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Title: Epidemiology of Leptospirosis in North-Central Italy: fifteen years of serological data (2002-2016)

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Abstract: Leptospirosis is a re-emerging bacterial zoonosisCentral Italy is characterized by a geographic area that promote Leptospira circulation.: Data on seroepidemiological survey carried out from 2002 to 2016 in Central Italy were reported and discussed. Overall, 709 out of the 8488 (8.35%) tested sera were positive for Leptospira at the cut-off titer (1:100) and 218 (2.57%) at higher titer ( $\geq$ 1:400). The highest percentages of positivity was recorded for coypus (22.86%), swine (19.74%) and bovine (, 13.03%). Pomona and Bratislava resulted the serovars more often detected, followed by Harjo and Icterohaemorragiea; while, a low number of positive sera was detected for serovars Ballum, Canicola and Tarassovi.. .. Percentage of positive sera for each year slightly decreased from 2002 to 2008 and rose from 2009., a rise in this; particularly, high percentages of positive reaction were recorded in 2014 (17.23%), 2015 (19.61%) and 2016 (38.05%). Bertelloni et al., *Epidemiology of Leptospirosis in North-Central Italy: fifteen years of serological data (2002-2016)*.

### <u>Highlights</u>

- Fifteen years of serological data on *Leptospira* in animals were presented;
- An increase positivity to *Leptospira* in last years of investigation was recorded;
- Rise or fall of some serovars were observed;
- Some changes in hosts/serovars relations were detected;

# 1 Epidemiology of Leptospirosis in North-Central Italy: fifteen years of serological

# 2 data (2002-2016)

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- 14
- 15 **Running Title**: *Leptospira* in North-Central Italy 2002-2016
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#### 34 **1. Introduction**

35 Leptospirosis represents one of the most important widespread re-emerging bacterial zoonosis [1–3]. Several outbreaks accounting for thousands of deaths worldwide highlight the importance of leptospirosis 36 37 as a severely neglected infectious disease [4,5]. Leptospirosis has a wide distribution and occurs overall in 38 tropical, subtropical and temperate zones, favoured by a large variety of both wild and domestic mammals 39 which can play the role of natural reservoirs of Leptospira [1,6]. Some animals are asymptomatic renal 40 carriers of this bacterium and they contribute to maintain the infection in a particular environment by constantly shedding Leptospira with urine [7,8]. Accidental contact with Leptospira infected urine causes 41 42 the incidental infection and produces clinical diseases. While, specific Leptospira serovars which show close 43 relationship with particular animal species develop host-maintained infection. The maintenance host 44 generally does not develop symptoms, except after long time, but it acts as a natural source of a specific 45 serovar [6,8]. In fact, Leptospira epidemiology is strictly related to the presence and widespread of the 46 maintenance hosts species [9]. In recent years, some serovars seem to be prevalent and emerging, 47 especially among wild animals, but also in domestic species. This occurrence suggests that the 48 epidemiology of leptospirosis may change over time in animals as well as in humans [10]. 49 Central Italy, and in particular Tuscany, is a geographic area characterized by some peculiarities which 50 promote the presence and persistence of *Leptospira* in hosts and in environment: a) presence of wild 51 animals which could represent potential reservoirs; b) presence of domestic animals raised in semi-52 extensive or extensive farms, which promotes contact with wild species; c) a significant presence of hunting 53 activity; d) abundance of wetlands such as marshes, ponds and irrigation canals. 54 The main purpose of the present work was to refer data on a sero-epidemiological survey carried out in 55 Central Italy, particularly in Tuscany, on serum samples collected from 2002 to 2016 in order to assess the 56 prevalence of leptospirosis in domestic and wild animals and to compare it with the data from a previous 57 epidemiological surveillance investigation carried out in the same area between 1995 and 2001 [9]. 58

#### 59 2. Material and Methods

60 From January 2002 to December 2016, 8488 serum samples were analysed for Leptospira. Sera were 61 collected from healthy and ill animals belonging to 23 different species: sheep (Ovis aries, 2682 sera), swine 62 (Sus scrofa, 1332 sera), bovine (Bos Taurus, 1328 sera), dog (Canis lupus familiaris, 1144 sera), wild boar 63 (Sus scrofa, 479 sera), goat (Capra hircus, 327 sera), european brown hare (Lepus europaeus, 162 sera), red 64 fox (Vulpes vulpes, 94 sera), horse (Equus caballus, 74 sera), roe deer (Capreolus capreolus, 74 sera), coypus 65 (Myocastor coypus, 70 sera), fallow deer (Dama dama, 65 sera), donkey (Equus asinus, 2 sera), ferret 66 (Mustela putorius furo, 1 serum), cat (Felis catus, 4 sera), red deer (Cervus elaphus, 56 sera), wolf (Canis 67 lupus, 43 sera), rats (Rattus norvegicus, 34 sera), mouflon (Ovis musimon, 8 sera), mouse (Mus musculus, 8 68 sera), Guinea pig (Cavia porcellus, 2 sera), alpaca (Vicugna pacos, 1 serum) and bear (Ursus arctos, 1 69 serum). Furthermore, 329 sera were of human origin and collected from men with clinical symptoms 70 referable to leptospirosis. All sera were collected in North-Central Italy. Serological investigation was 71 carried out with Microscopic Agglutination Test (MAT) [11]. The following serovars were used as live 72 antigens: Icterohaemorrhagiae (strain Bianchi), Canicola (strain Alarik), Pomona (strain Mezzano), Tarassovi 73 (strain Mitis Johnson), Grippotyphosa (strain Moscow V), Bratislava (strain Riccio 2), Ballum (strain 74 Castellon 3) and Hardjo (strain Hardjoprajitno). Titers of 1:100 were considered positive; 2-fold serial 75 dilutions were tested to determine the endpoint titer.

