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This is the final peer-reviewed author's accepted manuscript (postprint) of the following publication:

Published Version:

Magri A., Galuppi R., Fioravanti M., Caffara M. (2023). Survey on the presence of Leishmania sp. in peridomestic rodents from the Emilia-Romagna Region (North-Eastern Italy). VETERINARY RESEARCH COMMUNICATIONS, 47(1), 291-296 [10.1007/s11259-022-09925-4].

Availability: This version is available at: https://hdl.handle.net/11585/906128 since: 2024-01-29

Published:

DOI: http://doi.org/10.1007/s11259-022-09925-4

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which has been published in final form at:

https://doi.org/10.1007/s11259-022-09925-4

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- 1 Survey on the presence of *Leishmania* sp. in peridomestic rodents from the
- 2 Emilia-Romagna Region (North-Eastern Italy)
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12 Abstract

13 Leishmaniasis is a neglected vector-borne parasitic disease caused in Italy only by the 14 species Leishmania infantum of the Leishmania donovani complex, which is the causative agent 15 of the zoonotic visceral leishmaniasis (VL), and the sporadic cutaneous leishmaniasis (CL) 16 in humans, and of the canine leishmaniasis (CanL). The disease is considered endemic in southern, central, and insular Italian regions and recognizes phlebotomine sand flies as 17 18 vector and dogs as main reservoir. However, a specific north-eastern region, namely Emilia-19 Romagna, always showed a peculiar epidemiological situation when compared to the other 20 northern Italian regions and recent studies are indeed questioning the role of dog as main 21 reservoir of *L. infantum*. Due to their synanthropic relationship with humans, rodents have been tested for Leishmania spp. in several European countries. The aim of this study was to 22 23 assess the presence of Leishmania spp. in peridomestic rodents in the Emilia-Romagna 24 Region. The study was carried out on 136 peridomestic rodents collected by professional rodent control services: 47 brown rats (Rattus norvegicus), 39 black rats (Rattus rattus) and 50 25 mice (Mus musculus). Specimens of earlobe skin, spleen, liver and prescapular lymph nodes 26 27 were tested with a real-time PCR. Fifteen (11 %) rodents, tested positive for L. infantum. Positivity was obtained from different target organs; notably 33% of the rodents tested 28 positive in earlobe skin samples. These findings revealed the presence of Leishmania spp. in 29 30 peridomestic rodents of the Emilia-Romagna Region, also in two species never tested before in Italy, namely *R. norvegicus* and *M. musculus*. 31

32 Keywords: Leishmaniasis, Italy, Mus musculus, Rattus norvegicus, Rattus rattus

34 Background

Leishmaniasis is a neglected vector-borne parasitic disease endemic in southwestern 35 Europe. With reference to Italy, Leishmania infantum of the Leishmania donovani complex is 36 37 the only species responsible for visceral leishmaniasis (VL), for sporadic cutaneous 38 leishmaniasis (CL) in humans and for canine leishmaniasis (CanL) (Gramiccia and Gradoni 39 2005; Rugna et al. 2018). The parasite is transmitted by phlebotomine sand flies, and in Italy 40 dogs have long been recognized as the major reservoir in southern, central and insular 41 regions, where the disease is considered endemic. Among the northern Italian regions, 42 Emilia-Romagna has always had a different epidemiological scenario: CL has been widely 43 reported since 1934, and between 1950-1958 up to 2,670 cases were diagnosed in the 44 province of Forlì (Pampiglione 1975). In contrast, until the early 1970's, in this region VL appeared sporadically, with only 4 autochthonous cases observed, one in the province of 45 Bologna and 3 in the province of Forlì. In the same period and within the same area, no 46 47 ascertained autochthonous cases of CanL were reported (Pampiglione 1975). In 1971-1972, 48 in two municipalities of Bologna province located in a foothill area a dramatic outbreak of VL was reported, involving 60 patients with a lethality of 21.7% (Pampiglione 1975). Since 49 50 then, the geographic distribution of human and canine leishmaniasis has notably increased 51 and the disease spread even in other regions of northern Italy, where many autochthonous 52 cases of VL, CL and CanL have been reported (Gaspari et al. 2017; Mendoza-Roldan et al. 53 2020). This epidemiological change may be due to environmental issues, occurrence of 54 competent insect vectors and movement of infected dogs from endemic areas (Santi et al. 55 2014). However, such changes might not be sufficient to explain the recurrent VL and CL

foci recorded in Emilia-Romagna Region (Gaspari et al. 2017), especially considering that molecular studies carried out on strains isolated from autochthonous cases of VL are questioning the role of dogs as reservoirs of *L. infantum* in this region, as earlier suggested (Pampiglione 1975; Rugna et al. 2018).

