

Examination of Helminth Parasites in the Fecal Samples of Magpies (*Pica pica*), Jackdaws (*Corvus monedula*), and Rooks (*Corvus frugilegus*) in the Vicinities of Van Province/Turkey

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Abstract: The present study was designed to examine helminths in the gastrointestinal tract of corvid species including *Pica pica*, *Corvus frugilegus*, and *Corvus monedula*. A total of 43 fecal samples were collected from *Pica pica*, 42 samples from *Corvus frugilegus*, and 42 samples from *Corvus monedula* species. Feces were analyzed using sedimentation and flotation methods. *Echinostoma* spp. (2.32%), *Capillaria* spp. (6.97%), *Syngamus* spp. (2.32%), and *Raillietina* spp. (6.97%) were detected in *Pica pica*, *Raillietina* spp. (2.5%) and *Capillaria* spp. (10.0%) were detected in *Corvus monedula*, and only *Capillaria* spp. (9.52%) were detected in *Corvus frugilegus*. To our knowledge, this is the first study in Turkey identifying these helminth species in the fecal samples of these bird species.

Keywords: *Corvidae*, Helminth, *Pica pica*, *Corvus frugilegus*, *Corvus monedula*.

Van'da (Türkiye) Saksığan (*Pica pica*), Cüce Karga (*Corvus monedula*) ve Ekin Kargası (*Corvus frugilegus*) Dışkılarının Helminth Parazitler Açısından İncelenmesi

Öz: Bu çalışma Corvidae ailesine bağlı Saksığan (*Pica pica*), Cüce karga (*Corvus monedula*), Ekin kargası (*Corvus frugilegus*) türü yabani kuşlarda gastrointestinal sistem helmintlerini tespit etmek için yapıldı. Bu amaçla Saksığan'dan 43, Cüce kargadan 40, Ekin kargasından 42 adet dışkı toplandı. Dışkılar sedimentasyon ve flotasyon yöntemleri ile incelendi. İnceleme sonucunda Saksığan'da %2.32 oranında *Echinostoma* spp., %6.97 oranında %6.97 oranında *Capillaria* spp., %2.32 oranında *Syngamus* spp., %6.97 oranında *Raillietina* spp., Cüce kargada %2.5 oranında *Raillietina* spp., %10 oranında *Capillaria* spp., Ekin kargasında %9.52 oranında *Capillaria* spp. türü helmint yumurtaları tespit edildi. Dışkı muayenelerinde yumurtalarını tespit ettiğimiz helmint türleri *Pica pica*, *Corvus monedula* ve *Corvus frugilegus*'da, Türkiye'de ilk kez bu çalışma ile bildirilmektedir.

Anahtar kelimeler: *Corvidae*, Helminth, *Pica pica*, *Corvus frugilegus*, *Corvus monedula*.

1. Introduction

Parasites are frequently seen in wild birds and these birds are known to harbor a wide variety of parasites throughout their life span. Nevertheless, there is little documentation of parasites harbored by wild birds and there is no taxonomy for most of these parasites. Moreover, although parasitic protozoa that reside in blood have been investigated extensively due to the ease of collecting blood samples from living birds, little is known about other species such as intestinal parasites (Moore & Clayton, 1997). The helminth communities of birds are closely associated with water due to their intermediary hosts and these communities are generally more diverse than those of fish and mammal hosts (Kennedy, Bush, & Aho, 1986; Bush, Aho, & Kennedy, 1990; Poulin, 1997).

Turkey is home to a wide variety of species due to its geographical location, topographic conditions, habitat biodiversity, and climatic richness (Kızıroğlu, 2001; Balkaya & Çelikoba, 2005; Akın, 2009). On the other hand, birds play a vital role in the dissemination of parasites and both birds and their hosts are affected by migration and habitat quality (Webb, 2010; Kanarek & Zalesny, 2014).

The corvid species examined to date have been

shown to live in a half-wild state. These species can disseminate in cultivated areas and natural habitats and typically feed on local trash sources and urban dump sites. These animals breed in the wildlife and may also reproduce in urbanized areas and get adapted to these environments. In their daily life, they keep contact with humans and domestic animals by frequently leaving their feces in their living quarters.