76

#### 77 **3. Results**

78 Overall, 709 out of the 8488 (8.35%) sera resulted positive for Leptospira at the breakpoint titer (1:100). All 79 samples collected from the following species resulted negative: alpaca, donkey, guinea pig, mouse, rat, 80 ferret, cat, fallow deer, red deer, roe deer, mouflon, bear and wolf. Two hundred and eighteen sera (2.57%) 81 resulted positive at high titer (≥1:400). Table 1 reports the number of positive sera grouped by animal 82 species. The highest percentages of positive sera were recorded for coypus, swine and bovine at low 83 (22.86%, 19.74% and 13.03%, respectively) and high titer (5.61%, 6.31% and 3.16%, respectively). None 84 horse sera showed positivity to titer of 1:400 or higher. Coypus, wild boar, fox and hare resulted the unique 85 wild species that showed positive reactions. In particular, 22.86%, 8.56%, 3.04% and 1.85% of sera resulted

86 positive at titer ≥1:100 for coypus, wild boar, fox and hare, respectively. Among the 329 human sera 87 examined, 10 (4.26%) resulted positive and 3 (1.06%) showed a titer of 1:400 or higher. 88 Table 2 shows the distribution of positive sera grouped by serovars and animal species. Pomona and 89 Bratislava resulted the serovars more often detected, with 267 and 224 positive reactions (titer 1:100), 90 respectively. Moreover, for the same serovars, 68 and 67 sera, respectively showed positivity to high titer 91 (≥1:400). A low number of positive sera was detected for serovars Ballum, Canicola and Tarassovi (13, 28 92 and 32 sera, respectively). Hare was the only species in which positivity to serovar Icterohaemorrhagiae 93 was never found. Similarly, goat resulted the only species in which serovar Bratislava was never detected. 94 Positivity to serovars Icterohaemorrhagiae and Canicola was found mainly in dog sera. Serovars Pomona 95 and Bratislava resulted more often associated with swine. Most of the sera positive to Tarassovi were from wild boar. Positivity to serovars Grippotyphosa and Hardjo was detected especially in bovine sera. 96 97 Among human sera, positivity to six serovars was recorded: Icterohaemorrhagiae, Canicola, Pomona, 98 Grippotyphosa, Bratislava and Ballum. Icterohaemorrhagiae was the most detected serovar. Positive 99 samples were never found among human sera from 2004 to 2012. In 2013, 2014 and 2016 positive sera 100 were 1/20, 4/9 and 1/11, respectively; all these sera showed an antibody titer <1:400. 101 Percentage of positive sera for each year slightly decreased from 2002 to 2008; in particular in 2008 only 1 102 out of 597 (0.17%) sera resulted positive at titer of 1:100. From 2009, the percentage of positive sera 103 increased; particularly, in 2014, 2015 and 2016 a high percentage of positive sera was recorded: 17.23%, 104 19.61% and 38.05%, respectively. Figure 1 reports the percentage of positive sera to *Leptospira interrogans* 105 and to each serovar detected in each year of investigation. 106 Table S1 (supplementary material) reports the distribution of positive sera at titer  $\ge 1:100$  and  $\ge 1:400$  in 107 relation to animal species, *Leptospira* serovars and year of detection.

108

#### 109 4. Discussion

110 Leptospirosis is a worldwide public health and veterinary problem, frequently underestimated,

111 characterized by a downward trend [12]. Climatic changes, modifications of ecological niches, emergence

112 of new potential maintenance-hosts could represent the most important factors involved in Leptospira 113 epidemiology. The environmental and geographic features of North-Central Italy area can be considered as the optimal conditions for Leptospira spreading [13–21]. Tuscany, the main region involved in this 114 115 serological survey, is characterized by some peculiarities which promote the presence and persistence of 116 Leptospira in hosts and in the environment. Increasing presence of wild animals, potential reservoirs, 117 presence of animals raised in semi-extensive or extensive farms, hunting activity and presence of wetlands 118 such as marshes, ponds, lakes and irrigation canals are the main factors involved in enzootic trend of 119 leptospirosis in Italy.

120 In this study, serological results obtained whit a panel of eight *Leptospira* serovars were reported. Strains

121 employed as live-antigens was chosen considering the serovars more often detected, by isolation or

serology, in Italy [10,13,17,18,22–24]; indeed, positivity to other serovars were rarely reported [25].

Moreover, these strains were employed routinely in our Laboratory for research and diagnosis, and, for this
 reason, all serum samples considered were tested with same MAT antigens.

125 In this survey, the total apparent prevalence of antibodies against Leptospira registered during 2002-2016 126 was 8.65%. In a previous investigation, carried out in the same area during 1995-2001, the total apparent 127 prevalence was 6.81% [9]. It is not possible to exclude that this increase could be due to the different 128 number of available samples and the proportion and representation of the different animal species 129 included in the study. However, the increase could be also related to a changing of some environmental 130 conditions (rainfall, temperature, ecc...) which promoted the rise and spreading of new serovar/strains or 131 the re-emerging of endemic strains. The observed increase of positive sera was mainly related to the last years of this investigation (2013-2016) and could be due to an increase of rainfall in the investigated area, 132 133 especially in wetlands.

134 Serological investigations highlighted different apparent prevalence trends for each of the eight *Leptospira* 

serovars tested. Apparent prevalence of positive sera decreased from 1995-2001 to 2002-2016 for

136 Icterohaemorrhagiae (from 22.24% to 13.78%), Bratislava (from 55.04% to 24.69%) and Hardjo (from

137 22.08% to 14.22%). Furthermore, apparent prevalence increases for Canicola (from 0% to 3.08%), Pomona

(from 0.64% to 29.43%), Grippotyphosa (from 0% to 9.81%), Tarassovi (from 0% to 3.52%) and Ballum (from
0% to 1.43%) [9]. The results obtained from previous and from this investigation could be different for
some reasons. In the previous work [9], carried out in the same investigated area, employing the same *Leptopsira* serovars and strains as antigens, the threshold titer was 1:400, conversely in this investigation
the threshold titer used was 1:100. Furthermore, the number of tested sera for each animal species could
have influenced the detected total apparent prevalence.

