Version of Record: https://www.sciencedirect.com/science/article/pii/S0167587722000642 Manuscript 8f8021593fb0d2761495756c320704bf

1 Title: How far away do you keep your equines? Estimation of the equine population's spatial distribution in

2 France

3

7

10

13

14

15

16

19

20

21

22

23

24

26

4 Author names and affiliations: Halifa FARCHATI a.c.d, Benoit DURAND b, Maud MARSOT b, David GARON c,

5 Jackie TAPPREST ^a, Carole SALA ^d

6ª Laboratory for Animal Health in Normandy, Physiopathology and Epidemiology of Equine Diseases Unit,

French Agency for Food, Environmental and Occupational Health & Safety (ANSES) F14430 Goustranville,

8 France

9^b Epidemiology Unit, Laboratory for Animal Health, French Agency for Food, Environmental and Occupational

Health and Safety (ANSES), 94700, Maisons-Alfort, France

11° Normandy University, UNICAEN, Centre F. Baclesse, UR ABTE EA 4651, 14000 Caen-France

12^d University of Lyon- Epidemiology and Support to Surveillance Unit, French Agency for Food, Environmental

and Occupational Health & Safety (ANSES), 31, avenue Tony Garnier, F69364 Lyon Cedex 07, France

Keywords: Spatial distribution, Equine, Owner, Bayesian model, Surveillance

17 Acknowledgments

18 We would like to thank the French horse and riding institute (IFCE) and the French Agency for Food,

Environmental and Occupational Health & Safety (ANSES) for their financial support. We are also very

grateful to the data providers: IFCE, Le Trot, France Galop, the French ministry for agriculture and food. We

would also like to thank the owners and keepers who answered our surveys. We need to thank various

people having contributed in different ways to the completion of this work: Jean-Philippe Amat, Marie

Delerue, Mathilde Dhollande, Romain Dhollande, Xavier Dornier, Joelle Dop, Arnaud Duluard, Bénédicte

Ferry, Giovanni Lo Lacono, Carole Margelidon, Aurélie Merlin, Marion Wujek Moreau, Julie Schneider,

25 Caroline Teyssier, and Sonia Wittreck.

1	Title: How far away do you keep your equines? Estimation of the equine population's spatial distribution in
2	France
3	
4	
5	Keywords: Spatial distribution, Equine, Owner, Bayesian model, Surveillance
6	

Abstract

7

8

9

10

11

12

13

14

15

16

17

18

19

20

21

22

23

24

25

26

27

28

29

30

It is essential to have an accurate picture of the spatial distribution of equines to be able to monitor equine health events effectively. In France, this information is only available for certain categories of live equines kept in professional structures and for dead equines removed by renderers. This limits the surveillance, prevention and control methods able to be used to prevent the spread of equine diseases. Our study aimed to provide a realistic estimate of the spatial distribution of the French equine population at the detailed scale of the French commune (France's smallest administrative unit). For this purpose, we adapted the Bayesian method used by Lo Lacono et al., based on the distance between the owner's location and the location of his/her equines, and on the percentage of urban coverage. To assess whether the location of dead equines could be representative of the location of live equines, the distribution of distances between equines and owners was calculated from a sample of live equines on the one hand, and a sample of dead equines on the other, both accurately located. We also tested two different assignment methods for locating equines: Method 1 assigned to each owner a single holding commune, while Method 2 allowed more variability in holding communes for owners associated with multiple equines. A marked difference was observed between Methods 1 and 2 regardless of the sample used, with only 2.4% and 4.3% respectively of the communes having the same number of equines. Conversely, little difference was observed in the results whether the live or dead equine sample was used, with approximately 45% of the communes having the same number of equines. Regarding differences in spatial distribution, Method 1 based on the live equine sample estimated higher local densities of equines without considering urban areas. In contrast, Method 2 provided more dispersed maps, with low densities in the densest urban areas. In conclusion, dead equines appeared to be representative of live equines and some of our estimates are consistent with the information collected by the French horse and riding institute (IFCE). These results now have to be compared with field data to test their relevance.

1 Introduction

31

32

33

34

35

36

37

38

39

40

41

42

43

44

45

46

47

48

49

50

51

52

53

54

55

56

Data on geographic location and interspecies contacts are often considered key factors in developing accurate predictive models of pathogen spread or establishing effective and appropriate methods for the surveillance, prevention or control of animal diseases (House and Keeling, 2008a, b; M. J Keeling et al., 2010). The major influence of the population's spatial structure on the spread of infectious diseases is well known in epidemiology, whether for humans or domestic/wild animals (Ferguson et al., 2005; Ferguson et al., 2001; Gudelj and K.A.J., 2004; Keeling et al., 2001; Swinton et al., 1998). Depending on the different ways that diseases can spread and the density of animal populations, a disease can spread locally then extend either slowly or quickly over very long distances, especially when sick animals are moved around. The effectiveness and success of measures to control the speed and spread of disease depend on the existence of high quality spatial and epidemic data, in addition to the traceability of animal movements. The example of the equine influenza outbreak in Australia in 2007, when nearly 4,500 equine premises were infected in less than two months (Brendan et al., 2009; Callinan, 2007) demonstrated how the lack of accurate spatial data on the equine population and its movements can greatly complicate the implementation of surveillance methods and control measures to combat and control diseases. It also showed how traceability information is helpful in rapidly implementing effective control measures (Garner et al., 2010). More recently, in the South of France, management of the nine outbreaks of equine infectious anemia virus (EIAV) occurring between 2017 and 2019 systematically required use of a specific epidemiological survey to identify neighboring structures and test equines that had potentially been in contact with equines already identified as being sick (Amat et al., 2018; Merlin et al., 2021). In the absence of information on precise location of French equines, these investigations are time consuming and delay both the identification of sick animals and implementation of control measures. This example shows that the traceability of equines in France needs to be improved in order to effectively take measures against diseases. Currently, only the movements of racehorses in training have to be traced for anti-doping tests. For all other equines, current French regulations only require registration of equine

premises in the SIRE database, which is France's national identification system. They do not, however require

the registration of the equines kept there. The spatial distribution of French equines is currently estimated from surveys of professional structures, and identification checks carried out by the French horse and riding institute (IFCE) (Ecus-Ifce). However, the representation obtained is partial, as the majority of equines are kept in non-professional structures, often on the horse owners' premises.