This study aimed to examine the helminths in the fecal samples of corvid species including *Pica pica*, *Corvus frugilegus*, and *Corvus monedula* which are commonly seen in urbanized areas in Van province. We also investigated the role of these bird species in infections caused by helminths.

2. Material and Methods

Fecal Samples: Samples of the three bird species were collected from their nesting sites at 18 different locations in Van Province between March and May 2019 (Fig. 1).

Samples were collected from the fresh fecal matter underneath the nests of the birds. Throughout the sample collection processes, no contact was made with the birds and no bird was injured. A total of 43 fecal samples were

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collected from *P. pica*, 42 samples from *C. frugilegus*, and 42 samples from *C. monedula* species. Birds selected for the study were identified by an expert ornithologist using the relevant literature (Svensson, Mullarney, & Zatterström, 2009). All the samples were placed in sealed specimen containers with labels indicating the date, location of sample collection, and the species. The samples were transferred to the laboratory on the same day. Feces were analyzed on the same day using sedimentation and flotation methods. In the flotation method, hazelnut-sized feces were homogenized in 33% Zinc Sulphate in the tube. The top of the tube was covered with a coverslip and after

30 minutes, the coverslip was taken on the slide. In the sedimentation method, the feces were suspended with tap water or saline and filtered. The top liquid was poured after centrifugation at 400-500 g for 2 minutes. By taking some amount from the sediment, the smear was prepared. The thin layer was coated with another slide and then was examined under a microscope (Leica DM500) at 10× and 40× objective lens. The helminth eggs detected in the samples were identified based on the literature data and then were photographed (Mshot Md 90) (Kaufman, 1996; Fernando & Barta, 2008).