144 Considering the distribution of positivity detected year by year, it is possible to observe a decrease in the 145 percentage of positive sera detected from 2002 to 2008 and an increase starting from 2009. In particular,

146 from 2014 it was registered an impressive increase of *Leptospira* positive samples (Figure 1a).

147 Annual fluctuations in *Leptospira* spreading and seropositivity is well documented [26] and could be related

to many factors as climate changing, rains and drought springs. Fluctuations in observed data could be

related not only to environmental modifications, but also to hosts changing: variations in exposure to

150 *Leptospira* by animal, introduction of new wild animal species, changing in herds management. As concerns

151 this last point, in lasts years breeding management changed from indoor intensive to extensive or semi-

152 extensive with outdoor access in order to improve animal welfare. Furthermore, it could be related to

153 modifications of *Leptospira* epidemiology: introduction of new serovars/strains or change in host specificity

154 range by classical serovars/strains.

155 As for serovar Icterohaemorrhagiae, seropositivity reflected more or less the observed global trend.

156 However, positivity to this serovar was constantly detected, even if during some years with only few

157 positive samples (Figure 1b). This is an expected result, indeed, Icterohaemorrhagiae is maintained by rats

and it is the serogroup most often involved in animal and human infection in many parts of the world [27].

159 Only few samples scored positive to serovar Canicola during the first years of investigation and no positive

sera were detected between 2006 and 2012 (Figure 1c). It seems that this serovars re-emerged from 2013.

- 161 In nature, Canicola is maintained by dog and vaccination programs carried out for about 50 years in this
- species led to its disappearance [28]. Data obtained in this investigation showed in recent years an increase

of positivity to Canicola in animals different to dogs. This finding could suggest a possible change in hostrange of this serovar.

Serovar Pomona seemed to be almost disappeared from the investigated area for more than fifteen years 165 (Figure 1d). Ten positive reactions were recorded from 2002 to 2013, furthermore, a previous investigation 166 167 conducted in the same geographical area [9], reported an apparent prevalence of 0.64% (4 out of 9885 168 sera) between 1995 and 2001. From 2014, the number of positive sera increased, and Pomona was the 169 most detected serovar in 2016. In the last years of the investigation, a large number of swine sera were 170 analysed, and this could have influenced the improved detection of positivity to Pomona. However, a 171 considerable number of positive reactions was found also in sera of species other than pig, supporting the 172 hypothesis of the rise of this serovar in investigated area.

173 Circulation of serovar Grippotyphosa in the investigated area was slightly documented during this

investigation. This is in accordance with previous reports conducted in the same region and in Italy [9,10].

175 From 2013, a slight increase in the detected number of positive reactions was registered (Figure 1e). This

trend is in accordance with many studies conducted in Europe, where Grippotyphosa is considered an

177 emerging serovar [29–32]. Considering the studied area, it is not possible to exclude that this occurrence

178 could be related to the import from East Europe of wild animals for hunting purposes, in particular hares179 [33,34].

Positivity to serovar Tarassovi was never detected from 2002 to 2013 (Figure 1f). This trend is in line with other National surveys [9,10], indeed, in past years, Tarassovi showed a very limited diffusion in Italy. In 2014, a peak of positivity was registered which could resemble an epidemic event. Positive reactions were found mainly in wild boar, but also in domestic animals. In 2015 and 2016, the number of positive sera decreased and probably it will return to zero. These results suggest that Tarassovi is not disappeared from our territory, and infections could occur, since Tarassovi is probably maintained in the environment by *reservoirs* other than swine, as suggested by previously reported data in Italy [10].

As regard serovar Bratislava, its trend reflects the global trend observed. Every year, positivity to this
 serovar was detected, with exception of 2008 and 2011 (Figure 1g). This finding could probably be expected

considering the high number of positive *reservoir* hosts present in investigated area, as hedgehog and wild
 boar [35,36]. Generally, Bratislava is reported as an emerging serovar, but considering the obtained data it
 could be considered as endemic in investigated area.

192 Considering the abundance of cattle and, especially, sheep herds in the studied area, the low number of 193 positivity for serovar Hardjo for many years was unexpected. Indeed, this serovar were frequently detected 194 in Italy and sometimes it was involved in clinical outbreaks [9,10,25]. As for the other serovars, after a silent 195 period of about 12 years, from 2014, an increase of positive sera for Hardjo was registered, suggesting a 196 restart of the circulation of this serovar in investigated area. More focused investigations should be 197 probably required to better understand this finding. It may be supposed it could be related to the increase 198 of outdoor herds, especially for dairy cows, related to the increased attention for animals welfare. 199 Furthermore, animal species different to domestic ruminants could be infected and this could contribute to 200 the rise of serovar Harjo. 201 During the investigated period, serovar Ballum showed a very limited diffusion. In 2014, a slight increase of

seropositivity was registered, but it was a limited event. In Italy, this serovar was rarely detected in
serological investigations [9,10,13,18,22], even if it was recently isolated from small mammals in Tuscany
[24]. These observations could suggest a low virulence for animals and man of Ballum strains circulating in
our territory.