All around the world, surveillance plan strategies and outbreak management efforts in the equine sector are hampered by the lack of information on the precise location of equines and their movements. The recent enforcement in April 2021 of the European Union animal health law (EU's AHL) on transmissible animal diseases (Regulation (EU) 2016/429 (Anonyme, 2016)) should improve the traceability of equines, imposing mandatory notification of equines' usual location. Nevertheless, the operational implementation of this measure should take many years, even in countries with near-complete equine identification and a centralized identification database. The regulatory registration of the location of all equines is not expected to be effective for some time.

In the meantime, it would be very useful to managers, epidemiologists, and researchers to have realistic estimates of equine densities at detailed spatial scales in order to improve checks, surveillance, and equine disease modeling. With this in mind, Giovanni Lo Lacono et al. (Lo lacono et al., 2013) implemented in their research a Bayesian method to estimate the spatial distribution of horses in Great Britain, based on the distribution of owner-equine distance. They used a sample of owner-equine pairs that were both precisely located to estimate the probable distance between an owner and his/her equine. Additionally, an environmental component (proportion of urban coverage) modulated the location probability so as to assign to the equine the most plausible location(s). The final result was a theoretical map of the British equine population.

Adapting the approach of Giovanni Lo Lacono (Lo Lacono et al., 2013), our objective was to estimate the spatial distribution of the equine population in metropolitan France at the *commune* scale (a *commune* being the smallest French administrative unit). In our case, for comparison purposes and to assess whether the location of dead equines can be representative of that of live equines, two samples of live equines and dead equines were used like training datasets. To assign a location to live equines registered in the central

database from the SIRE database, two different assignment methods were realized. The first one assigned a single holding *commune* for all equines of each owner and the second method allowed one, two or three holding *communes* of owners, depending on the number of linked equines (one *commune* for one equine, two *communes* for two equines, or three *communes* for three or more equines). Different data sources and hypotheses that will be detailed later were used for this purpose, to obtain multiple results that are thereafter discussed. We will discuss in particular the value of combining multiple data sources and the contribution of using two samples and two methods.

2 Materials and Methods

2.1 Data sources and information

Several data sources were used to train the model and estimate the spatial distribution of the French equinepopulation.

2.1.1 Training datasets

The model was calibrated using information from three datasets:

- A survey sample dataset of 2,353 precisely located live equine-owner pairs. This sample came from a 2019 national survey of owners and keepers registered in the SIRE database (Farchati et al., 2020). The main characteristics of this sample are available in Table 1. The sample was obtained by linking each owner with one or more *communes* of equine ownership, with as many pairs as different *communes* of ownership for each owner. Racehorses in training were excluded from this dataset because they were considered differently in this study. This dataset is referred to as the live equine sample in the following.
 - A sample of 13,595 dead equine-owner pairs extracted in April 2019 from the fallen stock data interchange (FSDI) database, which has been centralizing information on all removed (dead) animals—animal identification, *commune* of death, and information on the person who requested the removal (name, surname or company name, *commune* of residence)—since 2011. The pairs included were those for which i) the equine identification number was traceable in the SIRE database (n=97,327, i.e. 33% of 298,733 initial records in the extraction of FSDI), ii) there was a high probability that the applicant was the owner of the dead equine (n=44,771 among the 97,327) and iii) the applicant had filled out the information on the cadaver online (n=13,595 among the 44,771) (through ATM¹ services managed by the IFCE). The selection was made by cross-referencing the FSDI and SIRE databases, and comparing the identification and location information for owners, keepers and removal applicants. The intention in limiting the sample to owners who used the online system

¹ ATM-équidés ANGEE (ATM) is an association created in 2009, following privatization of the public rendering service, to organize cadaver removal. It offers owners negotiated prices for cadaver removal in order to limit illegal burial and provide rendering companies with a guaranteed payment for removals.

was to increase the likelihood that the owners' home *communes* as registered in the SIRE database were up to date (as the recording of information at ATM allows it to update SIRE). This sample of dead equine-owner pairs is referred to as the dead equine sample in the following and is described in Table 1.

Data extracted from the CORINE Land Cover (CLC) (Cover) geographic database, to calculate the proportion of urban coverage for each *commune* in metropolitan France. CLC is a biophysical inventory of European land use and its changes according to a nomenclature of 44 items. To take into account the urban/rural land use aspect for equines, we calculated the proportion of continuous and discontinuous urban coverage relative to the total surface area of each *commune*.

