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First detection of herpesvirus and frequency-prevalence of
mycoplasma infection in free-ranging Hermann's tortoises (*Testudo*
hermanni), and in potential pet vectors

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22 **Abstract**

23 Two types of pathogens cause highly contagious upper respiratory tract diseases (URTD) in
24 Chelonians: testudinid herpesviruses (TeHV) and a mycoplasma (*Mycoplasma agassizii*). In
25 captivity, these infections are frequent and can provoke outbreaks. Pet trade generates
26 international flow of tortoises, often without sanitary checking; individuals intentionally or
27 accidentally released in the wild may spread pathogens. A better understanding of the
28 transmission of infectious agents from captivity to wild tortoises is needed. Many exotic
29 individuals have been introduced in populations of the endangered western Hermann's
30 tortoise (*Testudo hermanni hermanni*), notably spur-thighed tortoises (*Testudo graeca*). We
31 assessed the presence of TeHV and mycoplasma in native western Hermann's tortoises and
32 in potential pet vectors in south-eastern France. Using a large sample (N=572 tortoises), this
33 study revealed, by PCR, the worrying presence of herpesvirus in 7 free-ranging individuals (3
34 sub-populations). Additionally, *Mycoplasma agassizii* was detected, by PCR, in 15 of the 18
35 populations sampled with a frequency ranging from 3.4% (1 of 29 tortoises) to 25% (3 of 12
36 tortoises). Exotic spur-thighed tortoises showed high frequency of mycoplasma infection in
37 captivity (18.2%) and in individuals (50%) found in native Hermann's tortoise sub-
38 populations, suggesting that this species could be a significant vector. The paucity of
39 information of TeHV on European tortoise' URTD in natural settings, especially in
40 combination with mycoplasma, prompts for further studies. Indeed, sick tortoises remain
41 concealed and may not be easily detected in the field. Our results indicate that both the
42 prevalence and health impact of URTD are high and should be scrutinized-screened in the
43 field.

Met opmerkingen [PG1]: The results indicate a good health for most infected tortoises?! delete?

Met opmerkingen [PG2]: .. as well as in captivity? (see last sentence of ms)

44
45 **Key words:** Emerging infectious diseases (EIDs), *Mycoplasma agassizii*, Testudinidae, upper
46 respiratory tract diseases (URTD), reptiles.

47

48 Introduction

49 Emerging infectious diseases (EIDs) represent a growing challenge for biodiversity
 50 conservation (Daszak et al., 2000; Deem et al., 2001). During the last decades, rapidly
 51 spreading diseases are suspected to have wreaked havoc worldwide among amphibians and
 52 reptiles (Daszak et al., 2000). In tortoises, *Testudinid herpesviruses* (TeHV) and *Mycoplasma*
 53 spp. are two dangerous pathogens for wild populations (Origgi, 2012; Marenzoni et al.,
 54 2018). Both are highly contagious and are involved in the Upper Respiratory Tract Disease
 55 Syndromes (URTD); infection ~~to~~with these pathogens they can provoke disease/illness
 56 entailing with high morbidity and mortality (Brown et al., 1994; Goessling et al., 2019). For
 57 example, in the 80's, mycoplasma epizooties were responsible of multiple collapses of
 58 desert tortoise populations in North America (Jacobson et al., 1991; Brown et al., 1994).
 59 Currently, several tortoise species such as Gopher tortoise (*Gopherus agassizii*), or captive
 60 spur-thighed tortoise (*Testudo graeca*) are impacted by URTD (Marschang and Schneider,
 61 2007; Weitzman et al. 2017). Monitoring the health status of free-ranging chelonians, with a
 62 focus on two major agents of/causing URTD, is thus a conservation priority.

63 On the other hand, global trade of reptiles is flourishing; tens of thousands of individuals
 64 from an increasing number of species are displaced among continents under minimal (or
 65 non-existent for illegal trade) sanitary monitoring (Auliya et al., 2016). The resulting flows of
 66 individuals open major routes for the expansion of EIDs (DiGeronimo et al., 2019). Pet
 67 tortoises host both TeHV and *Mycoplasma* spp. (Martínez-Silvestre et al., 1999; Sandmeier
 68 et al., 2009; Lecis et al., 2011; Salinas et al., 2011; Origgi, 2012). The primary route of
 69 transmission of Herpesvirus herpesvirus and mycoplasma is believed to be horizontal via
 70 contact between individuals (DiGeronimo et al., 2019). Interspecific transmission has been
 71 demonstrated (Origgi et al., 2004; Soares et al., 2004; Salinas et al., 2011). Tortoise species

Met opmerkingen [PG3]: More recent reference ? ("During the last decades...")