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#### 207 <u>4.1. Bovine</u>

The total apparent prevalence in bovine increase from 0.005% (1995-2001) [9] to 13.03% (2002-2016). In cattle, higher prevalence of serovar Hardjo has been detected: 7.08%8 (94/132) and 2.26% (30/1328) sera scored positive at titer  $\geq$  1:100 and  $\geq$  1:400, respectively. These data confirm that bovine represents the main maintenance-host for serovar Hardjo (serogroup Sejroe) [7]. Relative high number of positive reactions was also detected for serovars Pomona and Grippotyphosa in cattle sera. A percentage of 4.15% (55/1328) and 0.23% (3/1328) were positive for Pomona at titer  $\geq$  1:100 and  $\geq$  1:400, respectively. Severe infections in cattle due to this serovar is uncommon and usually occur in young animals [7]. Nevertheless,

215 based on a recent National survey, Pomona resulted the second representative serovar in cattle in Italy 216 [10]. The relative high number of positivity could be related to the semi-extensive or extensive farms. These 217 types of breeding promote the contact with wild animal, in particular wild boars. As concern serovar 218 Grippotyphosa, 3.61% (48/1328) and 0.38% (5/1328) sera scored positive at titer  $\geq$  1:100 and  $\geq$  1:400, 219 respectively. Positivity to this serovar resulted higher than that observed in previous investigations 220 conducted in Italy [9,10]. However, in recent years, in other European countries, serovar Grippotyphosa 221 was often detected in cattle and occasionally involved in clinical leptospirosis outbreaks [26,29,30]. In 222 accordance with these Authors, our data seem to identify Grippotyphosa as an emerging serovars in cow. 223 Low level of positivity was scored for serovars Icterohaemorrhagiae, Canicola, Tarassovi, Bratislava and 224 Ballum, in accordance with other studies [9,10].

225

226 <u>4.2. Dog</u>

227 The total apparent prevalence in dog increased from 5.42% (1995-2001) to 8.65% (2002-2016) [9]. 228 Icterohaemorrhagiae and Bratislava resulted the most represented serovars, with 5.07% (58/1144) and 229 4.19% (48/1144) sera positive at titer  $\geq$  1:100, respectively. These results are in accordance with data 230 available in literature [7,9,10]. Percentages of 1.40% (16/1144) and (0.17% 2/1144) of sera scored positive 231 for serovar Pomona at titer ≥ 1:100 and ≥ 1:400, respectively. Infection by serovar Pomona in dog produce 232 a severe disease characterized by lethargy, fever, inappetence, diffuse haemorrhage, renal and liver failure 233 [37,38]. In Europe, infections in dogs caused by this serovar are rare and reported only in few countries of 234 East-Europe, such as Romania [7]. For this reason serovar Pomona was not included in dog vaccines [39]. 235 Our data seem to suggest an increasing incidence of this serovar in dog during last years (Table S1) 236 [9,10,13]. Taking into account the increase of Pomona positivity in dog and the severe symptoms, the 237 research on this serovar could be intensify. As regards serovar Canicola, 1.22% (14/1144) and 0.09% 238 (1/1144) sera resulted positive at titer  $\geq 1:100$  and  $\geq 1:400$ , respectively. This result confirms the decreasing 239 of this serovar in many European countries during the last years due to the use of vaccines [7,40]. Low 240 number of positive sera was observed for Grippotyphosa and Tarassovi with 0.35% (4/1144) and 0.09%

(1/1144) of positive sera at titer ≥ 1:100, respectively. In Europe, Grippotyphosa is considered an emerging
serovar in dogs [13,29,31,32] and was included in leptospirosis dog vaccine. Contrary to what is generally
observed in Europe, data obtained by this investigation suggest a limited spreading of Grippotyphosa in
dogs among investigated area. In Europe, Tarassovi is rarely reported in dog [10,17]. This is in accordance
with our data, since only one serum was found positive to this serovar, even if at high titer (1:800)
suggesting a recent and probably acute infection. No positivity was encountered for serovars Hardjo and
Ballum, which are rarely detected in dog as suggested by several investigations [7,10,40].

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249 <u>4.3. Swine</u>

250 The total apparent prevalence in swine increased considerably from 9.16% (1995-2001) [9] to 19.74% 251 (2002-2016). It is known that swine is the maintenance host for serovar Pomona (serogroup Pomona), 252 serovar Tarassovi (serogroup Tarassovi) and serovar Bratislava (serogroup Australis) [7]. As expected, 253 Pomona and Bratislava were the serovars more often detected in this species. For Pomona, 12.01% 254 (160/1332) and 3.38% (45/1332) of swine sera have been detected positive at titer  $\ge 1:100$  and  $\ge 1:400$ , 255 respectively. While for Bratislava, 9.83% (131/1332) and 2.33% (311332) of sera scored positive at titer ≥ 256 1:100 and  $\geq$  1:400, respectively. Bratislava is considered an emerging serovar and it could be the cause of 257 abortion and other reproductive disorder in swine. However, some strains become "pig-adapted", causing 258 subclinical infections [41]. In different European countries, during last years, Bratislava was the more 259 detected serovar in pig sera samples [10,42]. Despite the fact that swine represents the *reservoir* host for 260 Pomona, in last years, the seroprevalence of this serovar in pig was low [9,10,42]. It is noteworthy that, in 261 this investigation, the seroprevalence of Pomona in swine was increasing, in contrast with other surveys. 262 On the other hand, only 0.15% (2/1332) and 0.08% (1/1332) of sera were detected positive for Tarassovi at 263 titer  $\geq$  1:100 and  $\geq$  1:400, respectively. These data confirm that this serovar seems to disappear, as 264 suggested by other investigations [9,10,42]. The cause could be the wide use of vaccination program in 265 swine farm [43,44]. As regard serovar Canicola, 0.45% (6/1332) of sera scored positive at titer 1:100/1:200 266 (3 at titer of 1:100 and 3 at titer of 1:200). Some studies demonstrated that swine could be infected by

serovar Canicola and that intraspecies transmission is possible [42,45]. For this reason, pigs are considered
a new possible potential maintenance host for serovar Canicola, although its real epidemiological role is not
still clarified [46,47]. The results of this investigation confirm the circulation of this serovars among swine in
Italy too. No positive sera were detected for serovars Grippotyphosa and Ballum, and only 0.15% (2/1332)
and 0.08% (1/1332) of sera resulted positive for Icterohaemorrhagiae and Hardjo, respectively. These data
are in accordance with other studies [9,10,42].