2.1.2 Mapping datasets

115

116

117

118

119

120

121

122

123

124

125

126

127

128

129

130

131

132

133

134

135

136

137

138

139

140

In order to obtain realistic equine density maps, we assigned to each equine presumed to be living in metropolitan France a "holding" commune (where it was kept) out of the most probable ones, based on the model results. An initial identification of the equine presumed to be living in France was thus necessary. The SIRE database, managed since 1976 by the IFCE, centralizes declarative information on almost all of the equines living in France (equine identifying number, sex, breed, date of birth, date of death declaration (if relevant)), along with their owners (owner identifying number, commune for the owner's residence, date on which the equine owner became its owner, and date on which information was last updated) and in parallel information of keepers and equine premises (keeper identifying number, commune for the keeper's residence, commune of the equine premises and the date of the premises were opened and closed (if applicable)) without specific traceable link between keepers and the equines they keep, and therefore a lack of centralized equine location information in France. This database has also difficulty in identifying with certainty whether an equine is still alive or dead (it has been estimated that nearly 30% to 40% of owners have never reported the death of their dead equine(s) to the IFCE, or reported it long after the death (IFCE, personal communication)). The living equine population is consequently not well known. Four data sources were cross-checked to constitute the set of equine-owner pairs for which the equine and its owner were presumed to be living in Metropolitan France:

- Data were extracted from the SIRE database in late March 2019 as a basis for selecting appropriate data. This dataset included 3,031,446 equines owned by 790,051 owners located in 35,359 communes.
 - A dataset for 2015-2019 extracted from Europe's Trade Control and Expert System (TRACES) was used to identify equines that were no longer living in France because they had been exported.

 TRACES is a computerized system for international trade in live animals, animal products and certain non-animal products. It records, among other things, the SIRE identifier, transponder or international number and the date of departure of exported equines. The dataset extracted concerned 31,336 equines.
 - The FSDI database was used to identify the dead equines collected by rendering companies that had not been declared dead in the SIRE database. The whole database since its creation was used i.e., 298,733 removals.
 - A dataset of racehorses with the exact location of the horse corresponding to the commune in which it was being trained. This set included 37,776 trotters recorded as being in training at least once between August 2018 and August 2019, and 13,290 gallopers recorded as being in training at least once between July 2020 and October 2020. The dataset was compiled from monthly extractions of the databases for the two parent companies of horse racing in France: Le Trot (for trotters) and France Galop (for gallopers). For each horse in pre-training or training, each company records the SIRE identifier, the *commune* of the horse's location (premises where it is kept or where it is trained), and date of its entry/exit. Because these equines were accurately located, and the location of the owner(s) does not necessarily correspond to the location of the equine, due to the need for a specific (and rare) training structure, we did not apply the location algorithm to this population. However, we used the racehorse dataset to remove horses in training from the SIRE dataset and live equine sample.

From these four datasets, we obtained:

- A dataset of 1,068,142 equine-owner pairs corresponding to 1,068,142 equines presumed to be alive and belonging to 397,039 owners living in 32,837 *communes* in metropolitan France and resulting from the SIRE extraction after removing dead animals and cross-checking with the other data sources. The details of the different steps are shown in Figure 1. The dataset obtained is referred to as the equine dataset in the following.
- A dataset of 44,768 racehorses in training (12,785 gallopers and 31,983 trotters) along with their location, resulting from the removal of dead equines then the selection of a location for each equine in metropolitan France on the date closest to April 1st, 2019 (date on which the data were extracted from the SIRE database). This dataset is referred to as the racehorse dataset in the following.

2.2 Modeling

Using the approach of Giovanni Lo Lacono (Lo Lacono et al., 2013), we estimated the probability $P(r/r_0)$ that an equine is kept in *commune* of centroid r = (x, y) knowing that its owner resides in *commune* of centroid $r_0 = (x_0, y_0)$. $P(r/r_0)$ was expressed by the following equation:

179 equines in the owner's commune equines in a different commune from the owner
$$P(r/r_0) = \delta(|r-r_0|)P_0(r_0) + \frac{1-\delta(|r-r_0|)[1-P_0(r_0)]P_{sep}(r,r_0)}{[1-\delta(|r-r_0|)][1-P_0(r_0)]P_{sep}(r,r_0)}$$
(1)

with $P_0(r_0)$ the probability that the equine and owner are in the same *commune*, $|r - r_0| = \sqrt{(x - x_0)^2 + (y - y_0)^2}$ the Euclidean distance between r and r_0 , $P_{\text{sep}}(r, r_0)$ the probability that the equine and the owner are localized in *communes* remote from $|r - r_0| \neq 0$, and δ a Dirac function equal to 1 if the owner and the equine are in the same *commune*, and equal to 0 otherwise.

The probability $P_0(r_0)$ was related only to the urban coverage of the owners' commune of residence and obtained using the following equation:

186
$$P_{0}(r_{0}) = \begin{cases} n_{1} * \frac{1}{u(r_{0}) * \sigma \sqrt{2\pi}} * e^{\left(-\frac{\left[\ln\left(u(r_{0})\right) - \mu\right]}{2\sigma^{2}}\right)} & \text{if } u(r_{0}) \neq 0 \\ n_{2} * \Im & \text{if } u(r_{0}) = 0 \end{cases}$$
 (2)

with $u(r_0)$ the proportion of urban coverage of the owner's location, μ (the expectation) and σ (the standard deviation) the parameters of a lognormal distribution to be determined (Figure S-1 of supplementary

material), n_2 and n_2 the normalization constants that ensure that $\int_1^N P_0(r_0) = \alpha$ (N being the number of potential *communes* less one in metropolitan France and α the proportion of equines kept in the same location as the owner) and \Im the proportion of owners keeping their equines in their place of residence and with zero urban coverage.

Furthermore, both the live and dead equine samples showed that the distribution of distances between the owners' location and their equine was well described by a power law (Figure 2). The probability that the owner and the equine were distant from $|\mathbf{r} - \mathbf{r}_0|$ was modeled by:

196
$$P_{sep}(r,r_0) = A * ||r - r_0||^{-\beta} * P_0(r)$$
 (3)

with $|r - r_0|^{-\beta}$ the probability that the owner and the equine are distant by $|r - r_0|$, A a constant set by the constraint $\int_{D-r_0} P_{\text{sep}}(r, r_0) = 1$ (with $D - r_0$ the set of all potential *communes* in France except the one of the owner's choice) and $P_0(r)$ defined in the equation (2).