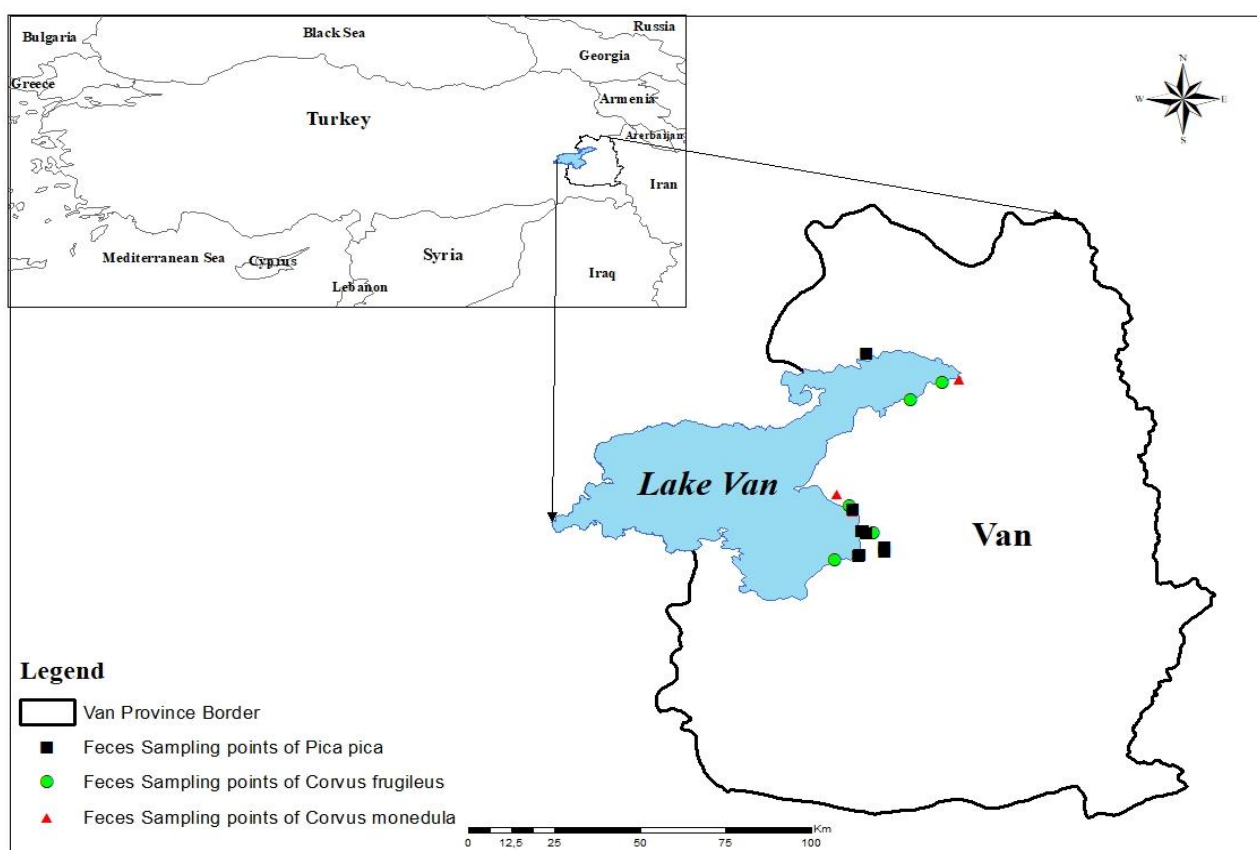


Figure 1. Sample collection sites

3. Results

It was revealed that all three corvid species nested on trees in urbanized areas. It was also found that both *C. frugilegus* and *C. monedula* established hundreds of breeding and feeding communities, while *P. pica* lived in individual communities. However, the territories of all three species overlapped with each other.

A total of four distinct helminth species were identified in the three bird species. *Echinostoma* spp., *Capillaria* spp., *Syngamus* spp., and *Raillietina* spp. were detected in *Pica pica*; *Raillietina* spp. and *Capillaria* spp. were detected in *C. monedula*; and only *Capillaria* spp. were detected in *C. frugilegus* (Table 1). The egg sizes of *Echinostoma* spp., *Capillaria* spp., *Syngamus* spp., and *Raillietina* spp. were $\sim 95 \times 119 \mu\text{m}$, $\sim 47 \times 25 \mu\text{m}$, $\sim 92 \times 47 \mu\text{m}$, and $\sim 45 \times 38 \mu\text{m}$, respectively (Fig. 2).

4. Discussion

Turkey is home to a wide variety of bird species due to its natural features. These species act as hosts to various

parasites. The biodiversity of bird species and their hosts are associated with numerous factors such as habitat quality and distribution as well as seasonal changes (Webb, 2010; Kanarek & Zalesny 2014).

In the present study, *Echinostoma* spp. were detected as a mixed infection with *Capillaria* spp. in 2.32% (n=1) of the fecal samples obtained from *P. pica* species. Similarly, in the previous studies, *Echinostoma* spp. have been shown to cause severe enteritis in both domestic and wild birds, particularly ducks and geese, by settling down in their cecum and rectum and have also been reported in humans and others mammals, though rarely. (Griffiths, Gonder, & Pomeroy, 1976; Hossain, Dewan, Baki, & Mondad, 1980). In Turkey, *Echinostoma* spp. have been reported in chickens, turkeys, and geese with an infection rate of 0.06-10% (Merdivenci, 1967; Kurt & Acici, 2008; Güçlü, 1992; Aydın, Göz, & Değer, 2010). In other countries, however, *Echinostoma* spp. have been mostly reported in birds and their most common species, *E. revolutum*, has been reported in ducks, wild water birds, and pigeons (Kaufmann, 1996; Farias & Canaris, 1986; Yousuf, Das,

Anisuzzaman, & Banowary, 2009). Another study detected three Echinostome species including *Patagifer bilobus*, *Petasisger neocomense*, and *Saakotrema metatestis* in wild water birds including *Platalea minor*, *Podiceps cristatus*, and *Egretta garzetta* (Choe et al., 2014). In a previous study conducted in Montana, Todd and Worley (1967) also detected *Echinostoma* spp. in *Pica picaludsonia*. Similarly, some previous studies identified *Syngamus* spp. in the trachea, bronchi, bronchioles, and rarely in the small intestine of numerous birds including chickens, turkeys, geese, guinea fowl, pheasants, peafowls, and quails (Norton & Ruff, 2003; Fernando & Barta, 2008). In Turkey, *Syngamus* spp. have only been reported in chickens with a prevalence of 0.2-3% (Aydın, Göz, & Değer, 2010; Kurt & Acici, 2008). However, in some other countries such as Canada, Britain, Norway, USA, and Czech Republic, *Syngamus* spp. have been reported in both domestic and wild birds (Fernando & Barta, 2008; Okulewica & Koubek, 1994). In the present study, *Syngamus* spp. were detected in only 2.32% of the fecal samples obtained from *P. pica* species. *Capillaria* spp. are small thin nematodes (hair worms) in the superfamily Trichinelloidea, family Trichuridae, and subfamily Capillarinae. In Turkey, these species have been reported in chickens, geese, ducks, turkeys, domestic and wild pigeons, and sea gulls (Güçlü, 1992; Kurt & Acici, 2008; Gıcık & Burgu, 2000; Kılınç, Çiçek, & Akkaş, 2011). In other countries, although *Capillaria* spp. have been reported in various bird species, these bird species and the infection rates have been shown to vary among countries. To illustrate, *Capillaria* spp. have been reported in 1.6-54% of chickens in Denmark, 2-72% of