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274 <u>4.4. Horse</u>

275 The total apparent prevalence obtained in horse was characterized by a remarkable decrease from 11.08% 276 (1995-2001) [9] to 2.89% (2002-2016). This species is susceptible to a wide range of incidental infections, 277 that are often characterized by absence of clinical symptoms. Bratislava is the most common serovar 278 detected in horses, but also Grippotyphosa, Pomona, Icterohaemorrhagiae, Autumnalis, Sejroe, Canicola, 279 and Ballum serogroups are occasionally reported [7,48]. The results of this investigation confirm this 280 evidence; indeed, positive reactions were detected only for Icterohaemorrhagiae (1.24% - 3/242 sera), 281 Bratislava (0.83% - 2/242 sera), Pomona (0.41% - 1/242 sera) and Hardjo (0.41% - 1/242 sera). The low 282 number of positive sera detected in horse could be related to few clinical samples or few specimens with 283 clinical manifestations. Also, the sera could have been collected after equine uveitis manifestation. Equine 284 uveitis, also known as "moon blindness", is an ocular disease consequent to Leptospira infection in horse, 285 that occurs after the acute phase of leptospirosis when the antibody titer decrease [48].

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#### 287 <u>4.5. Sheep and Goat</u>

The total apparent prevalence observed in sheep decreased from 12.13% (1995-2001) [9] to 3.13% (2002-2016). These results could seem unexpected. Indeed, in investigated area sheep flocks are breed extensively or semi-extensively and this could promote contact with wild animals. However, our data are in line with a recent survey conducted by the Italian reference center for leptospirosis [10]. Despite sheep represent the second maintenance host for serovar Hardjo [49], infection by this serovar could induce

293 subclinical or clinical disease [7], with abortion, stillbirth, birth of weak lambs, agalactia and infertility [50-294 52]. Although the results of this investigations do not show a high number of positive samples (0.71% -295 19/2682) for serovar Hardjo, almost all have a titer  $\geq$  1:400 (12 sera out of 19 positive). These data could 296 suggest the occurrence of clinical infection in sheep confirming the high virulence of serovar Hardjo for this 297 animal species. Sheep could be infected also by other serovars [7]. In this investigation, relative high 298 number of positivity was reported for serovars Pomona (1.12% 30/2682), Icterohaemorrhagiae (0.75% -299 20/2682) and Grippotyphosa (0.75% - 20/2682) at titer  $\geq$  1:100. These results are in disagreement with 300 previous investigations carried out in the same area [9] and in Italy [10,25]. Pomona and 301 Icterohaemorrhagiae could cause disease in sheep, in particular they have been associated with 302 reproductive disorders [53,54]. Furthermore, other positive reactions were found in sheep sera for 303 Bratislava (0.52%), Tarassovi (0.30%), Ballum (0.19%) and Canicola (0.04%) at titer≥ 1:100. Seropositivity for 304 these Leptospira serovars was previously reported in sheep where they could represent the cause of 305 accidental infections [55]. Even if the presence of these serovar was reported in other studies, the reactivity 306 to all tested serovars detected in sheep during this investigation seems unusual. It could be probably 307 explained by the presence of wild boars, hares, hedgehogs, rodents (as mice and rats) and other wild 308 animals, which represent maintenance hosts for different serovars, in the areas where the sampling was 309 conducted [23,29,33,40,56-58].

For goat sera, only 2.75% (9/327) of samples resulted positive for Icterohaemorrhagiae at titer  $\ge$  1:100, among them 6 were positives at titer  $\ge$  1:400. Our data confirm that goats are not very susceptible to *Leptospira* infection, as reported in literature by other study, where seropositivity was described for serovar Hardjo [51,59,60], Icterohaemorrhagiae [60,61] and Poi [25].

314

#### 315 <u>4.6. Wild boar</u>

The total apparent prevalence in wild board increased from 2.39% (1995-2001) [9] to 8.56% (2002-2016). The highest prevalence was detected for serovars Tarassovi and Bratislava. Similarly to swine, wild boar is a maintenance host for serovars Pomona, Tarassovi and Bratislava [7]. For Tarassovi, 3.76% (18/479) and 319 0.63% (3/479) of sera scored positive at titer  $\geq$  1:100 and titer  $\geq$  1:400, respectively. This result disagrees 320 with other studies previously conducted in the same area of investigation and in Italy. In Tuscany, from 321 1995 to 2001 no positive sera for Tarassovi has been detected in wild board [9,14]. In Italy, recent surveys 322 indicate the same trend: no Tarassovi seropositivity or few positive samples (2 out of 1987 sera) were 323 found in wild boar [10,22,62]. However, high seroprevalence for this serovar has been reported in different 324 European Countries [63–65]. Wild boar could represent the reservoir host of this serovar, disappeared in 325 other domestic animal, and it could contribute to maintain Tarassovi strains in environment. Furthermore, 326 the Tarassovi high prevalence in investigated area could be connected to the import in Tuscany of wild boar 327 from East Europe Countries for hunting [34,63,64]. The second most representative serovar in wild boar 328 was Bratislava: 2.51% (12/479) and 0.21% (1/479) of positive sera have been detected at titer  $\geq$  1:100 and 329 titer ≥ 1:400, respectively. Bratislava is one of the most worldwide spread Leptospira [26], consequently, 330 this result was expected. Indeed, this serovar is frequently detected in wild boar in Italy [9,10,14,22,62] and 331 in Europe [63–67]. Unexpected data has been recorded concerning positivity to Pomona, considering that 332 wild boar is a potential reservoir, as reported by other studies carried out in Europe [66]. Only 0.84% 333 (4/479) and 0.21% (1/479) of sera resulted positive for this serovar at titer  $\ge 1:100$  and titer  $\ge 1:400$ , 334 respectively. These data were unexpected, but similar to those reported by other Authors in Italy 335 [10,22,62]. As concerns serovar Hardjo, 2.92% (14/479) and 1.46% (7/479) of sera scored positive at titer ≥ 336 1:100 and titer  $\geq$  1:400, respectively. This serovar is generally associated to cattle and sheep [7] and its 337 detection in wild boar seems singular. In Italy, no Hardjo seropositivity has been founded previously in wild 338 boar [10,22,62]. However, Hardjo seems to be the most prevalent serovar in East Europe in wild boar [66]. 339 For this reason, our finding could be related to the import of animals from East Europe for hunting 340 purposes. Nevertheless, in investigated area there are a lot of free-range farms of cattle and sheep and it is 341 plausible to assume that wild boar became infected after direct or indirect contact with these animals. 342 Moreover, half of positive sera showed an antibody titer higher than 1:400, suggestive of recent and 343 possible acute infection. For all these reasons it could not be excluded a possible involvement of wild boar 344 in epidemiology of Leptospira serovar Hardjo, as maintenance or incidental host. Furthermore, positivity to