2.3 Estimation of the parameters

Parameters σ and μ of the lognormal distribution of the distances in equation (2) were estimated using the least squares adjustment method. Parameter β of equation (3) was estimated using a Bayesian approach, using Markov chain Monte Carlo (MCMC) methods to generate a distribution for the parameter. The prior considered was a non-informative uniform distribution and the likelihood was given by:

205
$$P(\text{data} \mid \beta) = \prod_{i=1}^{N_{\text{obs}}} P(r_i/r_{0_i})$$
 (4)

where N_{obs} was the number of owner-equine pairs in the sample, and r_0 and r_i the respective *communes* of the owner and his/her equine(s) for the ith pair.

We used the MCMCpack R package (Martin et al., 2011) to generate a chain of 6,000 iterations with a burnin of 1,000 iterations. The chain's convergence was monitored by Geweke's convergence diagnostic (geweke.diag (Geweke, 1991)). We estimated two distributions of parameters β and two values for each of parameters σ , μ , from both the live and the dead equine samples.

2.4 Assignment of holding *commune(s)* to owners and equines

211

212

213

214

215

216

217

218

219

220

221

222

223

224

225

226

227

228

229

230

231

232

233

234

235

236

For each of the 1,068,142 owner-equine pairs from the equine dataset, we assigned one or more probable holding communes for equines using the previously calculated parameters and the urban coverage of the owners' commune of residence. The data (owner-equine pairs) did not allow us to quantify exactly the number of different locations for the equines of the same owner, however, the survey showed us that almost (83%) of the owners hosted their equines in one commune, 12% in two communes and the rest in 3 communes (Farchati et al., 2021). Thus, we tested two different assignment approaches. In the first one, we assigned each owner a single holding commune regardless of the number of equines linked with that owner, i.e. the holding commune selected was the same for all the equines paired with the same owner. The second approach allowed more variability in the holding communes for owners linked with more than one equine. For them, we assigned up to two or three holding communes, depending on the number of linked equines (two communes for two equines, or three communes for three or more equines). The steps were similar in both methods: for each owner, located in commune C, we randomly sampled a commune. This latter could be C (same commune) with the probability $P_0(C)$ or C' (different to C) with the probability $(1-P_0(C))*P_{sep}(C,C')$. In Method 1, this draw was directed once for each owner, and for Method 2, once for owners with one equine, twice for owners with two equines and three times for owners with three or more equines. Thus, the equines of each owner were randomly assigned to the selected

2.5 Estimation of the number of equines per *commune*

For each owner, we applied both methods and used the two distributions of the parameters obtained from both the live and the dead equine sample. A distribution of their equines locations was estimated using the MCMC method. Then, the procedure of drawing lots for a location (*commune*) described above was used to calculate a realization of this distribution (i.e. a random draw of a *commune* in the distribution of locations): it was therefore necessary to make repetitions to add variability. We performed 100 iterations for each

communes. More detailed steps are available in supplementary material Method 1 and Method 2.

method and parameter set. Finally, the number of equines assigned per *commune* for each method and parameter set was obtained by averaging the numbers obtained over the 100 iterations and adding the number of racehorses from the racehorse sample information.

3 Results

3.1 Characteristics of the samples and datasets

A description of the live equine sample (n=2,353), the dead equine sample (n=13,595), and the equine dataset (n=1,068,142) is available in Table 1. The distribution of sex among the dead equines was almost identical to those of both the live equine sample and the equine dataset, with a slight under-representation of males and over-representation of females in the dead equine sample. Saddle horses were over-represented in the live equine sample, while draft horses were under-represented in both the live and dead equine samples. The age distribution differed in the dead equine sample, with an overrepresentation of old equines (over 20 years of age) and very young equines (less than two years old). The number of equines per owner differed between the three datasets, with more owners linked to more than one equine in the dead equine sample (74% vs. 59% in the live equine sample, and 36% in the equine dataset).

The proportion of owners who kept their equines in their commune of residence decreased as the percentage of the commune's urban coverage increased (Figure 2A). In addition, the distribution of the percentage of urban coverage was the same in both samples (live and dead equines, Wilcoxon test, p >0.05). Most of the owners kept their equines in their commune of residence, but when they did keep them in a different commune, it was less than 30 km away (Figure 2B), with no difference in the distance distributions

3.2 National estimation of the spatial distribution of equines in France

between the live and dead equine samples (Wilcoxon test, p >0.05).

The estimations obtained for β differed from 0.11 (95% credibility interval: 0.11-0.12) for the estimation resulting from the live equine sample to 0.05 (95%CI: 0.05-0.06) for the estimation from the dead equine

sample (supplementary material Figure S-1). However, the estimations obtained for σ and μ were quite similar with μ = -3.06 and σ =1.07 (supplementary material Figure S-2).

The average number of equines per *commune* varied greatly between *communes* depending on the method and sample used to estimate the distribution parameters (Table 2, Figure 3).

A big difference between assignment methods 1 and 2 was observed on the live and dead equine samples (Table 3; Figure 3A, B). Using the live equine sample, only 2.4% of the 32,837 *communes* had the same number of equines with both methods (Figure 3A). Method 1 assigned more equines in 66 % of the *communes*, with a mean difference of 28.9 equines and a median difference of 13 [IQR 5-31] equines. This indicates an asymmetric distribution of differences and large differences in some *communes* (maximum difference of 10.823 equines). In the 32% of *communes* where Method 2 gave a higher number of equines, the difference was more marked, with 75% of *communes* having more than 22 equines minimum compared with Method 1. Results were similar when using the dead equine sample, with 4.3% of *communes* having the same number of equines, and 73% having more equines with Method 1, though the difference is smaller than that observed with the live equine sample (median of 7[IQR 2-17] and mean of 16.4) (Figure 3B). The distribution of the difference when Method 2 assigned more equines was quite similar to that observed with the live equine sample.