pheasants and 3-73% of turkeys in Brazil, and 4-36% of domestic pigeons in Poland (Permin et al., 1997; Pinto, Tortelly, Menezes, & Gomes, 2004; Stenzel & Koncicki, 2007). In our study, *Capillaria* spp. were detected as a mixed infection with *Raillietina* spp. in 6.97% (n=2) of the fecal samples obtained from *P. pica* species. Additionally, *Capillaria* spp. were detected in 9.52% (n=4) of the fecal samples obtained from *C. frugilegus* species and 10% (n=4) of the fecal samples obtained from *C. monedula*. In the literature, *Raillietina* spp. have been shown to settle down in the small intestines of chickens, turkeys, guinea fowls, partridges, quails, and pigeons and to cause weight loss and diarrhea in infected animals (Permin & Hansen, 1998; Baker, 2007). In Turkey, *Raillietina* spp. has been reported 3-12% of chickens, turkeys, partridges, and pigeons (Güçlü, 1992; Kurt & Acici, 2008; Gıcık & Burgu, 2000; Köroğlu & Taşan, 1996; Aydın, Göz, & Değer, 2010). *Raillietina* spp. is a cosmopolitan species and have been identified in various bird species in Africa, Asia, America, and Europe and is more common in young animals (Permin & Hansen, 1998; Permin et al., 1997; Magwisha, Kassuku, Kyvsgaard, & Permin, 2002; Shahin, Lebda, Abu-Elkheir, & Elmeligy, 2011; Radfar et al., 2011; Adang et al., 2008). In a study conducted on pigeons in Bursa province, *Raillietina echinobothrida* species was reported (Yıldırımhan, Gürkan, & Altunel, 2009). In a postmortem study on 10 *P. pica* in Ankara, it was reported that *Dispharinx nasuta* nematodes and *Trichomonas* sp. type protozoans were found in one magpie (Çetindağ & Brykoğlu, 1997).