other serovars was detected at titer ≥ 1:100:0.62% (3/479) of sera for serovar Icterohaemorrhagiae, 0.21%
(1/479) of sera for serovar Canicola, 0.21% (1/479) of sera for serovar Grippotyphosa and 0.21% (1/479) of
sera for serovar Ballum . These animals could easily come in contact with *Leptospira* due to their lifestyle
and positivity to many different serovars is sporadically, but constantly reported [10,22,63–67].

349

#### 350 <u>4.7. Other wild animals</u>

- 351 Excluding wild boar, seropositivity in wild animals was recorded only for hares, foxes and coypus. It is well
- documented that hare could be infected by different *Leptospira* serovars, especially Grippotyphosa
- 353 [22,33,56,68]. Only 1.85% (3/162) of hares sera scored positive in this survey, 0.62% (1/162) for Bratislava

354 (titer 1:200) and 1.23% (2/162) for Ballum (titer 1:100 and 1:200, respectively).

355 Fox was reported as incidental host for different serovars, such as Icterohaemorrhagiae, Ballum and

Bratislava [10,58,69]. Low number of fox sera examined in this investigation resulted positive for

357 *Leptospira*, in particular for serovar Icterohaemorrhagiae (1.06% - 1/94) and Bratislava (3.19% - 3/94). Only

1.06% (1/94) of sera resulted positive at titer  $\geq$  1:400, for serovar Bratislava.

359 For both these species, a low percentage of positive sera was detected compared to the other reported

surveys. This could suggest a weak involvement of these two animal species in *Leptospira* epidemiology in
 investigated area.

362 As regard coypus, some studies highlighted the circulation of the same serovars among these animals and

its possible role as reservoir host [18,70–72]. In accordance with these studies, an high percentage of sera,

364 32.86% (23/70), analyzed in this investigation scored positive to *Leptospira*. Positivity was recorded for

365 Grippotyphosa (20.00% - 14/70), Bratislava (11.43% - 8/70) and Icterohaemorrhagiae (1.43% - 1/70).

Furthermore, 4.29% (3/70) and 2.86% (2/70) of sera exhibited a titer ≥ 1:400 for serovars Bratislava and

367 Grippotyphosa, respectively; these data could suggest a recent infection, confirming the circulation of these

368 serovars in coypus. Positivity to Bratislava and Icterohaemorrhagiae was frequently reported, while

369 positivity to Grippotyphosa, which is considered an emerging serovar in Europe, could open new interesting

370 epidemiological scenarios.

- 371 No positive reactions were detected in sera from wild ruminants: roe deer, red deer, fallow deer and
- 372 mouflon. Our data confirm the marginal role that these animals have in the epidemiology of *Leptospira*

according to the low number of positivity reported in literature [9,10,17,73,74].

No positivity was detected in sera collected from other wild animals (wolves, rats, bear and ferret).

However, small number of samples were analyzed for these species and it is not possible to advance robust
epidemiological hypothesis.

377

378 <u>4.8. Human</u>

379 The total apparent prevalence observed in man slightly decreased from 5.60% (1995-2001) [9] to 4.26% 380 (2002-2016). Humans are incidental hosts for *Leptospira* and they could be infected by serovars maintained 381 by animals in a particular geographical region. In past years in Europe, human leptospirosis had a 382 fluctuating trend, but different confirmed cases were always registered. In particular in Italy about 40 383 human cases were annually recorded from 2008 to 2015 [75,76]. According to the present survey, 384 Icterohaemorrhagiae remained the most detected serovar from human sera. However, positivity to all 385 tested serovars, with exception of Tarassovi and Hardjo, was recorded. This finding could suggest a change 386 also in human leptospirosis related to a modification of *Leptospira* epidemiology in investigated area.

387

#### 388 5. Conclusion

389 Leptospirosis is probably the most widespread, (re-)emerging and prevalent zoonotic disease in the world. 390 However, due to the difficult to exactly diagnose the disease clinically and by laboratory test, sometime, it 391 could be not recognized and consequently severely neglected. For this reason, the true spread and increase 392 of leptospirosis remains probably unknown [12]. Considering that many domestic and wild mammals 393 represent natural carriers of pathogenic leptospires, acquire epidemiological information on animal 394 leptospirosis could be helpful for both human and breeding animal health. Despite that isolation and strains 395 characterization should have an highest diagnostic value, serology represent for some disease, such as 396 leptospirosis, the best instrument for epidemiological purpose. The results of this investigation provide

397	information on Leptospira epidemiology in a defined geographical area involving many different animal
398	species, maintenance end accidental hosts, and humans. Moreover, a long period of time was considered
399	and this allowed to put in evidence a fluctuation in <i>Leptospira</i> positivity recovery, as also suggested by
400	other Authors; this trend involved both the total percentage of positive animals registered year by year and
401	the serovars encountered. Our results seem to highlight an increase of Leptospira in North-Central Italy and
402	a change in serovars potentially involved in animal and human infection.
403	Several animals resulted infected by unusual Leptospira serovars and this finding could suggest a change in
404	host range for some serovars, that may promote the adaptation to new hosts. Constant serological
405	monitoring results essential to control the evolution of the dynamics of <i>Leptospira</i> epidemiology and it
406	could represent the basis to lead future investigations focused on specific animals and that must include
407	both serological that isolation or molecular techniques.
408	
409	
410	Conflict of interest statement
411	All authors declare no conflict of interests.
412	