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72 are often mixed up in cages or enclosures. Overall, very high prevalence of infection in
73 captivity is expected (Kolesnik et al., 2017). Unfortunately, captive individuals are frequently
74 released into the wild, intentionally or accidentally, [where](#) they can settle in novel habitats
75 while carrying pathogens that may threaten native populations of tortoises (Sandmeier et
76 al., 2009; Jacobson et al. 2014; Whitfield et al., 2018). This process opens highways for
77 pathogens. The scarcity of investigations in natural populations, except in USA, means that
78 possible role of pet tortoises as agents of contamination is not quantified in Europe, as in
79 most parts of the world (Jacobson and Berry, 2012; Kane et al., 2017; Orton et al., 2020).

80 Fragile inbred populations are particularly at risk. Reduced phenotypic diversity and
81 genetic depression often hinder physiological and demographic resistances to diseases
82 (Frankham et al. 2002; Spielman et al., 2004). Many species are subjected to all the
83 threats above, including the not yet evaluated risk of simultaneous contamination by TeHV
84 and mycoplasma. The Hermann's tortoise provides a typical example of such a situation.
85 Two Hermann tortoise subspecies are currently recognized (Pérez et al., 2013): the Western
86 Hermann's tortoise (WHT, *Testudo hermanni hermanni*) that occurs west of the Po Valley in
87 Italy (e.g. Italian Peninsula, Sardinia, Corsica, southeastern France, northeastern Spain) and
88 the Eastern Hermann's tortoise *T. h. boettgeri* (EHT) found in Mediterranean regions of the
89 Balkan Peninsula and in small islands spread in the eastern Mediterranean sea. The two
90 subspecies come into contact in north eastern Italy where they possibly hybridize naturally
91 (Pérez et al., 2013). The WTH is severely threatened by habitat loss and fragmentation,
92 frequent fires, illegal harvesting and predation by feral animals. As a result, continental
93 populations drastically decreased (Livoreil 2009). Previously abundant in continental
94 southeastern France, relict isolated WHT sub-populations persist in the Massif des Maures
95 (Var district, 83) and in adjacent plains (Livoreil, 2009; Bertolero et al., 2011). Delayed

96 maturity, low fecundity and low population turn over mean that it is particularly sensitive to
97 a decrease of adult survival (Bertolero et al., 2011).

98 Intensive and long-lasting legal and illegal pet trades (CITES 2014) provoked substantial
99 introgression of EHT and of other tortoise species (e.g. spur-thighed tortoise) inside the
100 natural repartition area of WTH (Martínez-Silvestre et al., 2001), while various species
101 originating from other continents are sporadically found in the wild (unpublished data).
102 These exotic tortoises easily breed in captivity [and](#) are frequently owned as pets; they occur
103 in many properties spread across the entire (remaining) distribution range of the native
104 WHT. A considerable pool of captive individuals from various uncontrolled origins strongly
105 enhances the likelihood of contacts with free-ranging native WHT.

106 Cross-species transmissions have been documented both for TeHV and mycoplasma;
107 experiments demonstrated that these pathogens can infect Hermann and spur-thighed
108 tortoises (Origgi et al., 2004; Soares et al., 2004; Salinas et al., 2011). In Europe, TeHV-1 and
109 -3 affect most of the Testudinidae species raised in captivity (Origgi, 2012; Marschang and
110 Schneider, 2007). More generally, approximately 48% of individuals belonging to various
111 terrestrial and aquatic pet chelonians were positive for herpesvirus or mycoplasma while a
112 positive correlation was observed between the two pathogen detection frequencies
113 (Kolesnik et al., 2017). Like TeHV, mycoplasma was detected in tortoises kept in captivity
114 and in outdoor enclosures in Europe (Lecis et al., 2011; Salinas et al., 2011).

115 Mortality due to TeHV and mycoplasma is well documented in wild tortoises in USA
116 (Jacobson et al., 2012, 2014), but only in captivity in European tortoises (Soares, 2004;
117 Kolesnik, et al., 2017). In the latter, mortality rate caused by TeHV is higher in Hermann's
118 tortoise compared to other species, suggesting a recent and more deleterious contact
119 between the host and the pathogen (Soares et al., 2004). Possible occurrence and
120 consequences of co-infection by TeHV and mycoplasma are not documented, at least in the

121 Hermann's tortoise. Both frequency and possible severity of TeHV and mycoplasma
122 infections remain unexplored in native populations of tortoises in Europe. Overall, possible
123 impact of worrying EIDs has not been assessed in natural setting in the Mediterranean basin
124 that hosts many endemic tortoises (Gracià et al., 2020). This issue is urgent because
125 exogenous tortoises are frequently observed in the wild where they may carry new
126 pathogens (Lecis et al., 2011; Hidalgo-Vila et al., 2020).

