

1 **Lost and found: helminths infecting invasive raccoons introduced to Italy**

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27

28 **Abstract**

29 North American raccoons (*Procyon lotor*) have been introduced to several European countries,
30 where they may represent a sanitary threat as hosts of several pathogens such as the zoonotic
31 ascarid *Baylisascaris procyonis*. We carried out parasitological analysis on raccoons introduced to
32 Italy to verify whether the species had carried along *B. procyonis* or any other gastro-intestinal
33 helminths that may threaten humans, livestock or native wildlife. We examined 64 raccoons culled
34 in northern Italy during control activities and 3 roadkills opportunistically sampled from a separate
35 population located in central Italy. Helminths were collected from the gastro-intestinal tract
36 through standard parasitological techniques and identified based on a combination of morphology
37 and molecular methods. Overall, examined raccoons showed a poor parasitic fauna, with almost
38 30% of individuals free of any helminth infection. The most prevalent species were the nematodes
39 *Strongyloides procyonis* (26.9%), *Aonchotheca putorii* (25.4%) and *Porrocaecum* sp. (19.4%).
40 *Plagiorchis* sp. trematodes were also common (13.4%), whereas cestodes were scarcely
41 represented. With the exception of *S. procyonis* introduced from North America, all the other
42 identified taxa have either a European or a Palearctic distribution. Despite not finding any *B.*
43 *procyonis* in the examined raccoons, passive surveillance for this parasite should be implemented,
44 especially in Tuscany, since the limited host sample examined in the present survey does not allow
45 to exclude its presence.

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47 **Keywords**

48 alien species; macroparasites; biological invasions; gastro-intestinal helminths; *Baylisascaris*
49 *procyonis*; *Procyon lotor*

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51 The northern raccoon (*Procyon lotor*) is a carnivore native from North America that has been
52 introduced in several European countries, the Caucasian region, Japan and some Caribbean
53 islands, either deliberately as a game species or through releases from pet or fur trade [1]. Impacts
54 by invasive raccoons may include predation on native fauna and competition with other small- and
55 medium-sized carnivores, such as the endangered European mink (*Mustela lutreola*) [1]. From a
56 sanitary point of view, raccoons represent a risk towards humans, domestic animals and wildlife as
57 they are known to harbour a great variety of pathogens and are an opportunistic species that can
58 adapt to a wide range of habitats, including urban and peri-urban environments [2]. In its native
59 range the species is considered a reservoir for both rabies and canine distemper virus and is the
60 definitive host of the zoonotic ascarid *Baylisascaris procyonis* [2].
61 Adult *B. procyonis* inhabit the small intestine of raccoons and the infection is usually subclinical: in
62 North American populations, this nematode may naturally infect up to 90% of raccoons [3].
63 However, birds and mammals other than raccoons, humans included, may act as paratenic or
64 dead-end hosts and become infected with *B. procyonis* in the larval stage. In these hosts somatic
65 migration of larvae can sometimes cause extensive tissue damage and granulomatosis, potentially
66 resulting in visceral, ocular or neural Larva Migrans Syndrome (LMS) [3]. Compared to other
67 ascarids, *B. procyonis* larval migration appears particularly aggressive and leads more frequently to
68 severe or fatal neural damage [3]. Additionally, the opportunistic behaviour of raccoons coupled
69 with the high reproductive output of *B. procyonis* and the high resistance of its eggs to
70 environmental degradation, result in a relevant risk of exposure for humans and domestic animals
71 [3]. Other than in raccoons' native range, adult *B. procyonis* have been reported so far in feral
72 raccoon populations in Germany and Denmark [2,4] and eggs have been detected in raccoons'
73 latrines in Poland [5].
74 In Italy, raccoons have currently established two separate populations of different origin: a first
75 one that has been expanding since 2004 along the southern part of the Adda river (North Western
76 Italy, [6]) and a second one of more recent origin (2013) in Tuscany (central Italy, [7]). In the
77 European Union, the species is listed among the invasive species of European concern (EU
78 Regulation 1143/2014), for which management by member countries is mandatory. In 2016, Adda
79 Nord Regional Park and Lombardy region implemented a control program aimed at eradicating the
80 main raccoon population located in Northern Italy [6]. We surveyed the gastro-intestinal parasite
81 community of culled Italian raccoons, to verify if they introduced *Baylisascaris procyonis* or if they
82 harbour any other helminth that may represent a threat to humans, domestic animals or wildlife.

