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First microscopic, pathological, epidemiological, and molecular investigation of *Leucocytozoon* (Apicomplexa: *Haemosporida*) parasites in Egyptian pigeons

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Introduction: *Leucocytozoon* is an intracellular blood parasite that affects various bird species globally and is transmitted by blackfly vectors. This parasite is responsible for leucocytozoonosis, a disease that results in significant economic losses due to reduced meat and egg production. There is limited knowledge about the epidemiological pattern of leucocytozoonosis and its causative species in Egypt, particularly in pigeons.

Methods: The current study involved the collection of 203 blood samples from domestic pigeons from various household breeders and local markets across Qena Province, Upper Egypt. Samples were initially examined for potential *Leucocytozoon* infection using blood smears, followed by an evaluation of associated risk factors. Molecular identification of the parasite in selected samples (n = 11), which had initially tested positive via blood smears, was further refined through nested PCR and sequence analysis of the mitochondrial cytochrome b gene to ascertain the *Leucocytozoon* species present. Additionally, histopathological examination of the liver, spleen, and pancreas was conducted on animals that tested positive by blood smears.

Results: Interestingly, 26 out of 203 samples (12.08%) had confirmed *Leucocytozoon* infections based on microscopic analysis. Additionally, all 11 samples that initially tested positive via blood smears were confirmed positive through nested PCR analysis, and their sequencing revealed the presence of *Leucocytozoon sabrazesi*, marking the first report of this parasite in Egypt. The study into potential risk factors unveiled the prevalence of *Leucocytozoon* spp. seems host gender-dependent, with males exhibiting a significantly higher infection rate (33.33%). Additionally, adult birds demonstrated a significantly higher infection prevalence than squabs, suggesting an age-dependent trend in prevalence. Seasonality played a significant role, with the highest occurrence observed during summer (37.25%). Histopathological examination revealed

the presence of numerous megaloschizonts accompanied by lymphocytic infiltration and multiple focal areas of ischemic necrosis.

Conclusion: To our knowledge, this is the first study to shed light on the epidemiological characteristics and molecular characterization of leucocytozoonosis in pigeons in Egypt. Further research endeavors are warranted to curb the resurgence of *Leucocytozoon* parasites in other avian species across Egypt, thereby refining the epidemiological understanding of the disease for more effective control and prevention measures.

KEYWORDS

Leucocytozoon, pigeon, Egypt, epidemiology, molecular, phylogenetic, histopathology

1 Introduction

Pigeons are abundant and ubiquitous avian species, often found in urban environments. Since ancient times, pigeons have been regarded as symbols of various concepts, including deities, peace, messengers, pets, food, and spiritual sacrifice. In Egypt, pigeons are primarily raised to meet the protein needs of families during special occasions, serve as a source of income, for gaming, and ornamental purposes. However, pigeons can host numerous pathogens and serve as reservoirs for parasitic infections (1). Parasitism is a significant concern affecting bird production, leading to issues such as growth retardation, decreased vitality, blood loss, toxicosis, and poor health conditions. Ultimately, this reduces the quality and quantity of meat and egg production. Among various parasitic diseases affecting avian species, haemoprotozoan infections are predominant (2).

Avian haematozoa comprise a class of vector-borne parasites belonging to the apicomplexan group, which includes genera such as Plasmodium, Haemoproteus, and Leucocytozoon. These parasites are transmitted by blood-sucking dipteran vectors, including species of ceratopogonids (genus Culicoides), blood-sucking culicine mosquitoes (Culicidae), blackflies (Simuliidae), and hippoboscid flies, with birds acting as intermediate hosts (2). Among others, leucocytozoonosis is considered the most significant blood protozoan disease affecting birds, caused by approximately 60 species of parasitic protozoa of the genus Leucocytozoon. It affects wild and domestic avian species and is transmitted by biting blackflies such as Simulium venustum, S. croxtoni, S. euradminiculum, and S. rugglesi (3). The life cycle of Leucocytozoon species is complex, involving development both in tissues (exo-erythrocytic merogony) and blood cells. Before infecting blood cells and forming gametocytes, Leucocytozoon species undergo exo-erythrocytic merogony, which produces meronts in tissue cells. These meronts are the infective stage for vectors. Subsequently, sexual processes and sporogony occur in dipteran insects, producing infective sporozoites, which initiate new infections in vertebrate hosts (2, 4, 5). The pathogenic impact of Leucocytozoon infection on the host can potentially jeopardize productivity, reducing egg production and increasing mortality rates. Clinical signs of infection may include anemia, anorexia, green feces, and ataxia, although infections can be asymptomatic. Upon necropsy, common findings include fatty liver, splenomegaly, regressive reproductive organs, and other characteristic lesions.