127 ~~___ Infection to TeHV and mycoplasma does not necessary induce disease which can also~~
128 ~~be triggered by other etiological agents.~~ Accurate monitoring of the frequency of infections
129 ~~with these thes main agents pathogens TeHV and mycoplasma infections~~ and information
130 about the prevalence of URTD in the remaining populations of WHT is thus needed. ~~_ This,~~
131 notably because these pathogens-infectious agents can induce deleterious chronic diseases
132 that are not easily diagnosed in long-lived organisms (Sandmeier et al., 2013). ~~)-_ This study~~
133 reports results from the first comprehensive survey across the distribution range of the
134 WHT in continental France.

135

136 **Material and methods**

137 *Tortoise sampling*

138 From 2012 to 2016, 18 sites were monitored covering most of the distribution area of the
139 WTH subspecies in continental France (besides these surveys, individuals were also
140 opportunistically sampled throughout the distribution area) (Figure 1; Livoreil, 2009). Free-
141 ranging tortoises are cryptic; thus, in addition to visual searching, trained dogs were used
142 (Ballouard et al., 2019). Surveys were performed during the activity season of the species
143 (from March to October), mostly in spring (111 searching days in spring, 34 in summer and
144 15 in autumn). All tortoises sighted were captured: 457 free-ranging individuals (421 WHT
145 and 36 exotic specimens) were sampled (25.5 tortoises per site on average). In addition, 95

146 captive (pet) tortoises were sampled in 21 different properties in surrounding areas (5.2
147 individuals per property on average). Finally, 20 vagrant isolated individuals found in urban,
148 or peri-urban areas, were also tested; likely they were pets intentionally released or that
149 escaped from gardens. Most tortoises were adult (96%) and sex ratio was balanced (281
150 females, 267 males, 24 immatures). Overall, with respect to sampling context, we obtained
151 three categories of individuals: a) free-ranging, b) captive (pet), and c) vagrant (N total=572).

152 Each [sampling](#) category contained both native and exotic tortoises.

153 Individuals were assigned to species (e.g. *T. graeca* vs. *T. hermanni*) or subspecies (WHT
154 vs. EHT) according to their morphological characteristics. *Testudo* species are easily
155 distinguished (Bertolero et al. 2011), subspecies not. The following criteria were used to
156 discriminate WHT from EHT: yellow subocular scales, black continuous plastral bands,
157 narrow vertebral scute, supracaudal scute divided, long corneous tip of the tail, corneous
158 tubercles on the inner side of the thigh, ratio of pectoral vs. femoral seams (Bertolero et al.,
159 2011; Soler et al., 2012). Hybrids WHT x EHT displayed various combinations of phenotypic
160 characters and could not be identified with certainty (especially F2, unpublished genetic
161 results). Easily identifiable hybrids were brought to the Soptom rescue center.

162 Each tortoise was measured by straight carapace length (SCL), sexed when possible (small
163 immatures cannot be easily sexed), and weighted to the nearest gram. Individuals larger
164 than 100 mm in SCL were considered adult. Following blood, oral and nasal epithelium
165 sampling, individuals were subjected to clinical inspection (see below), and then they were
166 released at the place of capture, generally within 30 min. To ensure that researchers did not
167 spread pathogens and did not contaminate samples, they cleaned their hands and clothes
168 using Vircon spray 1% (Bayer®); equipment was cleaned with alcohol. Samples were stored
169 using one box per site.

170 Most individuals examined were free-ranging WHT (N=421; [table-Table1](#)). We identified
171 11 EHT: 2 free-ranging individuals introduced into WHT populations, 8 captive (pets), and
172 one vagrant. Thirty nine WHT x EHT easily identifiable hybrids were observed: 24 free-
173 ranging, 9 captive and 6 vagrants. Twenty four spur-thighed tortoises were examined: 10 of
174 them were free-ranging and thus have been introduced into WTH populations, 12 were
175 found in private properties (pets), and 2 were vagrant. Finally, 1 captive marginated tortoise
176 (*Testudo marginata*) was sampled.

177

178 *Tissue sampling*

179 Blood (0.4 - 0.7 ml) was collected from the subcarapacial plexus (Hernandez-Divers *et al.*
180 2002) using 1 ml syringes (Injekt-F – B Braun) and sterile needles (26G to 27G, Terumo
181 Neolus, adjusted to the size of the animal). Subcarapacial plexus delivers various mixtures of
182 blood and lymph, especially using needles larger than 27G (Bonnet *et al.* 2016); we did not
183 notice such mixture during sampling. Blood was immediately placed in Sodium or Lithium
184 heparin. Samples were gently homogenized and stored (max 4 hours) in ice-cooled
185 containers until centrifugation (1500 rpm for 5 min). Plasma was stored at -25°C until
186 analyses. Aliquots were distributed in two tubes: 50 µl for microbiological analysis and the
187 rest for biochemical analyses.