60 The role of wildlife has long been recognized as crucial in the transmission and maintenance of zoonotic agents and several sylvatic species are known to be susceptible to leishmaniasis. 61 62 Considering their synanthropic relationship with humans and their abundance the role of rodents as possible leishmaniasis reservoirs has been questioned in different European 63 64 countries (Alcover at al. 2021). Several studies established the presence of L. infantum in 65 these hosts in Greece (Papadogiannakis et al. 2009; Tsakmakidis et al. 2017), Portugal (Helhazar et al. 2013) and Spain (Navea-Pérez et al. 2015; Galán-Puchades et al. 2019; 66 67 Martín-Sánchez et al. 2020).

In Italy, a study performed in Sicily detected *L. infantum* by PCR in 45% of black rats, even
if in this region the role of the dog as reservoir has been well established (Di Bella et al.
2003). However, a study performed in Montecristo Island (Tuscany), revealed the presence
of *L. infantum* in the 15.5% of rodents examined, even in the absence of domestic carnivores
(Zanet et al. 2014).

73 The aim of this survey was to assess the presence of *Leishmania* spp. in peridomestic rodents74 collected in the Emilia-Romagna Region, Italy.

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78 Materials and Methods

From June 2019 to June 2021, 136 peridomestic rodent carcasses were sampled during pest control programs from the provinces of Ferrara, Forlì-Cesena and Ravenna (Emilia Romagna) (figure 1): 47 brown rats, *Rattus norvegicus* (20 females and 27 males), 39 black rats, *R. rattus* (21 females and 18 males), 50 mice, *Mus musculus* (22 females and 28 males) were collected from the territory by professional rodent control services and stored at -20 °C before processing.

The entire carcass was examined; species, sex and age classes were identified by morphological and metrical evaluation (CDC). Necropsies and samples collection were performed with sterile surgical instruments and when possible, according to the state of the carcasses, 25 mg of tissue were collected from earlobe skin, prescapular lymph node and liver, and 10 mg from the spleen (Helhazar et al. 2013). Due to the corruption of the remains, lymph nodes were not collected from 16 subjects. Samples were placed in sterile 1.5 ml tubes and stored at -20 °C.

92 DNA extraction was performed with PureLink® Genomic DNA Mini Kit (Invitrogen, ThermoFisher Scientific) following the manufacturer's instructions. For DNA amplification 93 94 a real-time PCR protocol was performed targeting a 71-bp region of the minicircle 95 kinetoplast DNA primer Leish71Up (5'using pair CCAAACTTTTCTGGTCCTYCGGGTAG-3') Leish71Do (5'-96 and 97 TGAACGGGATTTCTGCACCCATTTTTC -3') (Tsakmakidis et al. 2017). Reactions were carried out in a total volume of 20 µL with 10 µL of PowerUP[™] SYBR[™] Green master mix 98 99 (2X), 0.3 µM of each primer and 2 µL of DNA. The amplification was performed in 100 StepOnePlus Real-Time PCR System (Applied Biosystems) and the thermal cycling profile was as follows: 95 °C for 5 min, followed by 40 cycles at 95 °C for 5 sec., 60 °C for 30 sec. At 101 the end of the amplification, a melting curve analysis was performed from 60 °C to 95 °C, 102 103 with a slope of 0.3 °C to monitor primer dimers of non-specific product formation. Each 104 sample was amplified in triplicate, the average temperature of melting (Tm) observed was 79.39 ± 0.15 °C and the average standard deviation observed in cts was 0.65. The standard 105 curve was created with serial dilution of L. infantum DNA ranging from 10,000 to 0.1 106 parasites per reaction. Each reaction was carried out by three replicates per dilution, in three 107 independent experiments. The ct value cut-off was settled at mean ct value of 39.3 which 108 109 corresponds to 1 parasite per mL of the original parasite suspension.