This can result in a significant loss of production value in industrial settings and group deaths, as reported in various avian species (6, 7).

Direct microscopic examination of Giemsa-stained blood films was considered the most conservative diagnostic approach for detecting Leucocytozoon sp. infection. Additionally, identifying the parasite's genome using polymerase chain reaction (PCR) with primers derived from mitochondrial genes offers a more sensitive and accurate method widely employed in laboratory settings for precise analysis of infections. This molecular approach can provide high accuracy even in cases where blood smears are negative due to low parasitemia or early stages of infection in avian hosts (8-10). Understanding the epidemiological patterns of parasitic infections is essential for devising and implementing effective prevention and control strategies. Investigating the previous literature, very scant information is available on Egypt's Leucocytozoon sp. infection. Only one previous study in northern Egypt (lower Egypt) revealed the natural co-infection of poultry farms with *Leucocytozoon caulleryi* and chicken anemia virus (11). However, to the best of our knowledge, no specific investigations have been conducted to explore the incidence of blood parasitic infections, particularly Leucocytozoon species, in the country's southern region (Upper Egypt). Therefore, the current study was conducted to identify and determine the prevalence of Leucocytozoon species in pigeons from Upper Egypt and assess the associated risk factors through microscopic examination of stained blood smears. Additionally, this investigation examined the taxonomy of the identified leucocytozoids at the species level by analyzing the phenotypic characteristics of the cytochrome b gene (cytb) and reporting the major histopathological findings of the examined animals.

2 Materials and methods

2.1 Ethical statements

The present study received approval from the Ethics Committee of the Faculty of Veterinary Medicine at South Valley University, Egypt, per ethical regulations and animal research guidelines (permit code No. 84). Written and oral consent was obtained from each owner of the surveyed pigeons.

2.2 Study area

The study was conducted in Qena Province, situated in southern Egypt at coordinates 26°10′12″N 32°43′38″E. Renowned for its pottery, imposing mountains, and lush green landscapes, the province experiences a hot desert climate characterized by scorching summers and minimal yearly precipitation.

2.3 Birds and sample processing

Blood samples (n=203) were randomly collected from apparently healthy pigeons in Qena Province between November 2020 and October 2021, sourced from different household breeders and local markets. Information on age, sex, and sampling season was documented to evaluate potential associations with the presence and abundance of blood parasites. Each bird's sample (3 mL) was gathered in an anticoagulated test tube from the brachial wing vein using a sterile syringe and needle. These samples were transported to the Parasitology Laboratory at the Faculty of Veterinary Medicine, South Valley University, for parasitological analysis.

2.4 Laboratory analysis

Following collection, thin blood films were immediately prepared from each sample to identify blood protozoa. The smears were air-dried, fixed in absolute methanol three times for 10s each, and stained with Giemsa's stain (30%) for 10min. Subsequently, the slides were gently washed under running tap water, air-dried, and then subjected to microscopic examination, following established laboratory protocols (12). The stained slides were examined using an Olympus CX31 microscope at higher magnification (100X) to detect infections. The identification and intensity of recovered haemoprotozoa were recorded following established keys and descriptions outlined by Soulsby (13) and Levine (14).

2.5 Histopathological analysis

The examined pigeons were anesthetized using an equal mixture of ketamine and xylazine (0.0044 cc/kg), administered via injection into the pectoral muscle (15), then left to ensure complete euthanasia. Tissue samples, mainly liver, spleen, and pancreas, were then excised and prepared for histopathological examination (16). Approximately 1 cm sections of each tissue were collected and fixed in 10% neutral buffered formalin (pH=7.4). They underwent processing through ascending grades of alcohols, were embedded in paraffin wax, sectioned at a thickness of 5 μ m, and then stained with histochemical stains (Harries hematoxylin and eosin, Sigma-Aldrich) (17). The preparations were examined using a microscope (Olympus BX51, Tokyo, Japan) equipped with a camera (Olympus E-182330, Olympus Optical Co., Ltd., Japan), with five slides inspected for each block.