188 We sampled oral and nasal epitheliums and mucus. We injected ~0.5 ml of sterile saline
189 (0.9% sodium chloride, Lavoisier) to flush the nasal cavity. The resulting fluid was collected
190 with a syringe (0.1 ml) in each nostril and immediately stored in a 0.5 ml sterile conical tube.
191 Oral samples were collected with a brush (Cervibrush + LBC, Endocervical sampler, CellPath)
192 inserted inside the oral cavity: choana and mucosal surfaces of the tongue and of the beak
193 were targeted. Brushes were stored individually in a 0.5 ml sterile conical tube containing

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194 0.3 ml of sodium chloride (0.9%) to avoid desiccation of the mucus. All samples were placed
195 in ice-cooled containers in the field and stored at -25°C until analyses.

196 All samples were shipped frozen to Staaliches VetUAmt laboratory in Detmold, Germany,
197 for analyses.

198

199 *Pathogen screening*

200 The presence of Testudinid herpesvirus (TeHV-~~sero~~types 1 and 3) was assessed using both
201 polymerase chain reaction (PCR) ([see method in Teike et al., 2000](#)) and induced antibody
202 responses by serum neutralization test (SN) (Soares et al., 2004; Salinas et al., 2011; Origi,
203 2012). ~~PCR test can be applied to TeHV-1 and TeHV-3 in any species of tortoise (Salinas et~~
204 ~~al. 2011). A PCR test was considered positive for TeHV when DNA of pathogens was~~
205 ~~detected in oral mucus. However, the mucus of individuals that do not present clinical signs~~
206 ~~is generally less rich in viral DNA compared to the mucus of individuals displaying clinical~~
207 ~~signs (Origi, 2012). In addition, false negatives have been observed, likely due to sampling~~
208 ~~methodology difficulties or to oscillating elimination of some viruses such as [Herpesviruses](#)~~
209 ~~[herpesviruses](#) (Marschang 2019). Thus, we used SN testing as a complementary method to~~
210 ~~detect the presence of TeHV types 1 and 3 circulating antibodies on plasma aliquots (Origi,~~
211 ~~2001, 2012).~~

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212 The presence of *Mycoplasma agassizii* was assessed using PCR ([see method in Brown et](#)
213 [al. 1999](#)), no serological test being available in Europe during the study (2012-2016). Thus,
214 PCR were used to detect active infection by mycoplasma and TeHV (Soares et al., 2004).

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215 ~~Many tortoises were tested for both pathogens. Table 2 provides the numbers of tests~~
216 ~~performed on the different categories and taxa of tortoises.~~

217

218 *Clinical inspection*

219 All tortoises were visually inspected for clinical symptoms of URTD disease that are usually
220 associated with mycoplasma and herpesvirus infections (Jacobson et al., 2014): keratitis,
221 conjunctivitis, ocular and palpebral oedema, and nasal discharges (mucopurulent oculonasal
222 discharge), necrotic spots on the oral mucosa and on the tongue, rhinitis, lethargy and low
223 body condition (Brown et al., 2002; Berry and Christopher, 2001; Sandmeier et al., 2009).

224

225 Results

226 Many tortoises were tested for both pathogens. Table 2 provides the numbers of tests
227 performed on the different sampling categories (e.g., free-ranging, captive) and taxa of
228 tortoise species.

229 PCR testing revealed that seven free-ranging WHT (six adult females and one adult male)
230 were TeHV positive (2.8 %, n=7/253), and that 3 sub-populations (i.e. sites) were concerned
231 (Tables 2 and 3). SN tests for TeHV were all negative (table-Table2).

232 A total of 52 individuals (species and subspecies pooled) were positive for mycoplasma
233 DNA (prevalence Pr=9.2%) (table-Table2). ~~Most were free-ranging individuals (n=39, 75%)~~
234 ~~and most populations were infected (15 of 18) leading to a mean prevalence of 8.7.5%~~
235 ~~(range: 3.4% to 25%; Table 3). The numberproportion of individuals infected wasere not~~
236 ~~significantlyinfluenced different according theirby sampling categories situation ($\chi^2= 1.56;$~~
237 ~~df=2; p=0.45). We found a positivesignificant effect of the species on the observed~~
238 ~~prevalence of mycoplasma infection observed ($\chi^2= 20.41;$ df=3; p<0.001), ~~with the spur-~~
239 ~~thighed tortoises havingdisplaying the most highest rate of prevalence (two subspecies~~
240 ~~added,) (n=7, 13%, Pr=30%). ~~This species difference wasere also observed a~~Among free-~~
241 ~~ranging tortoises, ($\chi^2= 15.76;$ df=3; p<0.01), ~~b=most positive cases concerned WHT (n=34,~~
242 ~~87%; Pr=8.7%) followed by spur-thighed tortoises (two subspecies added) (n=5, 13%,~~
243 ~~Pr=50%). ~~But not among the captive All species pooled, 9 captive individuals were positive~~~~~~~~