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633

## **Figure**

637	Figure 1: Percentage of positive sera detected for L. interrogans and for each Leptospira serovars year by
638	year.

639 Note: Ic: Icterohaemorrhagiae; Ca: Canicola; Po: Pomona; Tar: Tarassovi; Gri: Grippotyphosa; Br: Bratislava; Ba: Ballum; Har: Har

			Tit	ter	
Species	Examined	≥1:100	%	≥1:400	%
Bovine	1328	173	13.03	42	3.16
Dog	1144	99	8.65	27	2.36
Goat	327	9	2.75	6	1.83
Horse	242	7	2.89	0	0.00
Wild boar	479	41	8.56	13	2.71
Hare	162	3	1.85	0	0.00
Coypus	70	16	22.86	4	5.71
Sheep	2682	84	3.13	38	1.42
Swine	1332	263	19.74	84	6.31
Fox	94	4	3.04	1	0.91
Human	329	10	4.26	3	1.06
Total	8488	709	8.35	218	2.57
	-	-		-	

**Table 1:** Number of positive sera to *Leptospira* at low ( $\geq$  1:100) and high titers ( $\geq$  1:400).

<b>Table 2:</b> Numbers of positive serological reactions detected for the different <i>Leptospira</i> serovars at low (≥
1:100) and high titers ( $\geq$ 1:400) in relation to animal species.

					Serc	ovar					
Animal species	Titer	lc	Ca	Ро	Gri	Tar	Br	Har	Ва	Total	Positive to more serovars
Bovine	≥ 100	21	4	55	48	3	3	94	4	232	44*
	≥ 400	4	1	3	5	1	1	30	2	47	3*
Dog	≥ 100	58	14	16	4	1	48	0	0	141	33*
	≥ 400	16	1	2	0	1	19	0	0	39	11*
Goat	≥ 100	9	0	0	0	0	0	0	0	9	0
	≥ 400	6	0	0	0	0	0	0	0	6	0
Horse	≥ 100	3	0	1	0	0	2	1	0	7	0
	≥ 400	0	0	0	0	0	0	0	0	0	0
Wild boar	≥ 100	3	1	4	1	18	12	14	1	54	11*
	≥ 400	1	0	1	0	3	1	7	1	14	1
Hare	≥ 100	0	0	0	0	0	1	0	2	3	0
	≥ 400	0	0	0	0	0	0	0	0	0	0
Coypus	≥ 100	1	0	0	14	0	8	0	0	23	7*
	≥ 400	0	0	0	2	0	3	0	0	5	3
Sheep	≥ 100	20	1	30	20	8	14	19	5	117	21*
	≥ 400	2	0	17	6	3	1	12	1	42	3*
Swine	≥ 100	2	6	160	0	2	131	1	0	302	40*
	≥ 400	0	0	45	0	1	41	0	0	87	3
Fox	≥ 100	1	0	0	0	0	3	0	0	4	0
	≥ 400	0	0	0	0	0	1	0	0	1	0
Human	≥ 100	7	2	1	2	0	2	0	1	15	2*
	≥ 400	2	0	0	1	0	0	0	0	3	0
Total	≥ 100	125	28	267	89	32	224	129	13	907	158
Note: Ic: Icterobaer	≥ 400	31	2	68	14	9	67	49	4	244	24*

Note: Ic: Icterohaemorrhagiae; Ca: Canicola; Po: Pomona; Tar: Tarassovi; Gri: Grippotyphosa; Br: Bratislava; Ba: Ballum; Har: Hardjo;

\*Some sera resulted positive to more than 2 different serovars

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#### Supplementary material

**Table S1:** Distribution of positive sera at titer  $\ge$  1:100 and  $\ge$ 1:400 in relation to animal species, *Leptospira* serovars and year of detection

Animal										Year								Total
species	Serovars	Titer	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	
Bovine	lc	≥ 100	0	0	0	0	0	0	0	0	3	0	0	0	18	0	0	21
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	4
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	4
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
	Ро	≥ 100	0	0	0	0	0	4	0	2	0	0	0	0	4	0	45	55
		≥ 400	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	2
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	3	0	0	3
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
	Gr	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	13	0	34	48
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	3	0	2	5
	Br	≥ 100	2	0	0	0	0	1	1	1	1	1	1	1	0	1	0	10
		≥ 400	0	0	0	0	0	1	1	1	1	1	1	1	0	0	0	7
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	4
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	2
	Har	≥ 100	0	0	0	0	0	0	0	2	0	0	0	0	4	88	0	94
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	2	27	0	29
Dog	lc	≥ 100	10	8	6	0	0	2	1	1	1	1	1	12	8	3	4	58
		≥ 400	5	1	5	0	0	0	0	1	0	0	0	2	0	2	0	16
	Ca	≥ 100	1	1	0	0	0	0	0	0	0	0	0	3	1	2	5	13
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1
	Ро	≥ 100	0	0	0	0	0	1	0	0	0	0	0	1	0	4	10	16
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	2
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
	Gr	≥ 100	0	0	0	0	0	0	0	0	0	0	0	2	1	1	0	4
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 100	6	9	6	2	2	1	0	1	2	0	1	6	3	4	4	47
		≥ 400	2	1	5	1	1	1	0	01	0	0	0	2	0	4	0	18
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Goat	lc	≥ 100	0	0	0	0	0	0	0	0	0	9	0	0	0	0	0	9