On the other hand, little difference was observed in the results according to the sample used when the same method was applied, with around 45% of *communes* having the same number of equines (Table 3; Figure 3C, D) and less variability with Method 2. This limited variability was confirmed by both the small differences observed in the number of equines, whether the live or dead equine sample was used, and the near-symmetric distributions of these differences (whether positive or negative) (median= 2/-2 [IQR 1-4] / [IQR 1-3] and mean =2.9/-2.9) (Figure 3D). With Method 1, greater variability was observed between the two samples. When using the live equine sample, Method 1 assigned more equines to a *commune*, the average difference being 36.4 equines with a maximum of 10,822 (Figure 3C). Negative differences were more limited, with an average and maximum difference of 21.4 and 37 equines respectively.

Regarding differences in spatial distribution, Method 1 resulted in higher local equine densities and did not seem sensitive to urban areas, where estimated equine densities were high (Figure 4A, C). The results were somewhat improved with the dead equine sample, which factored in urban areas better (Figure 4C). In contrast, Method 2 resulted in a more dispersed picture of equines, with low densities in most dense urban areas regardless of the sample used (Figure 4B, D). At a larger scale (*département* or *région*), regardless of the method or sample used, the estimates showed areas with higher densities in Normandy, Brittany, Paysde-la-Loire, Bouches-du Rhône, Pyrénées Atlantiques and a lower density in the Massif Central (Figure 4, Figure S-4). Finally, the addition of racehorse data to the estimates did not change the estimated spatial distributions, the areas of highest density in our estimates being those with the most racehorses (supplementary material Figure S-3).

4 Discussion

To our knowledge, this study is the first estimation of a national spatial distribution of French equines, including those kept by non-professionals, at the *commune* level. It supplements the studies conducted each year by the Economics Department of the IFCE, providing a map of the equine population (mostly kept in professional structures (Ecus-Ifce)) by *région* (the largest French administrative unit) and/or *département* (the medium French administrative unit). While promising, our results showed some sensitivity to the sample and the method used, and some of the spatial estimates obtained were quite unrealistic, with a high density of equines in highly urban areas. We discuss here the results obtained with regard to the imprecision of data used, the influence of the hypotheses and the methodology used.

4.1 Potential effect of data bias on the density estimates

Two samples were used as training data for estimating the parameters of the model, including the distribution of distances between owners and equines. The live equine sample had the same limitations as the survey itself, already mentioned in previous studies (Farchati et al., 2020, 2021). These limitations were mainly related to the bias in the representativeness of the respondents in terms of location and type of owners. Indeed, most owners in the sample kept their equine at home. Urban owners keeping horses on the outskirts of cities were thus very poorly represented, whereas they are in the majority in certain densely

populated urban areas. This probably explains the excessive number of equines obtained in urban areas, since it was considered that most owners— including those in the city—kept their equine at home. This error was exacerbated by the fact that Method 1 located all the equines of a single owner in the same commune. For example, 10,828 equines were located in Ile-de-France (Figure S-4) probably due to the presence in the Paris of the national guard equines. This hypothesis is supported by the fact that the estimate based on the dead equine sample better factored in the urban area with both methods. The dead equine sample was more variable than the live equine sample, with a few more distances over 30 km and probably more retired equines further away from their owners. Nevertheless, this sample probably overrepresented professionals and multiple owners, as we aggregated data from several years. Statistically, owners with more equines were more likely to i) be included, ii) use the ATM system for dead equine removal, iii) provide their equine identification number and iv) update their information in the SIRE database. This is confirmed by the difference in the number of equines kept, with only 25% of owners of dead equines having only one equine and 40% having more than six, compared with 41% of owners of live equines having only one equine and 18% having more than six. Additionally, despite close cross-checking of data and information to differentiate owners from keepers among the people requesting cadaver removal, some keepers (as opposed to owners) may have been retained in the dead equine sample, participating in the overrepresentation of multiple owners. This probably explained the very high density of equine estimates in French breeding areas (Normandy, Brittany, Bouches-du-Rhône and Pyrénées-Atlantique (Figure S-4)) when using Method 1 with the dead equine sample. Finally, the quality of the equine dataset used to assign the equines may have an impact on spatial estimates of equines. Indeed, a previous study of SIRE data quality (Farchati et al., 2020) indicated that nearly one-third of owners did not update their home address when they moved, while the location assignment of equines is based on this information. However, most of the owners who moved remained in the same département (Farchati et al., 2021), so the equine location error remains spatially localized (the equines distribution on a département scale (scale greater than the commune) is more robust). In addition, although we cross-

312

313

314

315

316

317

318

319

320

321

322

323

324

325

326

327

328

329

330

331

332

333

334

335

336

337

338

checked the data, it is likely that many equines in this dataset were actually dead. It is possible that the

their equine), in which case the overestimation of the number of equines is also likely to be evenly distributed across France.