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244 ($Pr=9.9\%$) and were sampled in five private properties (among 21). They belonged to almost
245 all species and subspecies found in the study area: 5 WHT ($Pr=7.7\%$), 2 spur-thighed
246 tortoises ($Pr=18.2\%$), one *Testudo marginata* ($Pr=100\%$), and one hybrid ($Pr=16.6\%$). ($\chi^2=$
247 4.61 ; $df=3$; $p=0.20$) or vagrant population tortoises ($\chi^2=2.44$; $df=1$; $p=0.11$). Among vagrants,
248 4 WHT were positives ($Pr=36.3\%$). Focusing on WHT (wild, captive and vagrant pooled)
249 tested positive for mycoplasma, there is was no statistical significant difference between of
250 the prevalence between the sexes (of 23 were female's (56.1% ; $Pr=9.67\%$, $23/238$), 18
251 were and male's (43.9% ; $Pr=7.9\%$, $18/227$); ($\chi^2=0.25$; $df=1$; $p=0.62$), and two individuals
252 were not sexed. No individual was tested positive for both pathogens.

253 Considering the whole sample, 28 individuals showed clinical symptoms of URTD; 4
254 displayed palpebral oedema, 1 with sunken eyes, 8 strong nasal discharges, and 14
255 abnormal mucus color (black, white or yellow), 1 exhibited abnormal mucus color + nasal
256 discharge. Among these 28 animals, 3 were tested positive for mycoplasma (2 with nasal
257 discharges, one with palpebral oedema) and none for TeHV. The general condition of the
258 tortoises was apparently normal otherwise and none exhibited ocular discharge.

259

260 Discussion

261 This study reveals the first cases of herpesvirus infection contamination and considerably
262 extends knowledge of mycoplasma infection occurrence (only one case previously known;
263 Mathes et al., 2001) in tortoises tested in natural populations in Europe (3 sub-populations
264 for TeHV, 15 sub-populations for mycoplasma). These issues come sharply into focus when
265 result obtained in captive and vagrant tortoises, native and exotic, are considered.

266 In this study, the high number of exotic tortoises observed vagrant (e.g., isolated
267 individuals walking nearby private properties) or settled in WHT natural populations
268 illustrate that uncontrolled introduction of exotic pet tortoises is an ongoing process.

Met opmerkingen [PG4]: Refer also to Table 3 in this paragraph, for difference between the sampling locations?

Met opmerkingen [PG5]: In contrast to the herpes infection, which occurred mainly in females.

269 Chelonian pet trades are intensifying worldwide (Stanford et al., 2020), including European
270 tortoises with massive exports of EHT and spur-thighed tortoises from Turkey, Balkan
271 countries or North Africa to France, Spain and Italy (Ljubisavljević et al., 2011). Our results
272 reveal substantial infection rates by mycoplasma both in captive and vagrant individuals of
273 different tortoise species sampled in the distribution range of the native species (WHT). This
274 situation poses the problem of possible ~~contamination-transmission~~ from captive toward
275 free-ranging tortoises, and perhaps conversely when free ranging WHT are illegally collected
276 and brought into captivity.

277 TeHV was not detected in captivity. The prevalence of mycoplasma infection was
278 relatively similar in free-ranging and captive tortoises (Pr=8.7% and Pr=9.9% respectively);
279 ~~TeHV was not detected in captivity.~~ Among pet tortoises, however, infection rate was higher
280 in exotic ~~species~~ and hybrid ~~s species~~ (22.2%, 4/18) compared to captive WTH (7.7%, 5/65).
281 In addition, high levels of infection were observed in exotic spur-thighed tortoises, both in
282 captivity (18.2%, 2/11) and among individuals introduced in the field (44%, 4/9) compared
283 to those observed in free-ranging WHT populations (~~138.57%~~, ~~3443/318391~~). This
284 disequilibrium between exotic and native tortoise's infection levels suggests that a common
285 exotic pet species may represent an important reservoir of mycoplasma. Wall-less
286 mycoplasma were thought to be highly fragile; but the ability of some species to form
287 resistant biofilms in vitro prompted a reconsideration of their ability to survive in the
288 environment (McAuliffe et al., 2006). If *M. agassizii* would be able to form resistant biofilms
289 in the environment, this may contribute to its persistent circulation even under such
290 ~~environmental resistance may contribute to persistent circulation of mycoplasma even~~
291 ~~under~~ low tortoise's population densities. Although preliminary, these results reinforce the
292 notion that pet tortoises, and thus international trade of chelonians, may represent a health
293 threat to native tortoises.