83 From 2017 to 2019, we examined a total of 64 raccoons (27 males and 37 females) culled in
84 Lombardy region. Additionally, we examined three road-killed animals opportunistically sampled
85 from the Tuscany population, where culling activities are not yet in place. All the animals from
86 Lombardy were captured with live-traps (model 205, Tomahawk Live Trap Co., Tomahawk,
87 Wisconsin, U.S.A.) baited with pet food and euthanised on the field by means of CO₂ overdose,
88 following AVMA guidelines [8]. Permits were issued by the Lombardy Region (permit n° 198,
89 13/01/2017) and ISPRA (Istituto Superiore per la Ricerca Ambientale – National Institute of
90 Environmental Research; permit n° 66230, 15/11/2016). Carcasses were stored at -20°C until
91 examination, during which the gastro-intestinal tract was removed, dissected longitudinally and its
92 content flushed with tap water through two sieves (lumen 0.4 and 0.038 mm, respectively), to
93 filter out coarser material. All the culled animals were in good condition and did not show any
94 clinical signs of disease at post-mortem examination. The whole gastro-intestinal content was
95 examined under a stereomicroscope and all the helminths counted and stored in ethanol 70% for
96 morphological identification based on keys by [9–11]. Inconclusive or uncertain identifications
97 were confirmed through molecular analyses. Based on morphological features, helminths found
98 within a single host were pooled in groups of 1-8 specimens, ethanol was removed and pools were
99 subjected to crude DNA extraction performed in 15 µl of Tris-HCl buffer (10 mM, pH 8) and 2 µl of
100 proteinase K (20 mg/ml), with overnight incubation at 56°C. Samples were put at 95°C for 10
101 minutes to inactivate proteinase K, then centrifuged at 20000 g for 10 minutes and stored at -20
102 °C until subsequent analyses. The supernatant (undiluted and 1:10, 1:100 serial dilutions) was
103 subjected to qualitative PCR analyses using primers reported in Supplementary Table 1 (primers
104 final concentration 1 µM). PCR products were loaded on agarose gel, excised and gel-purified
105 using the Wizard® SV Gel and PCR Clean-Up System (Promega) and Sanger sequenced. The
106 electropherograms were manually corrected and sequences were subjected to BLAST analysis
107 (<https://blast.ncbi.nlm.nih.gov/>).

108 Forty-eight (71.6%) out of the 67 examined raccoons (46/64 from Lombardy and 2/3 from
109 Tuscany) were infected by gastro-intestinal helminths. We found a total of **at least** 11 different
110 helminth taxa, with richness in infected animals ranging from 1 to 5 taxa (mean value: 1.2 ± 0.1
111 SE). *B. procyonis* was not detected in any of the examined individuals. Detailed results of
112 parasitological examination are reported in Table 1.

113 Overall, the most prevalent species found in Italian raccoons was the North American nematode
114 *Strongyloides procyonis* (prevalence 26.9%), which was also the only gastro-intestinal helminth

115 detected in roadkills sampled from the Tuscany population. The amplification and sequencing of
116 partial 18S rRNA, 28S rRNA and Internal Transcribed Spacer 1 (ITS1) showed 100% identity scores
117 with *S. procyonis* sequences present in GenBank (AB205054; [12]). Accession numbers of **of newly**
118 **obtained sequences at GenBank are shown in Supplementary Table 2**. *S. procyonis* is a nematode
119 specific to raccoons [13] and, to the best of our knowledge, this represents its first report in Italy.
120 The species has also been reported in invasive raccoons in Japan [12] and Poland [5]. With the
121 exception of *S. procyonis*, which has a clear North American origin, all the other taxa identified in
122 the present survey **have been recorded in Europe**, having a Palearctic or a wider **Holarctic**
123 distribution. This is the case for *Aonchotheca putorii*, which was the second most prevalent species
124 in our sample (25.4%). *A. putorii* is a capillariid nematode that frequently infects mustelids and
125 other medium-sized carnivores, raccoons included, in both North America and Eurasia [10,14]. The
126 genetic sequences of our specimens (Suppl. Table 2) showed respectively 100% and 98.12-98.44%
127 identity with 18S rDNA and cytochrome *c* oxidase subunit I (*COXI*) *A. putorii* sequences present in
128 GenBank (e.g. LC052363, KC355430). Larvae belonging to the cosmopolitan genus *Porrocaecum*
129 were also rather common in our sample (19.4%). The obtained 18S rRNA, 5.8S rRNA and
130 cytochrome *c* oxidase subunit 2 (*COXII*) gene fragment sequences were deposited in GenBank
131 (Suppl. Table 2). However, molecular analyses were not able to discriminate the analyzed
132 specimens at the species level. *Porrocaecum* spp. are ascarid nematodes with earthworms and
133 birds of prey as intermediate and definitive hosts, respectively [11]. By feeding on earthworms,
134 small and medium-sized mammals may become infected with the parasite in the larval stage and
135 act as paratenic hosts. Larval stages of *Porrocaecum* spp. have been reported also in raccoons
136 introduced in Japan [15]. Eight raccoons were found to harbour immature *Hystrichis tricolor*
137 individuals, which were identified based on keys by [9]. A fragment of the 18S rRNA of this species
138 was amplified, sequenced and deposited in GenBank (Suppl. Table 2). Similarly to *Porrocaecum*
139 spp., this nematode has earthworms as intermediate hosts and the infection in raccoons appears
140 of accidental nature, since adults of this species normally infect waterfowl and have been very
141 rarely reported in mammals [11].

