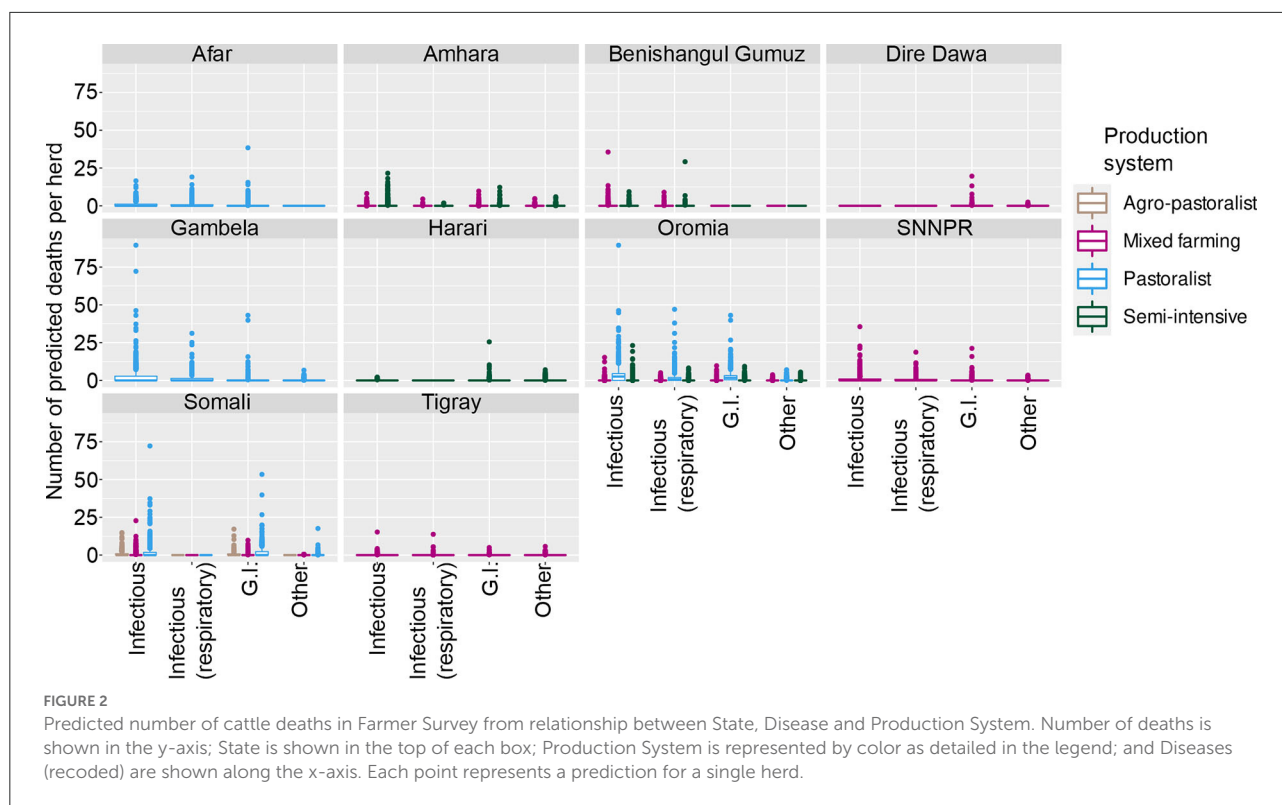
Farmer survey

Deaths in cattle

For cattle deaths (per annum), the final model was a hurdle model with the non-zero counts coming from a Poisson distribution, with a random effect for overdispersion. The final model selected was: the probability of observing zero or non-zero depended on Production System, State, Disease, Ecosystem and interactions between Production System and Disease, State and Disease, and Disease and Ecosystem. Taking this into account, the differences between non-zero counts could be accounted for by Production System, Ecosystem, Disease, Age of animal and interactions between Ecosystem and Disease, Disease and Age, and Ecosystem and Age. The complex relationships between State, Disease and Age group are shown in Figure 1, and State, Disease and Production System are shown in Figure 2.

From the total of 10,291 cattle in the analysis, 4,180 (41%) died (29% of herds (109 of 382) had no deaths reported for any of the diseases). Generally, a greater number of deaths was predicted for older age groups, the states of Oromia and Somali, infectious diseases and pastoralist systems (Figures 1, 2). Indeed, for a few pastoralist herds in Somali, Gambela and Oromia, the





model predicted up to 90 deaths in adult cattle and up to 50 deaths in younger cattle due to infectious diseases.

Abortions in cattle

The final model was a hurdle model with the non-zero counts coming from a Poisson distribution, with a random effect to account for overdispersion. The final models selected were: the probability of observing zero or non-zero events depended on State, Disease, Ecosystem, and interactions between State and Disease, Disease and Ecosystem, and State and Ecosystem; the count of non-zero entries depended on State, Production System, Disease, Ecosystem and an interaction between Disease and Ecosystem.

The results are presented as predictions. [Figure 3](#) demonstrates how abortions vary by state, ecosystem and disease.

Of all the herds, 252 (66%) had no abortions and 131 (34%) had at least one predicted due to any disease in any of the states in any ecosystem. The highest number of abortions predicted for a herd were caused by Brucellosis in Oromia and mainly in the lowland regions, followed by FMD and late abortions (> 6 months) in the same state and ecosystem.