As a positive control the reference strain *L. infantum* MHOM/TN/80/IPT1, kindly provided
by the Unit of Clinical Microbiology, Regional Reference Centre for Microbiological
Emergencies (CRREM), St. Orsola-Malpighi University Hospital, Bologna, Italy, was used.
Confidence intervals were calculated by R Studio (RStudio Team 2020).

114 **Results and Discussion**

Out of 136 subjects examined, 15 (11 %; 95% CI=5.7-16.3) were positive for *Leishmania* spp. In particular, 10.6% (95% CI=1.8-19.4) of brown rats, 12.8% (95% CI=2.5-23.7) of black rats and 10% (95% CI=1.7-18.3) of mice (Table 1). Of the five positive mice, three tested positive in two target organs - spleen and earlobe skin or spleen and liver or spleen and lymph nodes - the remaining two subjects tested positive only in lymph nodes or liver, respectively. The geographical distribution of the positive subjects appears homogeneous between the sampled sites (figure 1). The present survey assessed the presence of *Leishmania* spp. in synanthropic rodents of the Emilia-Romagna Region. The conditions settled by the WHO (2010) for a species to be recognized as reservoir is the prevalence of infection > 20% and the availability of the parasite in blood and skin in sufficient amount to be ingested by a sand fly. In the Mediterranean area such conditions were globally assessed only for *M. musculus*, while *R. norvegicus* and *R. rattus* showed lower prevalence of infections (16.4% and 9.9%, respectively) (Alcover et al. 2021).

The prevalence values observed in the current study are below the average found in 129 130 Portugal or Spain (Barcelona) where the 33.3% of examined rodents (M. musculus and R. 131 norvegicus, Helhazar et al. 2013; R. norvegicus, Galán-Puchades et al. 2019) resulted positive, 132 or the one reported in Granada (Spain) in mice (88.9%) (Martín-Sánchez et al. 2020) or in 133 different rodent species (R. rattus, M. musculus and Apodemus sylvaticus) (27%) by Navea-Pérez et al. (2015), whilst it is higher than the prevalence observed in brown rats (5.9%) in 134 Greece (Papadogiannakis et al. 2009). Further studies performed in Greece by Tsakmakidis 135 136 et al. (2017) on spleen of R. norvegicus, R. rattus and M. musculus revealed a prevalence of 137 19.58% comparable to the one herein reported. The majority of the studies evaluated the presence of the parasite in more than one target organ including skin, liver, spleen and blood 138 139 (Helhazar et al. 2013; Martín-Sánchez et al. 2020; Navea-Pérez et al. 2015; Tsakmakidis et al. 2017) while few studies examined only the spleen as target organ (Galán-Puchades et al. 140 141 2019; Papadogiannakis et al. 2009). Testing more than one target tissue allow to increase the 142 possibility to detect Leishmania spp. as observed also in our study. Three M. musculus here 143 examined showed the presence of the parasite DNA in two different target organs (spleen + lymph nodes and spleen + liver). Although the spleen is traditionally recognized as *Leishmania* spp. target organ for PCR in different animal species (Papadogiannakis et al.
2009), our results showed the presence of *Leishmania* spp. in the earlobe skin samples from
33 % of the positive rodents pointing out that this tissue should be also considered. In fact,
wild animals are frequently collected in decomposition state and the putrefaction of the
target tissues, like visceral organs, may affect the integrity of the kinetoplast DNA (MũnozMadrid et al. 2013).