142 Trematodes were found in 13.4% of examined raccoons but, based on molecular analysis, were
143 identified as *Plagiorchis* sp., a ubiquitous genus in European freshwater ecosystems [16].
144 Unfortunately, it was not possible to further define our specimens, due to contrasting sequencing
145 results of 28S rDNA, ITS2, and *COXI* (Suppl. Table 2). Finally, Cestodes were found only in five
146 hosts. Morphological identification and molecular analyses placed the cestode specimens

147 obtained from three of these raccoons in the genus *Dilepis*. However, the obtained gene
148 sequences (Suppl. Table 2) did not match with *Dilepis undula* (the only representative species
149 whose sequences are available in GenBank), showing the following identities scores with it:
150 99.64% for the 12S rDNA, 100% for the 18S rDNA gene, 92.49-95.43% for the *COXI* gene (reference
151 sequences used for comparison: L49457, AF286981 and EU665471, respectively). This genus
152 includes mostly species parasitising birds and small rodents [17] and, to the best of our
153 knowledge, has never been previously reported in raccoons. Unfortunately, the specimens found
154 in the two remaining raccoons were not identified due to morphological and DNA degradation.
155 Overall, our findings show that raccoons introduced to Italy have lost many gastro-intestinal
156 parasites commonly found in their native range (e.g. [18,19]). As a result, despite having acquired
157 a few European taxa, they have an impoverished community compared to North American
158 populations [cf. 18, 19], both in terms of species richness and parasite abundance. Indeed, almost
159 30% of examined individuals resulted entirely free from gastro-intestinal parasites. The exact
160 origin of the two Italian raccoon populations is unclear, but this parasite loss is likely the result of a
161 combination of founder effect and anthelmintic treatments, as the two populations were both
162 probably founded by a few pets escaped or released from captivity. Introduction of parasites by
163 invasive host species depends indeed on several factors, but it is first and foremost a stochastic
164 process [20]. For instance, as mentioned above, raccoons established in central Europe carried
165 along *B. procyonis* from their native range, but the infection appears to be absent from wild
166 populations in other introduction ranges, such as Japan [15, 21]. Whatever the cause, this reduced
167 parasite load could represent an advantage for invading hosts (enemy-release hypothesis, [20]).
168 The introduction of North American *S. procyonis* to Italy does not appear particularly alarming
169 from a sanitary point of view, as in humans this parasite is known to cause a mild dermatitis at the
170 most [13]. However, its presence deserves further attention because its potential impact on naïve
171 native hosts is unknown. In particular, the risk for spillover would be higher towards other small-
172 and medium-sized carnivores that share their habitat with the invader, such as stone and pine
173 martens, weasels, badgers, foxes or even domestic dogs. Indeed, the conspicuous presence of *A.*
174 *putorii* in our raccoon population suggests that interspecific transmission with native mustelids
175 and other wild carnivores is already occurring. Finally, our survey suggests that *B. procyonis* is
176 absent from Italian raccoons, but this result should be taken with caution. First of all, we cannot
177 draw any conclusion about the Tuscany population, because of the very limited number of
178 individuals examined through opportunistic sampling. Hence, further parasitological analyses need

179 to be carried out to specifically ascertain the absence of the ascarid from this nucleus, which has a
180 distinct origin from the North Italian one. Secondly, assuming intermediate to low prevalences of
181 *B. procyonis*, the Lombardy sample size appears adequate for a parasitological survey. However,
182 modelling data on *B. procyonis* dynamics suggest that, during the initial stages of host invasion,
183 the parasite might persist at very low prevalences that may hinder its detection [22]. As a
184 consequence, ongoing passive surveillance is recommended to rapidly detect any emergence of *B.*
185 *procyonis* in Northern Italy [23].

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258 **Table 1** Gastro-intestinal helminths detected in raccoons (*Procyon lotor*, N=67) introduced to Italy:
 259 number of infected hosts (n), prevalence of infection (% infected hosts/examined hosts) and mean
 260 intensity of infection (no. parasites/infected hosts); SE=Standard Error.

Taxon	n	Prevalence ± SE	Mean intensity ± SE
Nematoda			
<i>Strongyloides procyonis</i>	18	26.9 ± 5.4	11.7 ± 5.5
<i>Aonchotheca putorii</i>	17	25.4 ± 5.3	7.2 ± 1.9
<i>Porrocaecum</i> sp. (larvae)	13	19.4 ± 4.8	14.7 ± 5.3
<i>Hystrichis tricolor</i> (immature)	8	11.9 ± 4.0	8.5 ± 6.8
Trichuridae spp. ^a	2	3.0 ± 2.1	1; 2 ^b
Trichostrongylidae spp. ^a	2	3.0 ± 2.1	2; 3 ^b
Ancylostomatidae sp.	1	1.5	1 ^b
Oxyuridae sp.	1	1.5	1 ^b
Trematoda			
<i>Plagiorchis</i> sp.	9	13.4 ± 4.2	7.0 ± 3.9
Cestoda			
<i>Dilepis</i> sp.	3	4.5 ± 2.5	4; 1; 11 ^b
Unidentified	2	3.0 ± 2.1	1; 26 ^b

261 ^a uncertain number of species, single or two; ^b intensity of infection in each infected host

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