Deaths in small ruminants

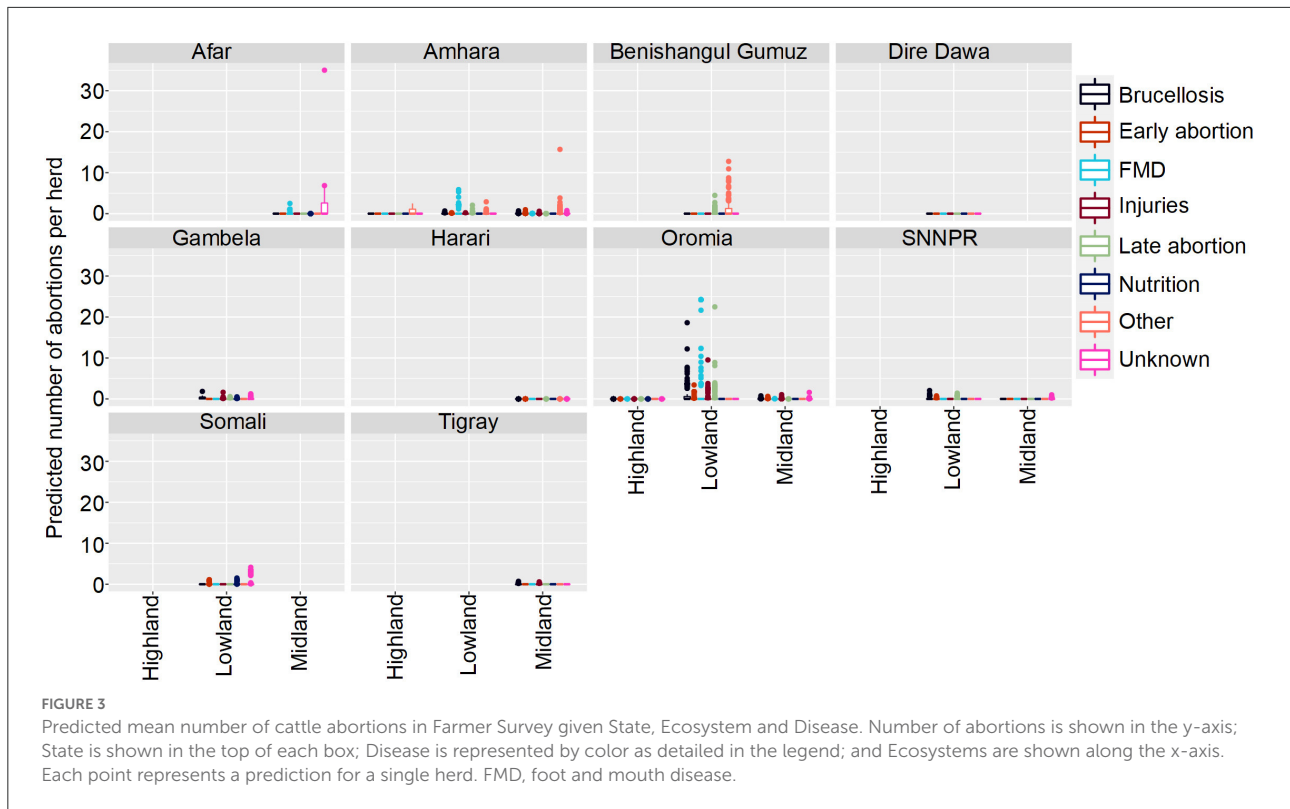
The final model was a hurdle model with the non-zero counts coming from a Poisson distribution with the total

number of animals reported used as a covariate, with a random effect to account for overdispersion. We included 16,587 small ruminants in the analysis, of which 5,611 (34%) died (57 of the 303 herds reported no deaths). Similarly to cattle, SR mortality was not uniformly distributed across states. Variables that were significantly related to mortality included Disease, State, Sex, Age, Total Number of animals, Production System and Ecosystem with several interactions between variables. As the fitted models included the effect of Total Number of animals held, this term was set to a standardized value of 10, close to the median value, to standardize predictions. [Figure 4](#) demonstrates how deaths vary by State, Ecosystem and Disease, and [Figure 5](#) demonstrates variation by State, Disease and Production System.

Most SR herds included in the survey were predicted to have no deaths due to any of the diseases. Increased mean number of deaths were associated with respiratory diseases, the states of Afar, Oromia and Somali, and pastoralist farming ([Figures 4, 5](#)). However, in Somali, increased mean number of deaths in pastoralist herds were associated with gastrointestinal diseases.

Abortions in small ruminants

The final model was a hurdle model with the non-zero counts coming from a Poisson distribution with the total number of animals reported used as a covariate, with a random effect to account for overdispersion. Of 2,128 small ruminants, 255 had an abortion. The best-fitting model for zero against non-zero observations included Disease,



Ecosystem, State, Total animals, and interactions between Disease and State, State and Total animals, Ecosystem and State, and between Disease and Total animals. The best-fitting model for number of cases, conditional on there being greater than zero cases, included Disease, State, Ecosystem, Total number of animals, and Production System, with no interactions. Results are presented as mean predictions across all herds (Figures 6, 7).

Most (74%) herds had no abortions due to any disease, in all states, ecosystems and production systems. In Amhara, herds situated in lowland and midland ecosystems were more likely to have abortions than herds in highland areas. Mean numbers of abortions due to brucellosis were predicted to be very high (>45 per herd) in lowlands and midlands, and mixed farming in Amhara. In contrast to the analysis of deaths, in Oromia there were marginally more abortions predicted in mixed farming than pastoralist systems.

Living standards measurement study (LSMS)