151 In Italy, the role of black rats in the transmission of *L. infantum* has long been investigated, starting from surveys performed in Tuscany in the 1980's (Pozio et al. 1985). Further studies 152 153 showed that *Phlebotomus perniciosus* and *P. perfiliewi* are attracted to *R. rattus* and that these sand fly species become infected when they feed on black rats experimentally infected with 154 155 L. infantum (Pozio et al. 1985). More recent study carried out in Calabria (Italy) by Di Bella et al. (2003) showed 45% positivity in the spleen of 22 R. rattus although in this region the 156 role of dogs as reservoirs has long been established. Zanet et al. (2014), reported 15.5% 157 158 prevalence in black rats examined in the Montecristo Island (Tuscany, Italy) where L. 159 infantum was recorded even in absence of domestic carnivore hosts. This value is similar to the one (12.8%) obtained in the same host in our study, that moreover provided also data 160 161 on *R. norvegicus* and *M. musculus* (10.6% and 10% respectively) species not previously tested for *L. infantum* in Italy. 162

Leishmaniasis in Emilia-Romagna has a peculiar epidemiological scenario compared to the
 other Northern Italian regions. Recently Rugna et al. (2018), by Multilocus Microsatellite
 Typing (MLMT) detected differences between *Leishmania* strains from men and sand flies to

the ones from dogs. The MLMT profiles showed all canine samples belonged to one group
genetically related to Mediterranean MON-1 strain and similar to the VL samples from other
Italian regions, while all but one VL Emilia-Romagna case, and the isolates from sand fly
fell into a different group. Therefore, in this region the co-circulation of two distinct groups
of *L. infantum* seems to occur, and the VL in humans could have different cycles involving *P. perfiliewi* as a vector (Rugna et al. 2018; Calzolari et al. 2019) and might include other
vertebrates, besides dogs, as reservoirs.

In two of the three provinces studied, Ravenna and Forlì-Cesena, foci of VL, usually located in hilly areas, were historically described. The rodent samples analyzed were collected in an area not higher than 50 m above sea level, where the density of phlebotomines is scant and, according to leishmaniasis regional control plan, in 2020 only CL cases have been reported (Santi et al. 2021). Further research should focus on studying which strains circulate in this area.

Also notable is the presence of a positive brown rat in the province of Ferrara, where autochthonous cases of leishmaniasis in both dogs and humans have never been recorded: the specimen was collected in a locality on the border between the provinces of Ferrara and Ravenna where the phlebotomine population is recorded as being moderate (Santi et al. 2021). This finding, considering the consistent increase in geographical distribution of the disease and its vector, will require further investigation.

L. infantum is a vector-borne parasite and in its epidemiology many mammal species are
involved, hence identifying which one may act as a reservoir in the Emilia-Romagna Region
is an ambitious task due to the presence of different environments i.e. hilly or flatlands and

different distribution of sylvatic and peridomestic animals, which may possibly be involved in the parasite cycle. Even if the presence of the parasite in mammalian hosts is crucial, in order to fully understand his meaning as main reservoir or epiphenomena it should be associated with studies on the blood preferences of the phlebotomine vector.

The total prevalence observed in the present study (11%), despite being lower to the one required from WHO (2010) to establish a role of reservoir is comparable to the Mediterranean's one. As reported in previous studies, this value is far from being trivial: considering their close relationship with humans, their ability to colonize new environments and their impact on human health, rodents should not be neglected for their potential role in the transmission of Leishmaniasis, especially in urban areas (Alcover et al. 2021).

Although these preliminary findings are not sufficient to prove the role of peridomestic
rodents as reservoirs of *L. infantum*, they nevertheless indicate the opportunity to further
investigate their possible role in the epidemiology of different strains of *L. infantum*circulating in the Emilia-Romagna Region.

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- 203 List of abbreviations
- 204 VL = Visceral Leishmaniasis
- 205 CanL = Canine Leishmaniasis
- 206 CL = Cutaneous Leishmaniasis

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- 208 Declarations:
- 209 -Ethics approval and consent to participate

- 210 No ethical approval is officially required since the rodents examined had been subjected to
- 211 pest control are considered pest species.
- 212 Consent for publication
- 213 Not applicable
- 214 Availability of data and materials
- 215 The datasets generated during and/or analyzed during the current study are available from
- the corresponding author on reasonable request
- 217 Competing interests
- 218 The authors declare that they have no competing interests
- 219 Funding
- 220 This study received no external funding
- 221 Authors' contributions
- 222 MF and RG conceived the study. AM performed field work. AM and MC performed
- 223 laboratory work and analyzed data. AM and MC wrote the first draft of the manuscript. MF
- and RG reviewed the manuscript. All authors read and approved the final manuscript
- 225 Acknowledgements

Emilia-Romagna Region.