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294 Finding that free-ranging tortoises can be infected by TeHV and mycoplasma in Europe,
295 although worrying, is not fully surprising. Negative SN tests for TeHV despite PCR positive
296 tests suggest possible reactivation or recent contact with the pathogen that has not yet
297 triggered immunity. Herpesvirus is a virus with a low environmental persistence capacity;
298 transmission requires direct physical contact between tortoises (Marschang 2019).
299 Numerous assessments performed in captive tortoises maintained in cages or in outdoor
300 enclosures (inaccurately categorized as “wild” or “free-living” individuals by various authors,
301 leading to misleading conclusions) showed that many individuals belonging to different
302 lineages were infected by various pathogens involved in URTD. For example, herpesvirus
303 was detected by PCR in 16.3% of individuals from different tortoise species maintained in
304 captivity in Belgium; a country voided of native chelonian however (Martel et al., 2009).
305 Kolesnik et al. (2017) tested by PCR 1,015 captive tortoises and terrapins originating from
306 different continents and kept captive (e.g. in cage, enclosure, park, garden) in different
307 European countries; ~~more than~~ 42.1% were infected by mycoplasma and 8.0% were
308 infected by herpesviruses. In Turkey, among 272 spur-thighed tortoises caught in the wild
309 and kept in captivity (sometimes during years) to supply a breeding program designed to
310 reinforce native populations, seroprevalences (SN test) were as follow: 37% positive for
311 herpesvirus serotype-I, 5.5% for serotype-II, and 5.2% ~~5.2%~~ for picorna-like ‘X’ virus and
312 4.9% for reovirus ~~(X and reovirus)~~ (Marschang and Schneider, 2007). Broadly similar PCR
313 result of 36.7% was obtained with mycoplasma (Lecis et al., 2011). ~~More generally, high~~
314 ~~infection prevalence for a wide range of pathogens has been observed in captive~~
315 ~~chelonians, notably in individuals intercepted during illegal trade (Brianti et al., 2010).~~

316 Herpesvirus infections have been documented more than 40 years ago in American and
317 in European chelonians (Origgi, 2012). However, previous assessments performed two
318 decades ago in tortoises actually sampled in the field in France and Morocco (*Testudo*

319 *hermanni*, *T. graeca*) failed to detect herpesvirus (Mathes et al., 2001; [Mathes pers. com.](#)). A
320 situation that contrasts with well-documented infections in free-ranging chelonians in North
321 America (~~Jacobson, 1994~~; Berish et al., 2000; Jacobson et al., 2012; Kane et al., 2017;
322 Weitzman et al., 2017; Lindemann et al., 2019; Orton et al., 2020).
323 Further studies are needed to determine to what extent the presence of URTD pathogens
324 results from recent transmissions from pet to wild tortoises in Europe, *versus* insufficient
325 investigations in natural settings. Unfortunately, monitoring infection prevalence and
326 severity is difficult in the field. Underestimations are likely because infected animals that
327 may die are quickly removed prior to sampling. Further, weakened sick tortoises may well
328 remain concealed and may not be easily detected in the field.

Met opmerkingen [PG6]: Could this explain the absence of co-infected individuals in your results?

330 ***Infection risks and diseases***

331 Lack of severe clinical signs should be treated with caution. Silent TeHV and *Mycoplasma*
332 infections have been reported in tortoises (Orrigi, 2012; Withfield et al., 2018). Multiple
333 strains of pathogens circulate worldwide (Salinas et al., 2011; Jacobson et al., 2014; Kolesnik
334 et al., 2017), they continuously evolve while pathology and coinfections risks increase with
335 pathogen diversity (Kari et al., 2008). Thus, constant introductions of infected pets into
336 natural populations represent a threat through the emergence of infectious diseases,
337 especially if exotic tortoises tolerate and thus carry pathogens that can cause outbreaks in
338 less-tolerant native species (~~Berish et al., 2010~~; Jacobson et al., 2014; Whitfield et al., 2018;
339 DiGeronimo, 2019; Goessling et al., 2019). Environmental factors, like seasonal variations of
340 immunity and physiological stress can perturb equilibriums with pathogens, tipping the
341 balance toward diseases; especially in case of multiple infections (Sandmeier et al., 2013;
342 Goessling et al., 2016).