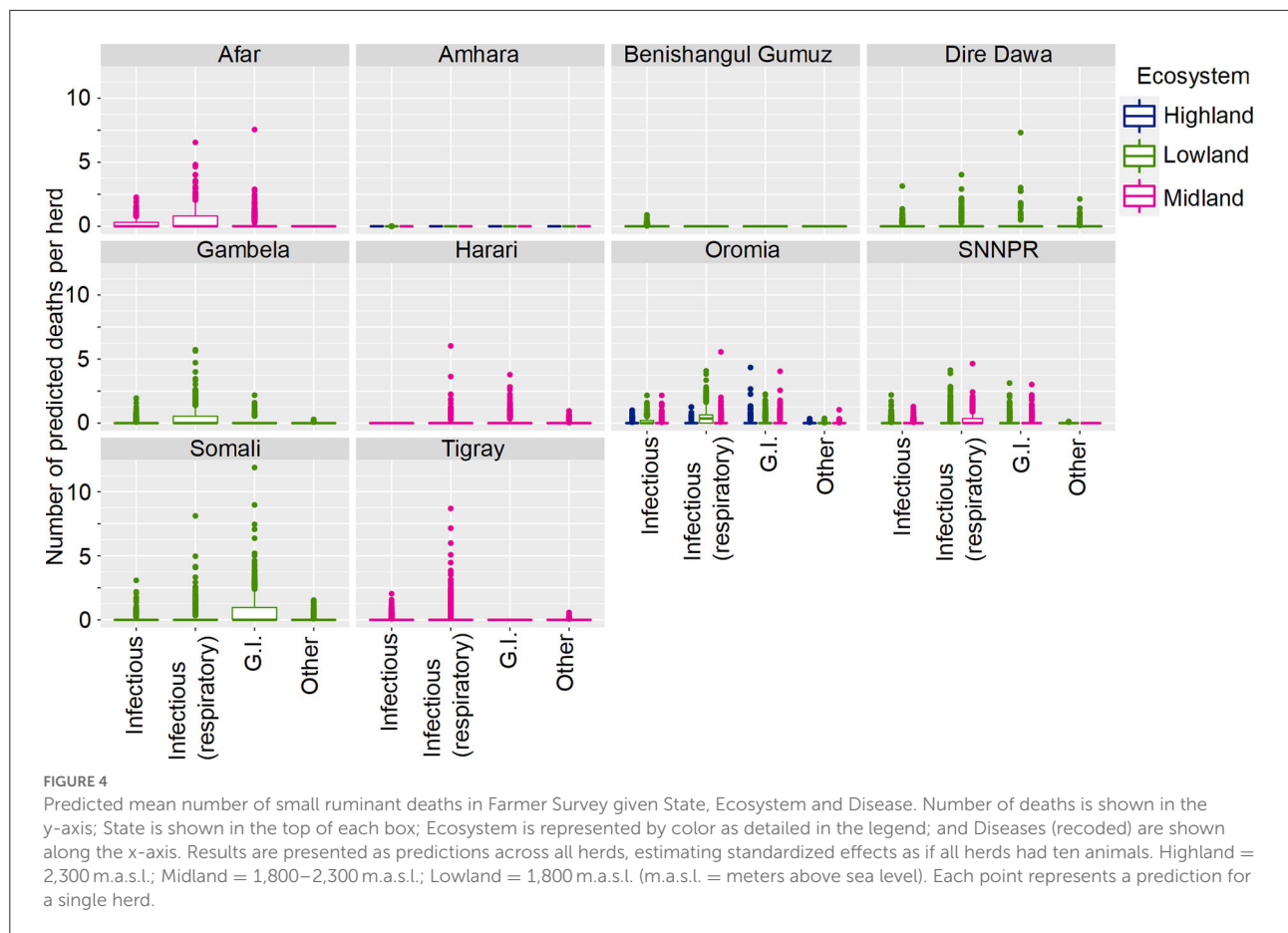
Cattle deaths

In 2011 and 2013, cattle mortality was 11.4% (1,252 of 10,989) and 6.3% (782 of 12,367).

The final model was a hurdle model with the non-zero counts coming from a Poisson distribution with the total number of animals reported used as an offset, with a random effect to account for overdispersion. The best-fitting model for zero vs. non-zero observations included State, Year, Longitude, Elevation, Agroecology, Sex, and interactions between Year and Longitude, State and Longitude, State and Elevation, Longitude and Agroecology and between State and Agroecology as terms. The best-fitting model for the non-zero counts of deaths included Latitude and Year. Results are presented as mean predictions across all herds, standardizing the effects as if all herds had ten animals (Figures 8, 9).

In the model predictions (for a herd of 10 cattle), the majority of herds had zero deaths for both years, in all states and in both sexes. Where there were deaths predicted, there are major differences between states and years, and the effect of year varies between states. Dire Dawa had the highest mean number of cattle deaths predicted in 2011, for both male and female cattle, whereas Gambela had the highest number in 2013, again for both sexes. The lowest numbers of predicted deaths, for both sexes, and in both years, were in Tigray.

The Southern Nations, Nationalities, and Peoples' Region (SNNPR) recorded data for six of the eight agroecological zones, the most of any state. Harari only recorded data for a single zone. The agroecological zone with the greatest risk of animals dying in 2011 was 'tropic-warm/semiarid' in SNNPR,



followed by ‘tropic-cool/semiarid’ in Dire Dawa, whereas in 2013, it was ‘tropic-warm/semiarid’ in SNNPR again, followed by ‘tropic-warm/subhumid’ in Gambela. Overall, there were more predicted deaths in 2011 compared to 2013, for all states.

Small ruminant deaths

In 2011 and 2013, mortality was 10.4% (650 of 6,200) and 11.1% (814 of 7,354) for sheep, and 12.8% (1,038 of 8,090) and 11.5% (1,160 of 10,133) for goats, respectively. The final model was a hurdle model with the non-zero counts coming from a Poisson distribution with the total number of animals reported used as an offset, with a random effect to account for overdispersion. The best-fitting model for zero vs. non-zero entries included State, year, Sex, Longitude, Agroecology, Elevation, and interactions between State and Elevation, State and Longitude, State and Agroecology, Longitude and Agroecology, Elevation and Agroecology, Year and Longitude, Elevation and Year, Elevation and Longitude, between State, Elevation and Agroecology and between State, Longitude and Agroecology as terms. This model therefore included six direct effects, eight two-way interactions and two three-way interactions.