4.2 Effect of the model, hypotheses and assignment methods

339

340

341

342

343

344

345

346

347

348

349

350

351

352

353

354

355

356

357

358

359

360

361

362

363

364

The main limitation in our adaptation of the work of Lo Lacono was the approximation of the location of owners and equines at the centroid of the communes because the exact (x, y) locations were not available. This probably had a strong effect on distance distribution estimates because equines and owners in the same commune can be far apart, depending on the size of the commune. Similarly, an owner and equine(s) located in different communes may be very close if they are both close to the inter-communal "border". In the first case the distance will be underestimated, while in the second case it will be overestimated. The effect of this approximation on the estimation of model parameters is not clear. The estimates of parameters σ and μ were quite similar regardless of the sample used. This similarity could be explained by the similarity of the two distributions of urban coverage for the owners who kept their equines in their commune of residence. In contrast, the estimates of the β parameter were very different from one sample to another. This difference could possibly be explained by the use of a Bayesian approach, which was sufficiently sensitive to the sample size and especially to the fact that its calculation took into account not only the urban coverage of the owner's commune of residence, but also the urban coverage of the equine holding commune, which differed for each owner-equine pair in the sample. When estimated from the dead equine sample, parameter β was much larger than when estimated from the live equine sample. It appears to be more realistic as fewer equines were assigned to high density urban areas. Furthermore, the estimation of the spatial distribution of equines was very sensitive to the assignment method used, despite a first random step to select, for each owner, a holding commune from a list of all plausible holding communes. Assignment method 1 assumed that all the equines of a given owner were kept in the same commune. This approach limited the variability of holding communes, and exacerbated the effect of the bias in the samples used to estimate β. The results showed areas of very high local equine density, including the densest urban areas such as large cities. The bias in the spatial estimation was more limited when using the dead equine sample, the dataset including more variability. Assignment method 2

allowed more variability in the holding *commune* for owners having more than one equine. This approach resulted in a highly dispersed spatial distribution regardless of the sample used. Interestingly, the high density observed in large cities with Method 1 when using the live equine sample was no longer observed. Besides, Method 2 appeared less sensitive to the sample used, as differences between the estimations were limited (around two equines on average when a difference was in fact observed). The variability probably counterbalanced the bias in the samples used to estimate the distribution of distances.

5 Conclusion and prospects

Three of the four estimations of the spatial distribution of French equines are consistent with information collected by the IFCE (Ecus-Ifce). It is probable that the reality lies somewhere between the results obtained using the two assignment methods. With very little difference between the results obtained using Method 2 with both samples and a more realistic map with Method 1 and the dead equine sample, our results show that the location of dead equines could be used to estimate the location of live equines. This could be extremely useful in a disease outbreak situation in order to effectively control the spread prior to implementation in the field of the EU's Animal Health Law. This work could be a source of inspiration for many countries affected by the same problematic as ours. An evaluation of the relevance of our estimates is therefore planned in the near future by confronting our estimates with the surveys carried out in the field to check the traceability of equines in France, thanks to the censuses carried out by the IFCE.

	1	Dead equ	ine sample	Live equ	uine sample	Equine	dataset
		(n=13,595)		(n=2,353)		(n=1,068,142	2)
Information	Modalities	n	%	n	%	n	%
Sex	NA	0	0.0	0	0	52	0.0
	Female	7 938	58.4	1234	52.4	575099	53.8
	Gelding	3 444	25.3	640	27.2	205426	19.2
	Male	2 213	16.3	479	20.4	287565	26.9
Age*	NA	5011	0.0	448	0.0	0	0.0
	0	115	1.3	0	0.0	0	0.0
	[1-2]	938	10.9	144	7.6	81324	7.6
	[3-10]	1850	21.6	678	35.6	402158	37.7
	[11-20]	2545	29.6	853	44.8	412812	38.6
	[21-30]	2820	32.9	226	11.9	151357	14.2
	>30	316	3.7	4	0.2	20491	1.9
Group of breeds	Donkey	822	6.0	70	3.0	46194	4.3
	Racehorse	1923	14.1	341	14.5	216041	20.2
	Pony	3178	23.4	486	20.7	150962	14.1
	Saddle horse	6360	46.8	1362	57.9	467460	43.8
	Draft horse	1312	9.7	94	4.0	187485	17.6
Number of equine(s)	1	1689	25.4	826	41.1	251939	63.5
per owner	[2-5]	2222	33.5	814	40.5	112468	28.3
	[6-20]	1801	27.1	290	14.4	26875	6.8
	[21-100]	828	12.5	76	3.8	5437	1.4
	>100	98	1.5	2	0.1	320	0.1

Total number of different owners 6638	100	2008	100	397039	100
---------------------------------------	-----	------	-----	--------	-----

* In years; age at death for a dead equine, or age at the date of data extraction for a live equine

TABLE 2. Distribution of the average number of equines (among the 100 simulations) per *commune* depending on the assignment method and sample used to estimate the parameters

Sample	Method	Minimum	1st quartile	Median	Mean	3rd	Maximum
23			•			quartile	
Live equines	1	0	6	13	30.7	33	10828
Live equilies	2	0	2	7	30.6	36	1313
Dead equines	1	0	5	24	30.8	37	2810
Dead equiles	2	0	2	8	30.6	37	1418

TABLE 3. Comparison of the results of the models: distributions of the difference in the average number of equines assigned to each *commune* according to the samples and models.

Comparison	Method or sample concerned	Distribution of the difference	No difference	Positive difference	Negative difference
Live equine vs.	Method 1	%(number of <i>communes</i>)	43.2 (15051)	20.8 (7257)	36.0 (12571)
dead equine		Minimum		1	-1
sample		1st quartile		5	-22
		Median		15	-24
		Mean		36.5	-21.4
		3rd quartile		38	-26
		Maximum		10822	-37
	Method 1	%(number of <i>communes</i>)	47.9 (16718)	26.0 (9073)	26.1 (9088)

			Minimum		1	-1
			1st quartile		1	-1
			Median		2	-2
			Mean		2.9	-2.98
			3rd quartile		4	-3
			Maximum		50	-143
Method 1 vs.	Live equine	%(number	of communes)	2.4 (841)	65.7 (22910)	31.9 (11128)
Method 2	sample		Minimum		1	-1
			1st quartile		5	-22
			Median		13	-42
			Mean		28.98	-59.2
			3rd quartile		31	-74
			Maximum		10823	-457
	Dead	%(number	of communes)	4.3 (1514)	73.4 (25612)	22.2 (7753)
	equine		Minimum		1	-1
	sample		1st quartile		2	-14
			Median		7	-33
			Mean		16.4	-53.5
			3rd quartile		17	-66
			Maximum		1392	-423