343 In European tortoises, the presence of herpesvirus and the high prevalence of
344 mycoplasma in natural populations were only suspected prior to the current study. But the
345 paucity of information regarding possible health impact of TeHV, in combination with
346 mycoplasma, along with the ~~spectre~~-spectrum of emerging serious diseases caused by
347 picornavirus in captive Hermann's tortoises in Spain for example (Martinez-Silvestre et al.,
348 2020), prompts for further studies.

349

350 ***Perspectives and recommendations***

351 French continental populations of Hermann's tortoise represent relicts of the distribution of
352 a previously widespread and abundant species. It is crucial to not add infectious burden to
353 the multitude of existing threats (e.g., habitat fragmentation, sprawling urbanization,
354 frequent fires, illegal harvesting, wild-boars, dogs). Both the prevalence and demographic
355 impact of URDI in the field should be carefully monitored. Pet owners should be informed
356 that reproducing tortoises in captivity and releasing individuals in the field may threaten
357 wild populations. Exotic tortoises should be removed from native populations. Long-term
358 mark-recapture surveys should be coupled with health checking, including monitoring of
359 most deleterious strains of TeHV (Gandar et al., 2015). Sanitary protocols are needed to
360 handle free-ranging tortoises; nasal and oral secretions, feces, sperm and even urine can be
361 contaminating (Origgi et al., 2012). Similarly, strict protocols are needed during conservation
362 translocations; notably because individuals often travel long distances before settling (Pille
363 et al., 2018), while stress associated with release may promote virus reactivation from
364 latent infection (Griffith, 1993; Martel et al., 2009; Jacobson et al., 2012). Furthermore,
365 captive individuals carrying a wide range of pathogens and parasites represent additional
366 risks to URDI transmission and should be treated specifically (Ahne 1993; Chávarri et al.
367 2012).

368

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378

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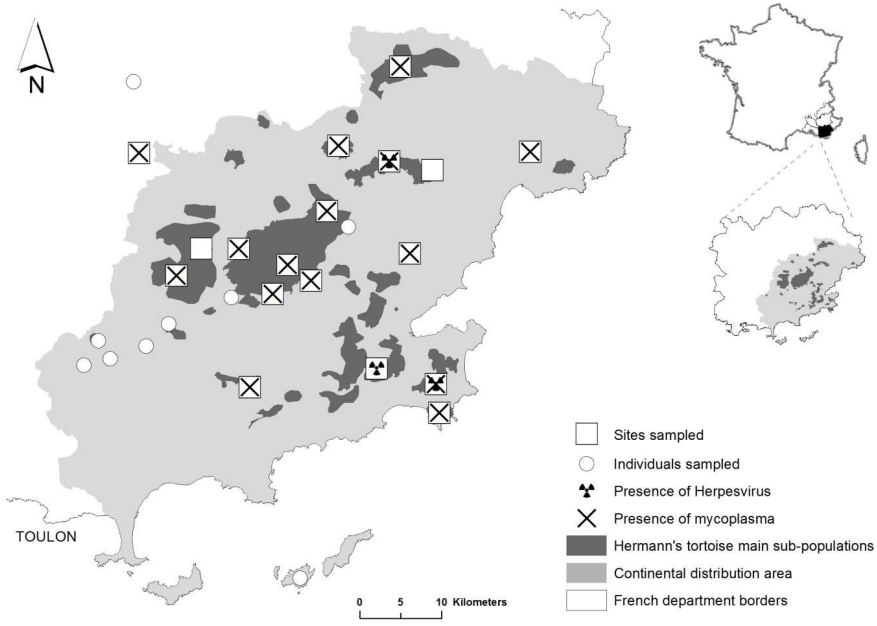
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591

592 **Figure 1:** Sampling sites inside the French continental distribution range of WHT.



593
594

595 **Table 1:** Numbers of individual tortoises sampled in function of species and subspecies
 596 (taxon), and in function of the situation where the animals were found (free-ranging, captive
 597 or vagrant, see text for details). *Testudo hermanni hermanni* is coded WHT, *T.h. boettgeri*
 598 EHT. WHT x EHT stands for hybrids. Different subspecies of the spur-thighed tortoises (*T.*
 599 *graeca*) are not distinguished.
 600

	Free-ranging	Captive (pet)	Vagrant	TOTAL
<i>T. h. hermanni</i>	421	65	11	497
<i>T. h. boettgeri</i>	2	8	1	11
<i>T. g. iberica</i>	1	-	-	1
<i>T. g. sp</i>	9	12	2	23
<i>T. marginata</i>	-	1	-	1
WHT x EHT	24	9	6	39
TOTAL	457	95	20	572

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603

604 **Table 2:** Results from the PCR tests for herpesvirus and mycoplasma infections according to
605 taxon and in function of the situation where the tortoises were found. TeHV stands for
606 Herpesvirus, SN for Serum Neutralization, PCR for polymerase chain reaction test,
607 Mycoplasma for *Mycoplasma agassizii*. Most individuals were tested for both pathogens
608 with PCR (TeHV + Myc). Sample size is provided (n); the sign + indicates the number of
609 positive tests associated. Percentages of positive tests are provided in brackets.