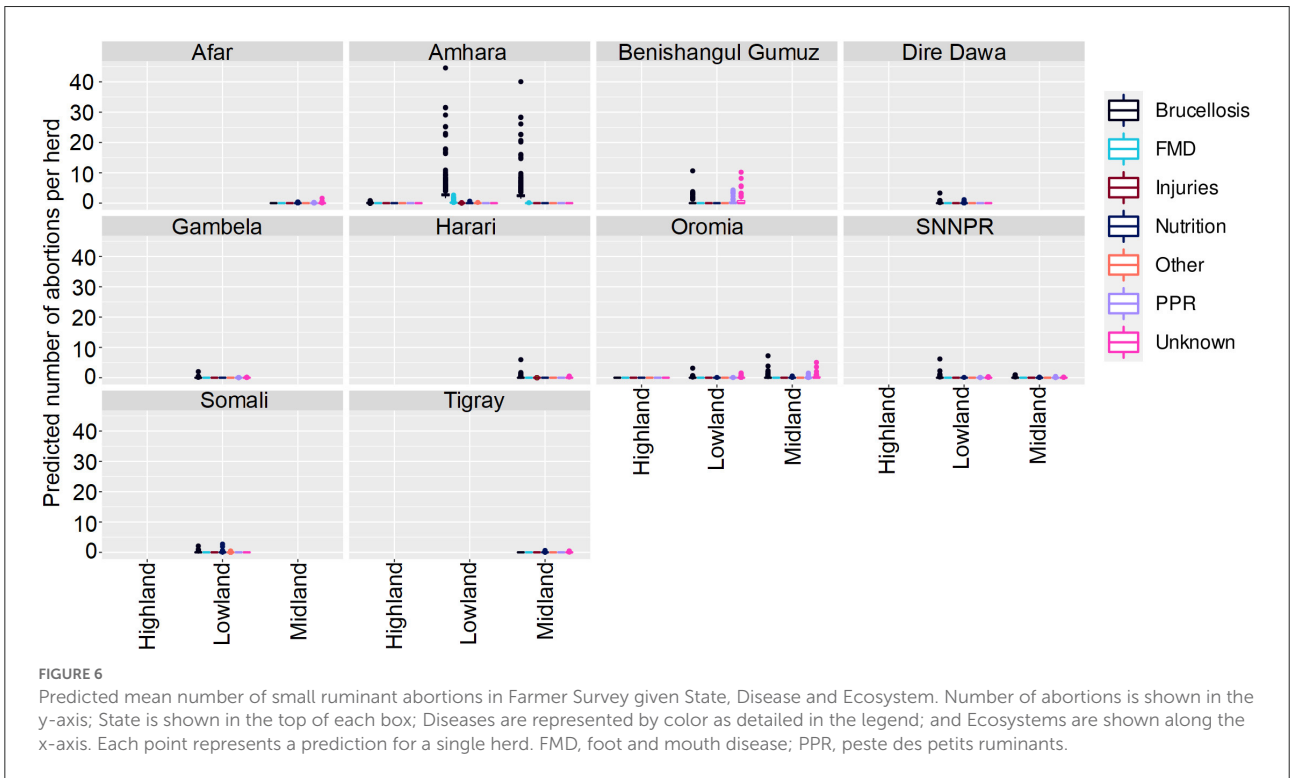
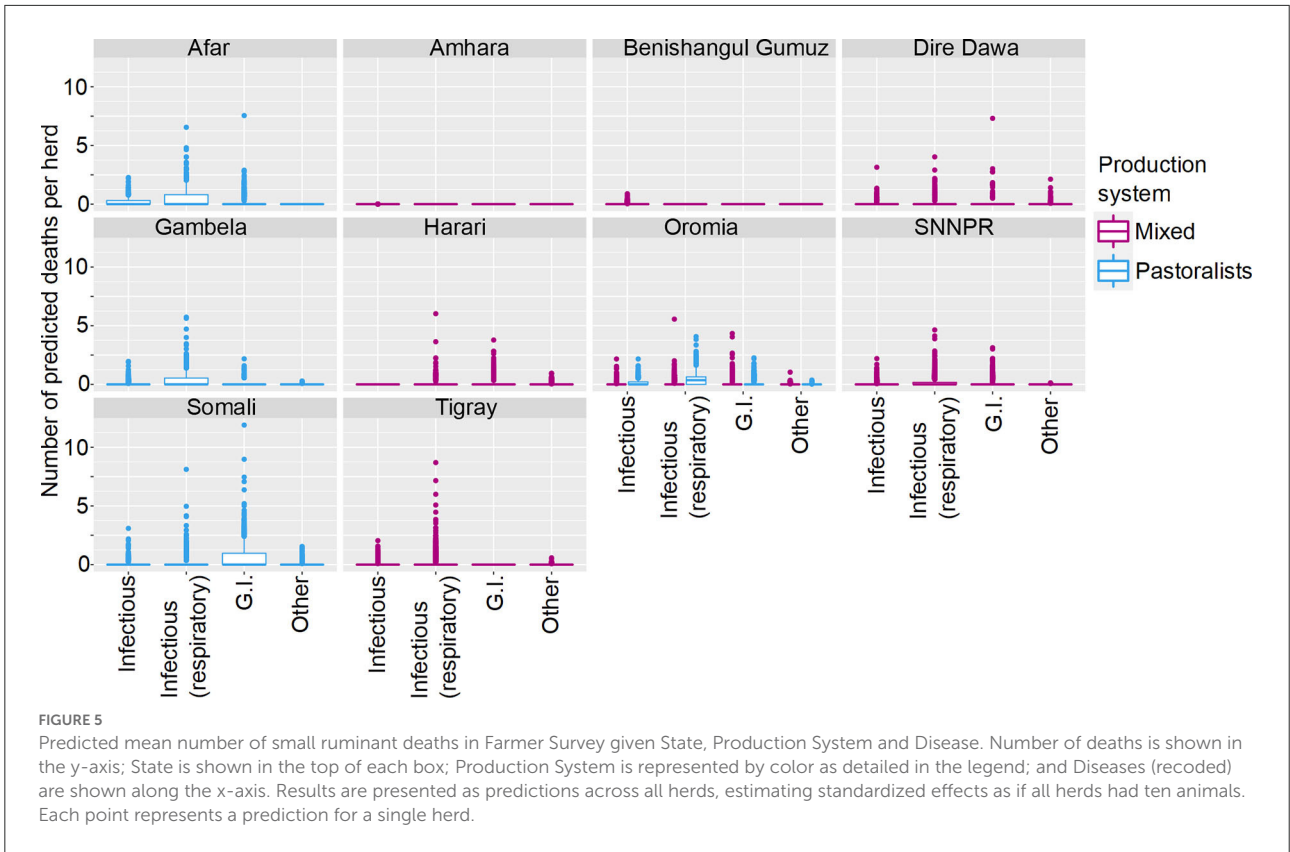
The best-fitting model for non-zero counts included Longitude, animal Sex, Agroecology, Year and an interaction between Longitude and Year as terms, with a random effect of herd, and an offset of the natural logarithm of the adjusted number of animals held ($n + 0.5$ to account for zeros). Results are presented as predictions across all herds, estimating the effects if all herds had ten animals (Figure 10).