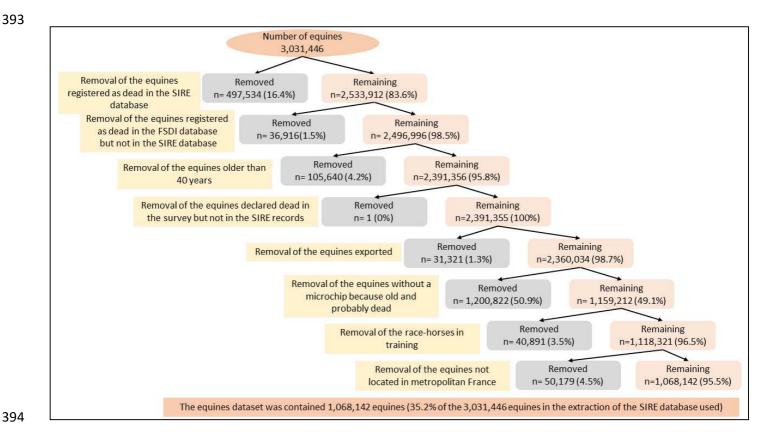


Figure 1. Different steps determining the number of equines assumed alive and used to build the equines dataset.

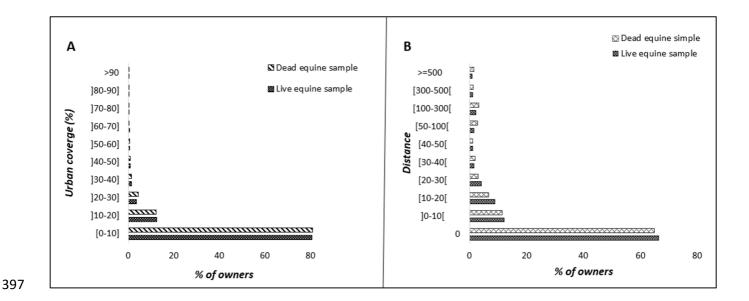


Figure 2. Distribution of the proportion of owners keeping their equine in their commune of residence according to the urban coverage (A) and distribution of the proportion of owners (B) according to the distance between the residence and holding communes in the live and dead equine samples

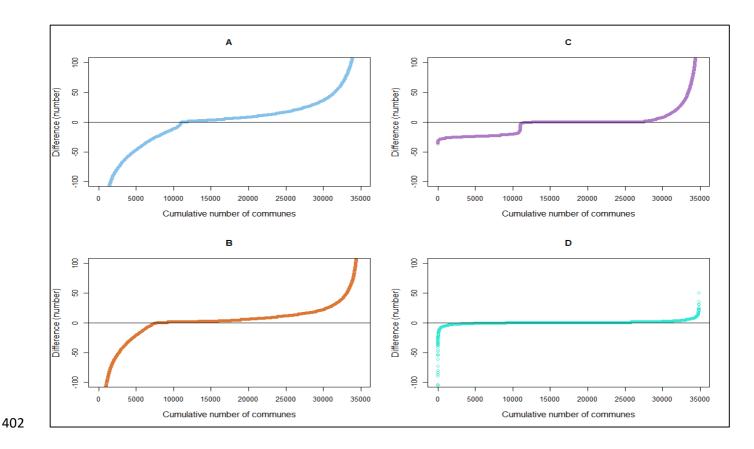


FIGURE 3. Difference in the average number of equines per *commune* according to the sample and method used (restricted to values between 100 and -100): (A) live equine sample Method 1 vs. Method 2, (B) dead equine sample Method 1 vs. Method 2, (C) Method 1 live equine sample vs. dead equine sample and (D) Method 2 live equine sample vs. dead equine sample

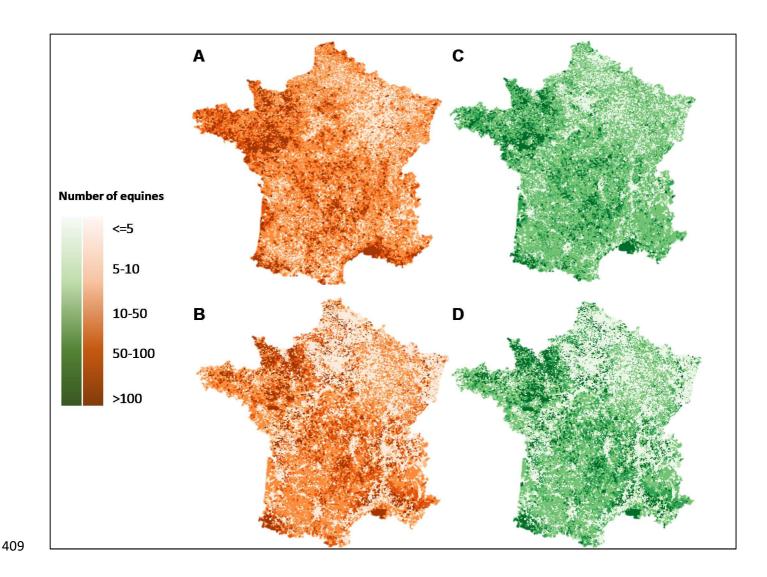


FIGURE 4. Spatial distribution of the equine population in France estimated from (A) Method 1 and live equine sample, (B) Method 2 and live equine sample, (C) Method 1 and dead equine sample and (D) Method 2 and dead equine sample (D) after inclusion of racehorse data.