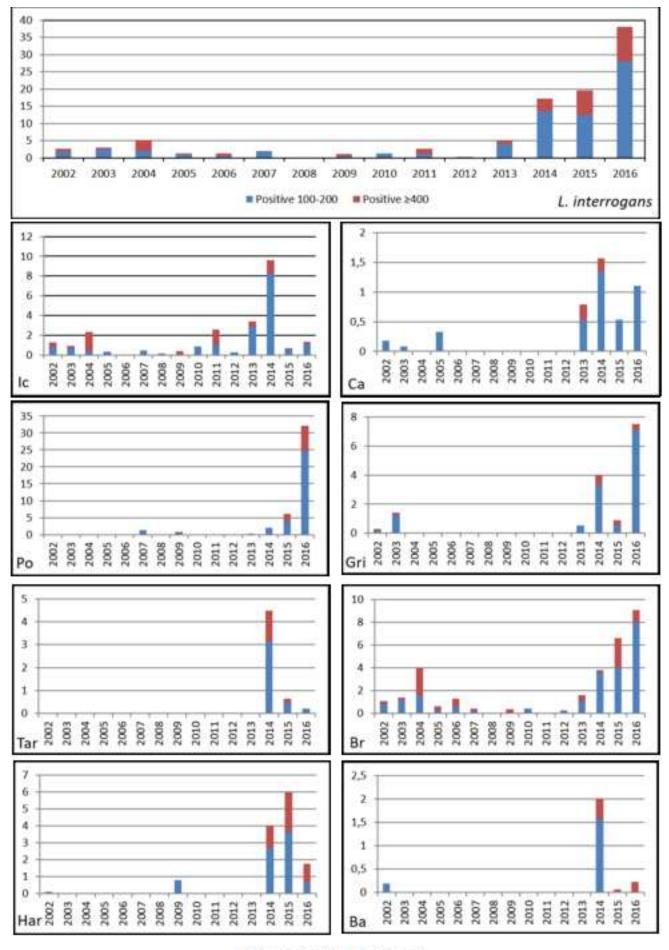
								-			-		-		-	-		
-	_	≥ 400	0	0	0	0	0	0	0	0	0	6	0	0	0	0	0	6
	Са	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Gr	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Horse	lc	≥ 100	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	3
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	141	≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Gr	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 100	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	2
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Da	≥ 100 ≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	lc	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	1	0	2	3
Wild boar		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 400 ≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	1	0	3	4
		≥ 100 ≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
	Tar	≥ 400 ≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	12	5	1	18
		≥ 100 ≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	3	0	0	3
	Gr	≥ 400 ≥ 100	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1
	5	≥ 100 ≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 400 ≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	2	5	5	12
		≥ 100 ≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	12
	Ва	≥ 400 ≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
	Dd	≥ 100 ≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
	llar																	
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	6	0	8	14

			•	<u> </u>				-	_	<u> </u>				-	_		-	-
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	2	0	5	7
Hare	lc	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Са	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Gr	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ba	≥ 100	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Coypus	lc	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Gr	≥ 100	0	14	0	0	0	0	0	0	0	0	0	0	0	0	0	14
		≥ 400	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	2
	Br	≥ 100	2	6	0	0	0	0	0	0	0	0	0	0	0	0	0	8
		≥ 400	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	3
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep	lc	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	12	7	0	20
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	2
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	1	0	0	0	0	0	0	0	0	0	0	4	1	23	29
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	17	17
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	5	3	0	8
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	21	1	0	22
	Gr	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	3	14	0	18
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0	6
	Br	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	11	2	0	14
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
			0	0	0	0	0	0	0	0	0	0	0	0	4	1	0	5

		> 400	0	0	0	0	0	0		0	0	0	0	0	0		0	
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
	Har	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	7	11	0	19
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	2	10	0	12
Swine	lc	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	2
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0	6
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	98	62	160
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	33	12	45
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	2
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
	Gr	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	99	32	131
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	38	33	71
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Fox	lc	≥ 100	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ca	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Gr	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Br	≥ 100	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	3
		≥ 400	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Human	lc	≥ 100	0	2	0	0	0	0	0	0	0	0	0	1	3	0	0	6
		≥ 400	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	2
	Ca	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	2
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ро	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Gr	≥ 100	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	2
		≥ 400	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
	Br	≥ 400 ≥ 100	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	2
	וט	- 100	U	Т	0	0	0	0	0	U	U	U	0	0	1	U	U	-

	1	1		1		1					r	-						r
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Ва	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Har	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Total	lc	≥ 100	14	11	6	1	0	2	1	1	4	10	1	13	43	12	6	125
		≥ 400	5	3	5	0	0	0	0	1	0	6	0	2	6	2	1	31
	Ca	≥ 100	2	1	0	1	0	0	0	0	0	0	0	3	7	9	5	28
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	2
	Ро	≥ 100	0	1	0	0	0	6	0	2	0	0	0	1	9	103	145	267
		≥ 400	0	0	0	0	0	0	0	1	0	0	0	0	0	34	33	68
	Tar	≥ 100	0	0	0	0	0	0	0	0	0	0	0	0	20	11	1	32
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	6	3	0	9
	Gr	≥ 100	3	17	0	0	0	0	0	0	0	0	0	2	18	15	34	89
		≥ 400	1	2	0	0	0	0	0	0	0	0	0	0	3	6	2	14
	Br	≥ 100	12	17	10	2	2	2	0	1	2	0	1	6	17	111	41	224
		≥ 400	3	3	6	1	1	1	0	1	0	0	0	2	1	43	5	67
	Ва	≥ 100	2	0	0	0	0	0	0	0	0	0	0	0	9	1	1	13
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	2	1	1	4
	Har	≥ 100	1	0	0	0	0	0	0	2	0	0	0	0	18	100	8	129
		≥ 400	0	0	0	0	0	0	0	0	0	0	0	0	6	38	5	49

Note: Ic: Icterohaemorrhagiae; Ca: Canicola; Po: Pomona; Tar: Tarassovi; Gri: Grippotyphosa; Br: Bratislava; Ba: Ballum; Har: Hardjo



■ Positive 100-200 ■ Positive ≥400

Bertelloni et al., *Epidemiology of Leptospirosis in North-Central Italy: fifteen years of serological data (2002-2016)*.

## Conflict of interest statement

All authors declare no conflict of interests.