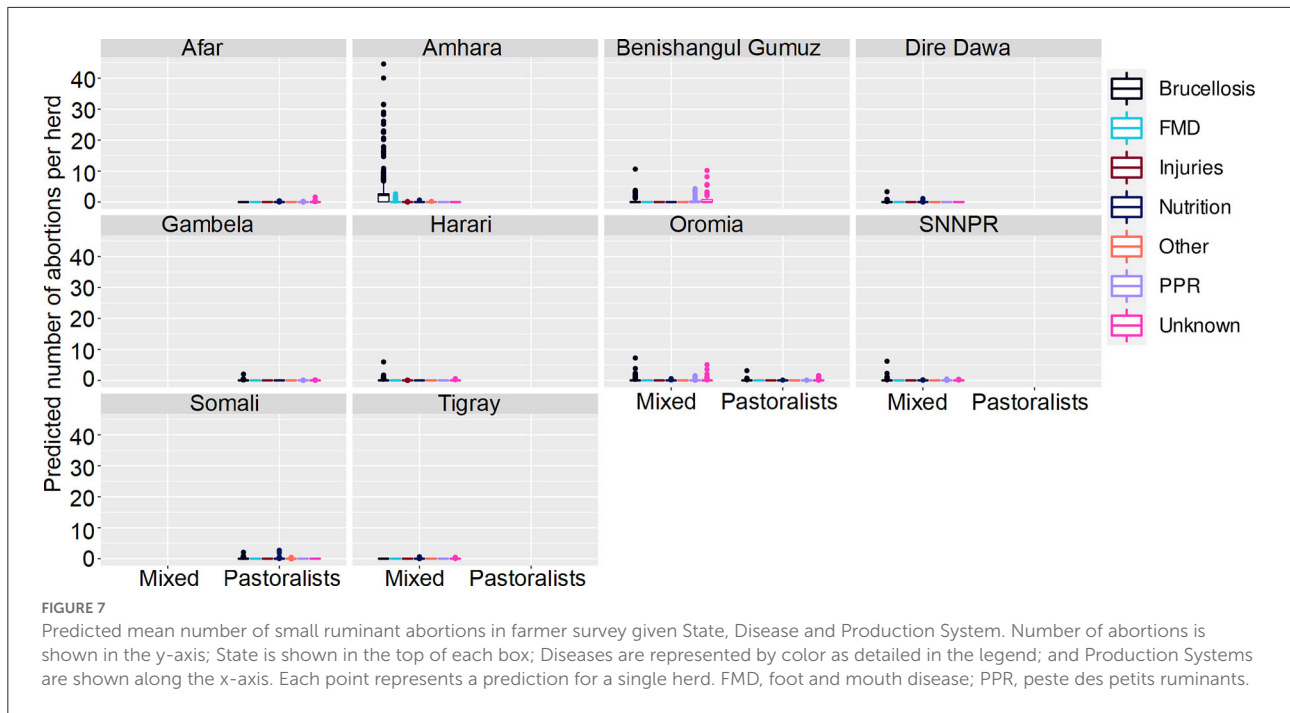
Median mortality in small ruminants for a herd of 10 animals was predicted to be under 3 for herds in all agro-ecological areas within the different states. Highest predicted mortalities indicated that complete herds perished. Increased mortality was seen in Benshagul Gumuz, Dire Dawa and Gambela, and female small ruminants were at slightly higher risk than males. There was no clear trend in the association of mortality with either humidity or temperature.

Disease outbreak and vaccination reporting (DOVAR)

Deaths in cattle

The total number of cattle deaths between 2009 and 2017 was 34,617. The final model was a simple generalized





linear model (GLM) with the counts coming from a Poisson distribution. The best-fitting model for whether any deaths were recorded or not depended on Year, State and Disease. Conditional models, that is models based using zero-truncated distributions on the non-zero data, failed to converge, so a Kruskal-Wallis test was used. Differences, statistically significant at the 5% level, were found between States ($p < 0.001$), and also between Diseases ($p < 0.001$), but not between years ($p = 0.15$). These differences are shown in Figure 11, where number of deaths is shown on the vertical axis on a log scale, as is appropriate for such skewed data.

Deaths in small ruminants

The total number of sheep and goat deaths between 2009 and 2017 was 22,059 and 23,674, respectively. The final model was a simple GLM with the counts coming from a Poisson distribution. The best fitting model for whether any deaths were recorded or not depended on Year, State and Disease. Again, conditional models failed to converge, so a Kruskal-Wallis test was used. Differences, statistically significant at the 5% level, were found between States ($p < 0.001$), and also between Diseases ($p < 0.001$), but not between Years ($p = 0.28$). These differences are shown in Figure 11, where non-zero number of deaths is shown on the vertical axis on a log scale, as is appropriate for Poisson or negative Binomial data.

We can interpret this analysis as indicating that whether or not an outbreak occurs varies depending on the particular combination of State, Year and Disease, but should an outbreak