- 226 Authors would like to thank social cooperative For.B and dr. Guglielmo Pampiglione for
- 227 kindly providing rodent carcasses from the provinces of Ravenna and Forlì-Cesena of
- 229

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ID	Specimen	Locality		Real-Time PCR			
ID			Earlobe Skin	Spleen	Liver	Lymph Node	
57	Mus musculus	Bizzuno (RA)	ct= 32.7 (87)	ct=29.68 (676)	Negative	NA	
59	Mus musculus	Bizzuno (RA)	Negative	ct= 30.77 (316)	ct= 31.97 (143)	NA	
67	Mus musculus	Bizzuno (RA)	Negative	Negative	Negative	ct= 33.61 (47)	
98	Mus musculus	S. Alberto (RA)	Negative	ct= 36.71 (5.8)	Negative	ct= 37.07 (4.5)	
111	Mus musculus	Bizzuno (RA))	Negative	Negative	ct= 35.9 (10)	Negative	
4	Rattus norvegicus	Ravenna (RA)	ct= 34.25 (30.9)	Negative	Negative	Negative	
86	Rattus norvegicus	Godo (RA)	Negative	Negative	ct= 36.47 (6.8)	Negative	
141	Rattus norvegicus	Ravenna (RA)	ct= 37.75 (2.9)	Negative	Negative	Negative	
175	Rattus norvegicus	Forlì (FC)	Negative	Negative	ct= 36.27 (7.8)	Negative	
178	Rattus norvegicus	Argenta (FE)	Negative	Negative	ct= 36.67 (5.8)	Negative	
37	Rattus rattus	Forlì (FC)	Negative	ct= 36.47 (6.8)	Negative	Negative	
60	Rattus rattus	San Pietro in Campiano (RA)	ct=36.86 (6.2)	Negative	Negative	Negative	
95	Rattus rattus	Montaletto di Cervia (RA)	Negative	ct= 37.44 (6.2)	Negative	Negative	
179	Rattus rattus	Montaletto di Cervia (RA)	Negative	Negative	Negative	ct= 36.63	
206	Rattus rattus	Longastrino (RA)	ct= 37.89 (2.6)	Negative	Negative	Negative	

Table 1: Real time PCR positive samples

Legend: Ct values are reported as mean ct of observed in different target organs with the estimated quantity of parasites/ml (mean

standard deviation observed ±0.65). Localities are as well reported with reference to the province: Ferrara (FE), Forlì-Cesena (FC) and Ravenna (RA).

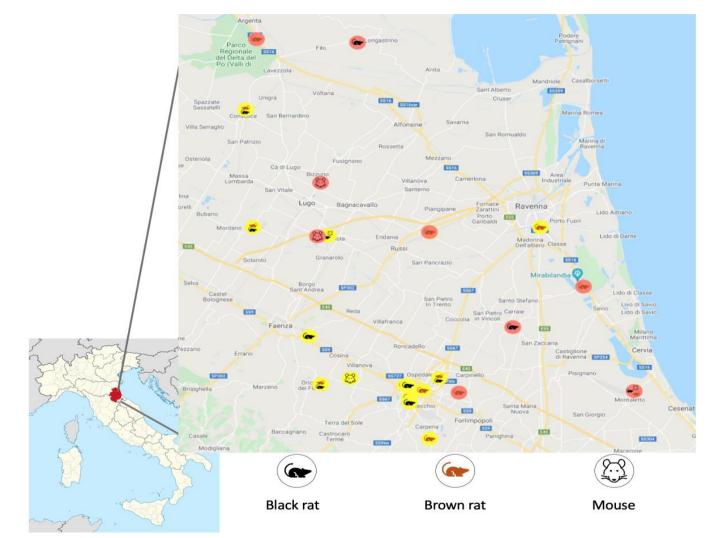


Figure 1. Map of the sampling area in the Emilia-Romagna Region. Dots are representative for sampling sites; red dots: at least one specimen positive, yellow dots: all the specimen negative.