416 Reference

- 417 Amat, J.P., Tapprest, J., Gaudaire, D., Merlin, A., Strugar, S., Coulibaly, E., Grandcollot-Chabot, M., Hans, A.,
- 418 2018. Deux cas d'anémie infectieuse des équidés détectés dans le sud-est de la France : point de situation
- 419 n°2 (27/03/2018) Platforme ESA.
- 420 Anonyme, 2016. Règlement 2016/429 du Parlement Européen et du Conseil du 9 mars 2016 publiée au
- **421** journal officiel de l'Union Européenne le 31 mars 2016.
- 422 Brendan, C., Michael, P.W., Samuel, H., Greame, G., 2009. The equine influenza epidemic in Australia: spatial
- and temporal descriptive analyses of a large propagating epidemic. Prev. Vet. Med. 92, 60-70.
- 424 Callinan, I.H., 2007. Equine influenza. The August 2007 outbreak in Australia. https://nla.gov.au/nla.obj-
- 425 961843962/view?partId=nla.obj-962288923#page/n5 /mode/1up, (accessed on 30 September 2021).
- 426 Cover, C.L., Corine Land Cover : données d'inventaire biophysique de l'occupation des sols et de son
- 427 évolution selon une nomenclature en 44 postes. https://www.statistiques.developpement-
- durable.gouv.fr/corine-land-cover-0, (accessed on 30 September 2021).
- 429 Ecus-Ifce, Annuaire écus, IFCE, Institut Français du Cheval et de L'équitation. https://www.ifce.fr/recherche-
- 430 transverse/?q=ANNUAIRE+%C3%89CUS&cb=https://www.ifce.fr/, (accessed on 30 September 2021).
- Farchati, H., Merlin, A., Sassac, M., Dornier, X., Dhollande, M., Garon, D., Tapprest, J., Sala, C., 2020. Is the
- 432 French SIRE equine information system a good basis for epidemiological and modeling research? Quality
- assessment using two surveys. Res. Vet. Sci 134, 96–101.
- 434 Farchati, H., Merlin, A., Sassac, M., Dornier, X., Dhollande, M., Garon, D., Tapprest, J., Sala, C., 2021. Home
- 435 Sweet Home: New Insights Into the Location of Equine Premises in France and Keeping Habits to Inform
- 436 Health Prevention and Disease Surveillance. Front Vet Sci. 23, 8:701749.

- 437 Ferguson, N.M., D.A.T., C., S., C., Fraser, C., Riley, S., Meeyai, A., Iamsirithaworn, S., Burke, D.S., 2005.
- 438 Strategies for containing an emerging influenza pandemic in Southeast Asia. Nature 437, 209–214.
- 439 Ferguson, N.M., Donnelly, C.A., Anderson, R.M., 2001. Transmission intensity and impact of control policies
- on the foot and mouth epidemic in Great Britain. Nature 413, 542–548.
- 441 Garner, M.G., Cowled, B., East, I.J., Moloney, B.J., Kung, N.Y., 2010. Evaluating the effectiveness of early
- vaccination in the control and eradication of equine influenza a modelling approach. Prev. Vet. Med. 99,
- **443** 15-27.
- Geweke, J.F., 1991. Evaluating the accuracy of sampling-based approaches to the calculation of posterior
- moments. In Bayesian Statistics 4 (ed JM Bernado, JO Berger, AP Dawid and AFM Smith). Clarendon Press,
- 446 Oxford, UK.
- 447 Gudelj, I., K.A.J., W., 2004. Spatial heterogeneity, social structure and disease dynamics of animal
- populations. Theor Popul Biol, 66, 139–149.
- House, T., Keeling, M.J., 2008a. Deterministic epidemic models with explicit household structure. Math.
- **450** Biosci. 213, 29–39.
- 451 House, T., Keeling, M.J., 2008b. Household structure and infectious disease transmission. Epidemiol. Infect.
- **452** 137, 654–661.
- 453 Keeling, M.J., Danon, L., Vernon, M.C., House, T.A., 2010. Individual identity and movement networks for
- disease metapopulations. Proc. Nat.l Acad. Sci. U.S.A. 107, 8866–8870.
- Keeling, M.J., Woolhouse, M.E., Shaw, D.J., Matthews, L., Chase-Topping, M., Haydon, D.T., Cornell, S.J.,
- 456 Kappey, J., Wilesmith, J., Grenfell, B.T., 2001. Dynamics of the 2001 UK foot and mouth epidemic: Stochastic
- dispersal in a heterogeneous landscape. Science 294, 813–817.

- Lo Iacono, G., Robin, C.A., Newton, J.R., Gubbins, S., Wood, J.L., 2013. Where are the horses? With the sheep
- 459 or cows? Uncertain host location, vector-feeding preferences and the risk of African horse sickness
- transmission in Great Britain. J. R. Soc. Interface 10, 20130194.
- 461 Martin, A.D., Quinn, K.M., Park, J.H., 2011. MCMCpack: Markov chain Monte Carlo in R. J. Stat. Software 42,
- **462** 323.
- 463 Merlin, A., Gaudaire, D., Letorey, L., Delerue, M., Amat, J.P., Lecouturier, F., Madeline, A., Deshiere, A.,
- 464 Aubert, C., Poudevigne, F., Quinio, C., Martin, F., Strugar, S., Coulibaly, E., Depecker, A., Grancollot-Chabot,
- 465 M., Hans, A., 2021. Synthèse des foyers d'anémie infectieuse des équines reportes en France entre 2017 et
- **466** 2019.
- Swinton, J., Harwood, J., Grenfell, B.T., Gilligan, C.A., 1998. Persistence thresholds for phocine distemper
- virus infection in harbour seal Phoca vitulina metapopulations. J Anim Ecol 67, 54–68.
- TRACE, The Trade Control and Expert System. https://ec.europa.eu/food/animals/traces_en, (accessed on 30
- **470** September 2021).