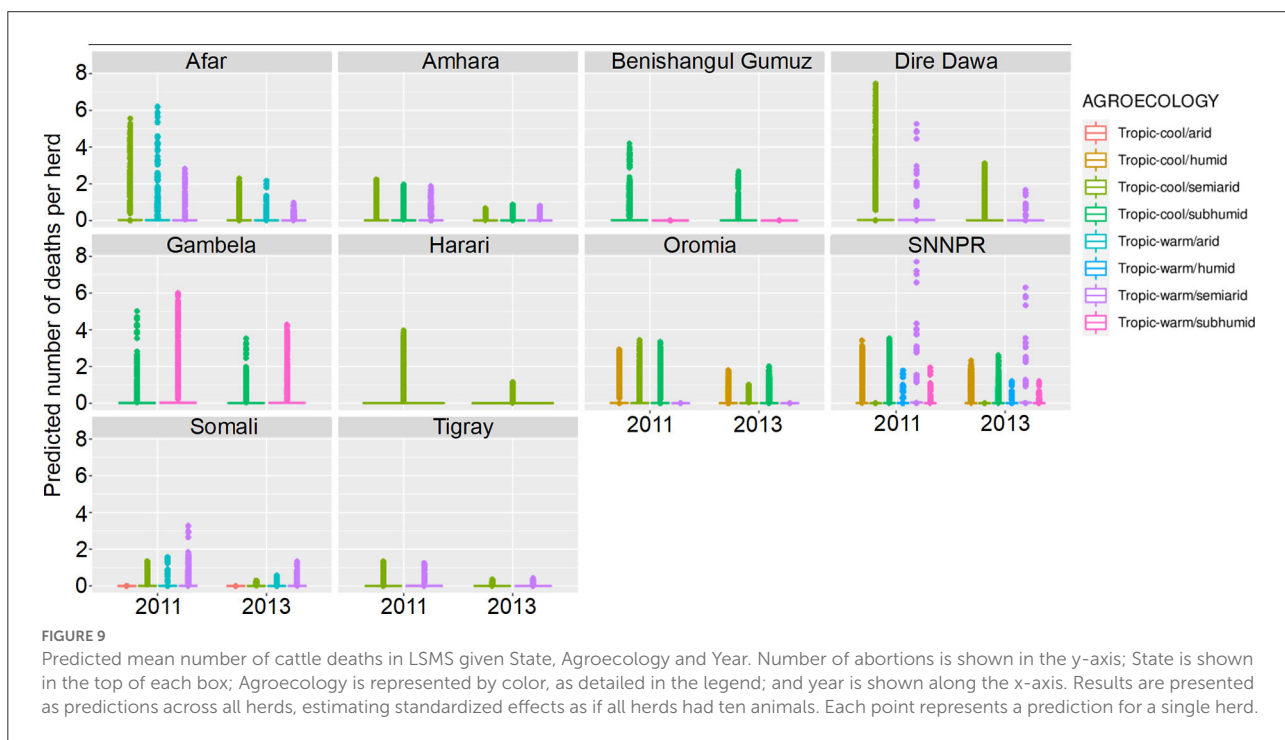
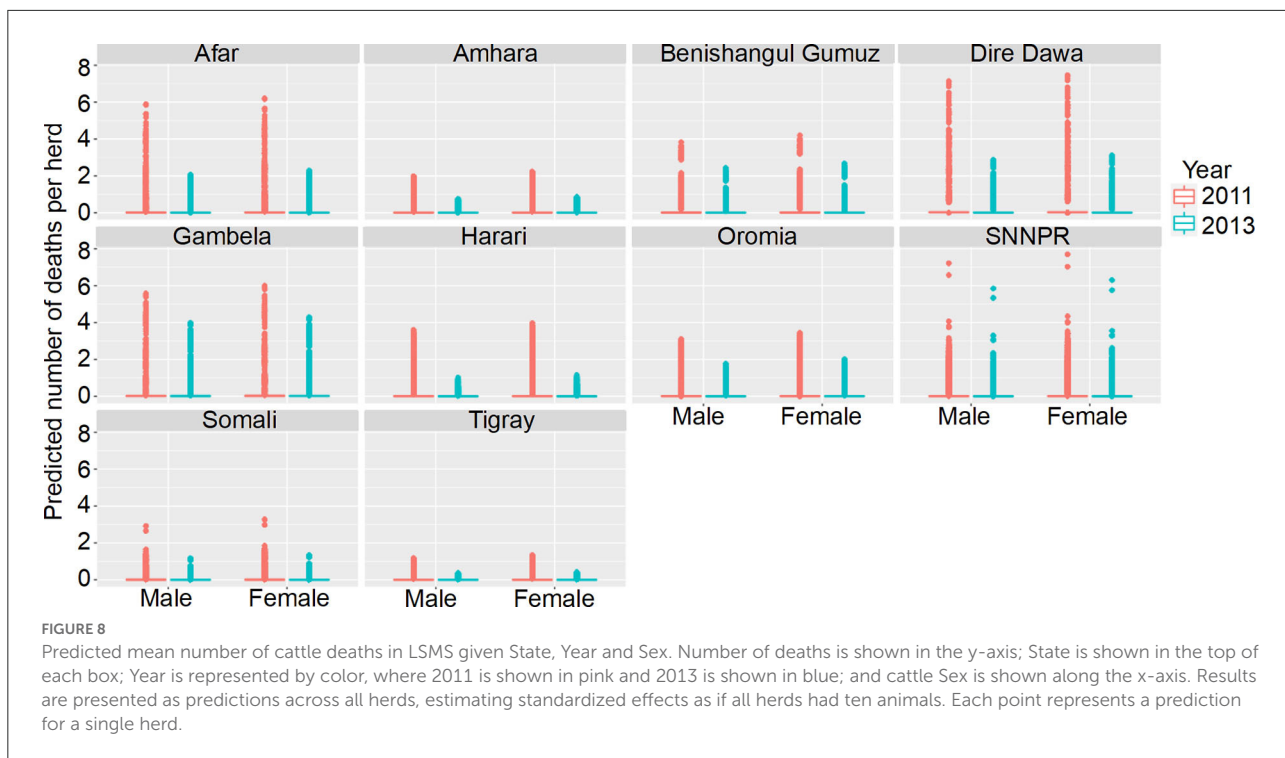
of a specific disease be recorded, the mean number of small ruminants to die will be relatively consistent (independent of state, year or of disease).

For both cattle and SR, moderate variability was predicted for the number of deaths caused by different diseases and in different states, with a median of 7 to 55 cattle deaths and 7 to 403 SR deaths per year for most of the diseases. Pasteurellosis was reported to cause the most deaths for all ruminants and very similar mortalities were reported for Anthrax and FMD in all ruminants; up to 3,000 deaths were due to FMD (Figures 11A,C). There were also similarities between states for all ruminants, with the most deaths in Amhara and Oromia (Figures 11B,D).

Discussion

This study aimed to assess if three different datasets could be combined to provide improved estimates of mortality in Ethiopia. Predictive models were created to analyse three datasets. Cattle were analyzed separately to sheep and goats, which were combined due to their similarities. Deaths and abortions were analyzed separately.

The three datasets each had their own benefits and limitations, but when combined helped to provide a more accurate picture than if assessed individually. Being veterinary-led, the Farmer Survey had a level of reliability, with farmers who were motivated to be involved, and provided targeted information, on abortions, for example. It was not nationwide, however, as it was a convenience sample. The LSMS dataset was nationwide and included apposite questions on mortality,



however it was not veterinary-led, with less motivated farmers, and so likely less reliable. Also nationwide was the DOVAR dataset, which had information gathered over a long history, but not targeted toward mortality as it was case-specific

during outbreaks. By combining the more reliable (Farmer Survey) with the nationwide (LSMS/DOVAR) and the historical (DOVAR) data, we are able to make a reasonable assessment of national mortality.