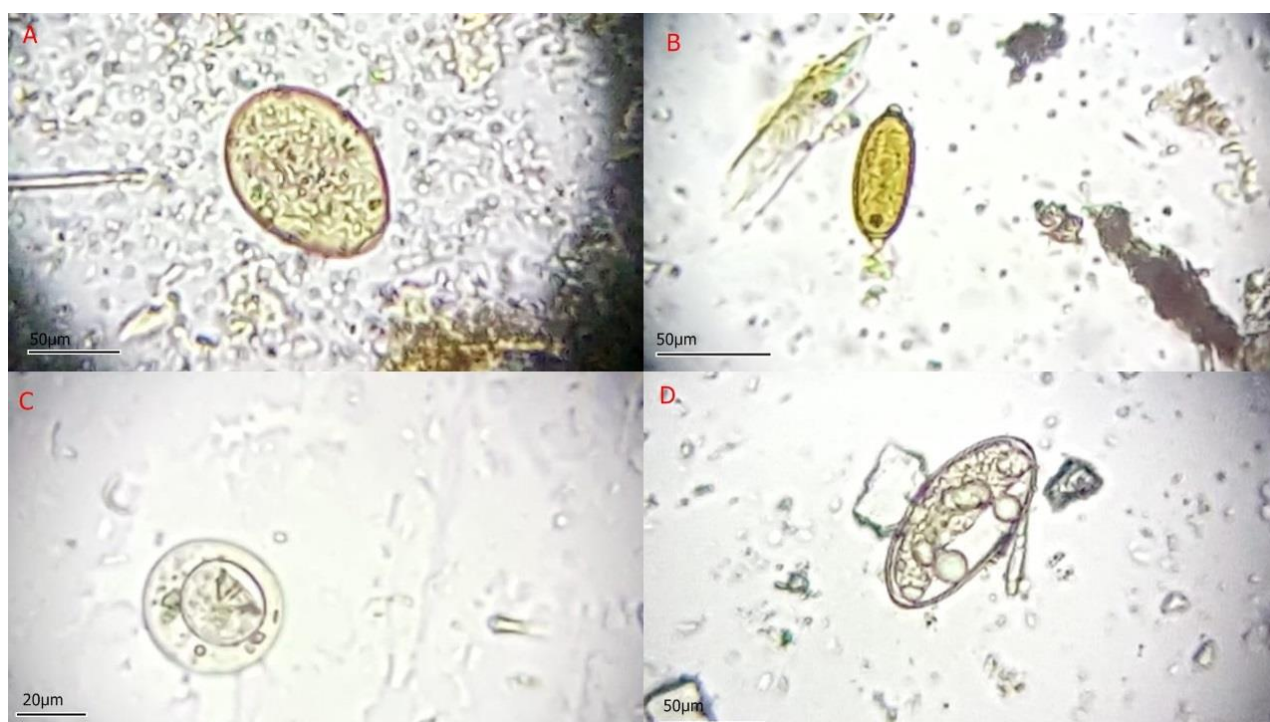


Figure 2. Helminth parasites eggs detected in fecal samples (A: *Echinostoma* spp. egg, B: *Capillaria* spp. egg, C: *Raillietina* spp. egg, D: *Syngamus* spp. egg)

In our study, however, *Raillietina* spp. were detected as a mixed infection with *Capillaria* spp. in a single fecal sample and as a single parasite in two fecal samples obtained from *P. pica* species.

Halajian et al. (2011) evaluated *P. pica*, *C. frugilegus*,

and *C. corone* in Iran and diagnosed *Syngamus trachea* in 6.3% of *P. pica* and 6.2% of *C. corone* and identified *Capillaria* spp. in 9% of *C. frugilegus*. The differences between these rates and the rates found in our study could be attributed to regional differences and to the fact that Halajian et al. (2011) examined the animals at post mortem.

Girişgin et al. (2019) conducted a helminthological examination on *P. pica* species at post necropsy and identified *Passerilepis* spp. and *Brachylaima* spp. in the animals. Interestingly, these helminth species were not detected in any of the animals examined in our study which could be associated with regional differences and with the fact that Girişgin et al. (2019) only examined eight animals that were brought to the animal hospital for

medical intervention. In our study, one trematode (*Echinostoma* spp.) one cestode (*Raillietina* spp.), and two nematodes (*Capillaria* spp. and *Syngamus* spp.) were identified in the helminth eggs detected in the fecal samples of *C. frugilegus*, *C. monedula*, and *P. pica* species (Table 1). To our knowledge, this is the first study in the literature identifying these helminth species in these bird species.

Table 1. Helminth species identified in the study

Bird species	Fecal samples (n)	Positive samples (n)	Helminth species	Infection Rate
<i>Pica pica</i>	43	1	<i>Echinostoma</i> spp.	2.32%
		3	<i>Capillaria</i> spp.	6.97%
		1	<i>Syngamus</i> spp.	2.32%
		3	<i>Raillietina</i> spp.	6.97%
		4	<i>Capillaria</i> spp.	9.52%
<i>Corvus frugilegus</i>	42	4	<i>Capillaria</i> spp.	9.52%
<i>Corvus monedula</i>	40	1	<i>Raillietina</i> spp.	2.5%
		4	<i>Capillaria</i> spp.	10.0%

5. Conclusion

In conclusion, wild birds harbor a wide variety of parasites throughout their life span and little is known about the importance of infections caused by these parasites in wild birds. Mixed parasitic infections are highly common in wild birds and the effects of these infections on wild birds are associated with the type of the parasite, coexisting parasitic diseases, and stress factors. The results indicated that the corvid species examined in the study had large populations in urbanized areas in the vicinities of Van province which could be related to the abundance of food in those areas. Additionally, this finding could most probably be related to the fact that these species need trees for reproduction and there are almost no trees in the areas other than those areas.

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