Test	Taxon	Free-ranging		Captive (pet)		Vagrant		TOTAL	
		n	+	n	+	n	+	n	+
TeHV SN	<i>T. h. hermanni</i>	253	0	48	0	8	0	309	0
	<i>T. h. boettgeri</i>	2	0	6	0	1	0	9	0
	<i>T. g. ibera</i>	1	0	0	0	0	0	1	0
	<i>T. g. sp</i>	8	0	10	0	2	0	20	0
	<i>T. marginata</i>	0	0	0	0	0	0	0	0
	WHT x EHT	32	0	6	0	3	0	41	0
	TOTAL	296	0	70	0	14	0	380	0
TeHV PCR	<i>T. h. hermanni</i>	253	7 (2.8)	58	0	10	0	321	7
	<i>T. h. boettgeri</i>	2	0	7	0	1	0	10	0
	<i>T. g. ibera</i>	1	0	0	0	0	0	1	0
	<i>T. g. sp</i>	9	0	12	0	2	0	23	0
	<i>T. marginata</i>	0	0	1	0	0	0	1	0
	WHT x EHT	35	0	4	0	5	0	44	0
	TOTAL	300	7 (2.3)	82	0	18	0	400	7 (1.7)
Mycoplasma PCR	<i>T. h. hermanni</i>	391	34 (8.7)	65	5 (7.7)	11	4 (36.34)	467	43 (9.2)
	<i>T. h. boettgeri</i>	2	0	8	0	1	0	11	0
	<i>T. g. ibera</i>	1	1 (100)	0	0	0	0	1	1 (100)
	<i>T. g. sp</i>	9	4 (44.4)	11	2 (18.2)	2	0	22	6 (27.3)
	<i>T. marginata</i>	0	0	1	1 (100)	0	0	1	1 (100)
	WHT x EHT	45	0	6	1 (16.7)	7	0	58	1 (1.7)
	TOTAL	448	39 (8.7)	91	9 (9.9)	21	4 (19.0)	560	52 (9.2)
TeHV + Myc PCR	<i>T. h. hermanni</i>	250	0	58	0	10	0	318	0
	<i>T. h. boettgeri</i>	2	0	7	0	1	0	10	0
	<i>T. g. ibera</i>	1	0	0	0	0	0	1	0
	<i>T. g. sp</i>	9	0	11	0	2	0	22	0
	<i>T. marginata</i>	0	0	1	0	0	0	1	0
	WHT x EHT	35	0	4	0	5	0	44	0
	TOTAL	297	0	81	0	18	0	396	0

610

611 **Table 3:** ~~results~~ Results from the PCR tests for herpesvirus (TeHV) and mycoplasma
 612 (*Mycoplasma*) infections (~~N=457 free-ranging tortoises~~ of free-ranging tortoises) in the 18
 613 sites sampled. Each site hosts a free-ranging population of WHT and often exotic tortoises.
 614 Other stands for tortoises opportunistically sampled in various locations. Numbers of
 615 individuals sampled and tested positive in each site are provided (n, +). The percentage of
 616 positive tests is indicated into brackets.

617

Sites	TeHV		<i>Mycoplasma</i>	
	n	<u>positive results+</u>	n	<u>positive results+</u>
3 caps	12	0	18	2 (11.1)
La Pardiguière	28	0	28	1 (3.6)
Callas	17	0	24	4 (16.7)
Carcès	10	0	10	2 (20)
Cogolin-La Môle	20	4 (20)	23	0
Estérel	18	0	28	5 (17.9)
Flassans calcaire	13	0	21	0
Lambert	15	0	23	1 (4.3)
Le Muy - St Luen	12	1 (8.3)	21	2 (9.5)
Les Arcs	13	0	12	3 (25)
Neuf riaux / Est PDM	11	0	29	1 (3.4)
Ramatuelle	9	2 (22.2)	16	1 (6.3)
Redon	18	0	24	2 (8.3)
Reserve National des Maures	18	0	21	3 (14.3)
Roquebrune sur Argens	11	0	17	0
Les Mayons	25	0	60	5 (8.3)
Sainte-Maxime	15	0	18	3 (16.7)
Vidauban	19	0	35	2 (5.7)
Other	16	0	20	2 (10)
TOTAL	300	7 (2.3)	448	39 (8.7)

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Tabel met opmaak