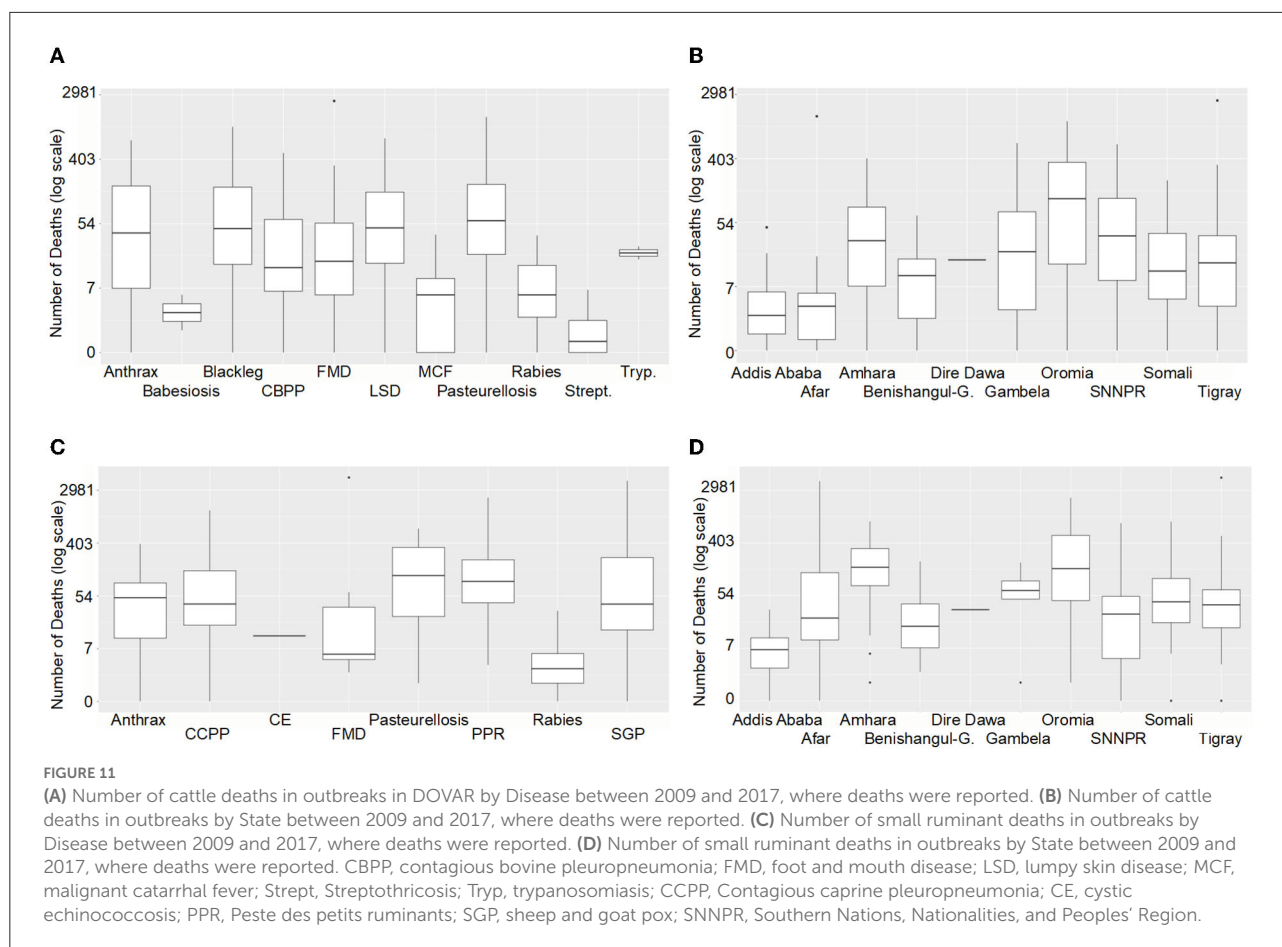
Across states and years in the Farmer Survey and LSMS data, most herds reported no deaths, with only rare occasions of large losses. In herds with increased mortality the cause of death in both cattle and SR varied by State, Years and Age groups and Production System. In cattle herds, mortality in young stock was more commonly associated with respiratory and gastrointestinal disease. In SR herds, mortality in both young stock and adults was commonly associated with respiratory and gastrointestinal diseases. In both cattle and SR herds, pastoralists reported the highest mortality. Wide variation in the predicted number of deaths in both cattle and SR was seen across agroecological zones; in 2011 in Oromia, most cattle deaths occurred in tropic cool semi-arid areas, compared to in 2013 when most occurred in tropic cool sub-humid areas. There was similar variation for SR deaths across agroecological zones but no clear trend in the association of mortality with either humidity or temperature.

Unsurprisingly, number and reasons for abortions in both cattle and SR were associated with disease, eco-system, time and place. Given relatively small herd sizes, we predicted zero abortion for most herds. Increased numbers of abortions were predicted for cattle herds in the lowlands of Oromia and Somali and predicted for SR in the lowlands of Amhara. Where the cause of the abortion was known, abortions in both cattle and SR were most likely caused by brucellosis while in cattle abortions were appreciably also caused by FMD.

The data can be considered heterogeneous, with such a high degree of variation that interpretation of them is challenging. It is necessary to question why the data are

heterogeneous; are the underlying data unreliable, or is natural/environmental variation the cause of the variability? With regards to the reliability of the data, there were indeed some areas of concern. From the Farmer Surveys, it was reported that sometimes respondents were not able to provide reliable information about the numbers of their animals and sometimes provided exaggerated mortality figures, not matching the actual herd size. The overall impression of the diagnostic laboratory data supplied was that it lacked detail and did not appear comprehensive, rather providing a reflection of the general causes of mortality and reproduction losses. A recent study reported that a lack of diagnostic laboratory resources, combined with the presence of most potential infectious and non-infectious causes of abortion made diagnosis particularly challenging in Ethiopia (31).

In both the Farmer Survey and DOVAR data, the greater mean number of deaths predicted for cattle and SR herds in Oromia are likely caused by the greater proportion of pastoralist herds with higher number of animals in that area. Other likely sources of natural variation in the datasets include different conditions for different years, such as droughts and epidemics (32, 33). That there was evidence of spatiotemporal variation of risk factors and causes of mortality is not surprising, as seasonal and environmental variation affects grazing availability and ensuing movement and mixing of animals (31, 34, 35) which can then affect disease transmission. A recent systematic review observed a scarcity of disease-associated mortality data for Ethiopia, with only 14 studies identified [(36), under review].



Droughts occur frequently in Ethiopia, causing severe damage and harm. Previously, droughts were reported to occur every 8–10 years for the whole country but recently it has been observed that the frequency of drought has shortened, with severe drought happening 3 years in succession, which was the situation in 2011 and 2015–2017 (37). Conversely, the relative lack of variability in the DOVAR dataset was surprising, and as this data collection only captured disease-related events, it is difficult to interpret.

While infectious causes are regarded as predominant in ruminant abortions (38), environmental factors are known risk factors (19), as are management systems and husbandry factors (39). As husbandry systems vary across agroecological zones (40) disease prevalence varies widely across agroecological zone, as shown by Welay et al. (41). Therefore, seasonality, agroecology and production systems all significantly affect abortions (42). A recent review of the causes for small ruminant abortion in Ethiopia reported several infectious diseases and non-infectious conditions, and highlighted evidence gaps on disease dynamics (20). It also identified the major risk factors for small ruminant abortion, including age, parity, genetics, geographical and environmental factors, and herd size in the varying production systems. In order to establish the source of the variability, it

would be necessary to compare the data with unbiased data, something that was not available. However, as multiple data sources were assessed and similar heterogeneity was observed in those analyzed, it is likely that the observed heterogeneity is real, but possibly confounded with, and compounded by data collection artifacts.

It was notable that the addition of some explanatory factors caused models to fail to converge, however, once other variables had been added it was sometimes possible to include them. The reported models should be viewed as being the optimum models to describe the data, given the datasets provided, and it cannot be guaranteed that replications of the datasets would result in the same models being selected.

Using the models to examine predictive outputs i.e., how many deaths or abortions we would expect from a similar set of herds across the various explanatory factors shows that in nearly all cases zero deaths or abortions would be observed. There are, however, occasional herds where relatively large numbers of deaths or abortions might occur; in the Farmer Survey, for example, up to 90 deaths were predicted in a single herd. The data did not provide the evidence required to explore whether this variability is consistent across time or not.

As stated, the Farmer Survey and LSMS analyses indicated that losses, from both deaths and abortions, in domesticated ruminants in Ethiopia demonstrate a large amount of variability, both geographically and across time. They represent losses from a number of diseases, and the major diseases recorded vary across both time and space. It is likely that the country-based mean death rate varies widely across different areas, management systems and from year to year. The fitted models assist our understanding of patterns of mortality, however, the degree of variation observed, particularly across time, suggests that these models are only really useful as a *descriptive* tool, and are unlikely to be useful *predictively*. The model predictions provide some insight into why mortality rates are highly variable and therefore unlikely to be a useful measure of production or change in health status. As most herds had no deaths or abortions, with a few herds having relatively high mortality, a single metric is very hard to interpret. This was reflected in our modeling strategies, and in particular in the choice of hurdle models and models allowing for zero-inflation and excess heterogeneity.

Summaries would be best based on data that are more likely to be stable, i.e., less likely to change due to random influences. Furthermore, for farmers to keep reliable records of outcomes, they should be encouraged to record outcomes that are of interest to the farmer. While deaths and abortions ought to be of interest to farmers, their rarity and high variability likely make them of less use to a researcher trying to improve productivity, and indeed they are only indirectly related to productivity as they represent a potential loss. Tsouloufi et al. recently established a systematic mapping protocol to review the available evidence on disease prevalence and associated mortality in ruminants in Ethiopia, in attempt to address the disparate data and to improve decision-making on disease control policies (43). A recent systematic review by Wong et al. (3) observed that the greatest proportion of mortalities in cattle, sheep and goats was reported to be in those under the age of 6 months. Interestingly, this study did not observe increased mortality in young stock compared to adults, as young stock mortality is generally considered high in sub-Saharan Africa, which leads us to question farmer recall and whether they remember adult losses more than young stock. Wong et al. (3) also highlighted that there was a high degree of variability in both causes and risk factors which meant that 'mortality rate' was an incomplete indicator, and as such, qualitative and quantitative data on mortality causes would improve the understanding and allow preventative actions. Information is required on the causes of mortality for decision-making, as well as the rates in different livestock species (44).

We would suggest, therefore, the investigation of alternative measures of productivity. For example, number of offspring raised to 1 year old per dam, or median or mean age of adults are likely to be more indicative and useful measures. The first

relates to mortality of young stock; the second to adult mortality. Both variates would be expected to be more stable over time than numbers of deaths or abortions, since all observational units (animals) contribute to the estimate, and not merely those with large losses. In addition, they are likely of more direct interest to farmers since they represent profit or productivity, and can be more easily equated to financial gain/loss.

The study had several limitations. The objectives were somewhat misguided in that they were initially to obtain numbers in order to inform interventions. None of the datasets provided could be reasonably expected to provide unbiased, accurate estimates of the death or abortion rate. The Farmer Survey was a convenience survey, using local experts to help define the sampling frame. Thus, the local experts might have chosen individuals known to have had problems, or not. Survey households were selected based on their accessibility and availability to the study teams (selection bias), which may affect the veterinary service delivery to these households and impact on mortality and abortions, however it was not logistically feasible to include less accessible households. Organizational constraints prevented the inclusion of Addis Ababa city administration in the Survey. Additionally, multifactorial etiologies were not considered as the survey depended on farmer recall. The LSMS data comes from an unknown sampling frame. It is believed that this sampling frame is carefully chosen, but it is meant to represent a country's economy, and is not necessarily representative of its agricultural sector or livestock population. Many LSMS records were considered invalid due to data discrepancies such as farmers reporting they owned livestock, but when asked how many animals they owned, zero were reported. Additionally, these data are collected retrospectively; all individuals are asked for their losses during the previous year, and hence are subject to recall bias. The DOVAR dataset requires a chain of events to occur before the data become available; the animal must first contract a notifiable disease; someone must recognize the disease, then decide to report it; the report must be passed through a reporting chain until it is collated into the country-wide statistic. Importantly, as a passive surveillance system of notifiable diseases it does not capture mortality due to non-notifiable diseases or due to, for example, malnutrition and, therefore, underestimates total mortality to an unknown and variable extent. However, information on mortality from DOVAR is extremely useful in explaining the year-to-year variation in total mortality as observed in the other datasets.

Conclusion

The study contributes to the understanding of the patterns of mortality in domesticated ruminants in Ethiopia. The

degree of variation observed, especially temporally, suggests that the predictive nature of the models is limited and that the estimated mortality rates are highly variable and therefore lack reliability as a metric of productivity. Unrecorded variables such as changeable seasons and environmental conditions likely affect the morbidity and mortality rates across space and time and thus the high variability observed is challenging to interpret. Nevertheless, whilst there are many limitations in the use of heterogeneous data, there are also many opportunities; indeed, heterogeneous data may be the best available, and can provide valuable insights otherwise untold of the situation in the field, especially in rural settings in LMICs, with limited resources. As such, this study should inform the choice of dataset for future metrics, such as number of offspring raised to 1 year old per dam, and the choice of statistical method used to calculate metrics that may better inform productivity and allow for suitable preventative actions and interventions to be implemented as required.

Data availability statement

The datasets for this study can be found in the GitLab <https://gitlab.bioss.ac.uk/giles/sebi-data>.

Ethics statement

The studies involving human participants were reviewed and approved by Human Ethical Review Committee (HERC), Royal (Dick) School of Veterinary Studies, University of Edinburgh. Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

Author contributions

CV and AP designed the study. SH and VN carried out the field work. GL, DE, IM, FA, and CS undertook analysis and interpretation. All authors wrote the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fvets.2022.986739/full#supplementary-material>

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