2.6 Molecular identification

2.6.1 DNA extraction

DNA was extracted successfully from 11 positive blood samples via microscopic examination. This extraction was performed using a

QIAamp DNA mini kit (1,043,368, Qiagen, United States) following the manufacturer's instructions, and the extracted DNA was stored at -20° C until PCR analysis.

2.6.2 PCR amplification

Two pairs of specific primers from Macrogen (Korea) were utilized to amplify the *cytb* gene via nested PCR. The first step of amplification employed the primers LsF1 (5'-CATATATAAGAGAATTATGGAG-3') and LsR1 (5'-ATAAAATGYTAAGAAATACCATTC-3'). In the second step, the primers LsF2 (5'-TAATCACATGGGTTTGTGGA-3') and LsR2 (5'-GCTTTGGGCTAAGAATAATACC-3') were utilized. The expected size of amplification products was 248 bp. The reaction was conducted in a 25 μ L volume containing 12.5 μ L of DreamTaq Green PCR Master Mix (2X) (K1081, ThermoFisher, United States), 1 μ L of each primer (20 pmol), 5.5 μ L of water, and 5 μ L of DNA. PCR was performed using an Applied Biosystems 2,720 thermal cycler with the following program: 40 cycles of denaturation at 94°C for 1 min (1st step) and 20s (2nd step), annealing at 50°C for 1 min (1st step) and 30s (2nd step), followed by a final extension at 68°C for 5 min (1st and 2nd steps).

2.6.3 Visualization of the PCR outcomes

The PCR products were subjected to electrophoresis on a 1.5% agarose gel (Agarose, Universal, PeqGold, Peqlab. Germany) in 1x TBE buffer. About 20 μ L of the PCR products was loaded onto the gel. A 50bp DNA ladder gene marker (PeqGold 2kb DNA-Ladder, Peqlab., VWR) was utilized to determine the size of the amplicons. The gel was stained with ethidium bromide (0.5 μ g/mL) and visualized under UV light. The results were documented using a gel documentation system (Geldoc-it, UVP, England).

2.6.4 Sequencing and phylogenetic analysis

The PCR products were purified using a QIAquick PCR product extraction kit (Qiagen, Valencia) and Centrisep spin columns. The gel documentation system (Geldoc-it, UVP, England) was utilized to capture the sequence reaction, and analysis was conducted using Totallab analysis software.¹ The identity of the obtained DNA sequences from the ABI PRISM[®] 3,100 Genetic Analyzer (Micron-Corp. Korea) was confirmed through BLAST analysis (18). The phylogenetic tree was constructed using the MegAlign module of Lasergene DNA Star version 1.83 software, based on the *cytb* gene sequences (19). The analysis was conducted in MEGA11 using the accession number (ON399180) with maximum likelihood method (20). The sequences, along with their corresponding host and locality or country, were downloaded from GenBank and used in the tree construction, as depicted in Table 1.

2.7 Statistical analysis

Statistical analysis was conducted to assess the variation in *Leucocytozoon* spp. incidence among pigeons, considering the epidemiological data. The chi-square (χ^2) test was employed using IBM SPSS Statistics for Windows, Version 21.0 (IBM Corp., Armonk,

¹ www.totallab.com, Ver. 1.0.1

Accession number	Species	Host origin	Location/country
MW316431	Leucocytozoon sabrazesi	Chicken	Thailand
MW316432	Leucocytozoon sabrazesi	Chicken	Thailand
MW316434	Leucocytozoon sabrazesi	Chicken	Thailand
KT290930	Leucocytozoon sabrazesi	Gallus gallus spadiceus	Malaysia
MZ634390	Leucocytozoon sabrazesi	Gallus gallus	Thailand
KT290929	Leucocytozoon sabrazesi	Gallus gallus domesticus	Malaysia
AB299369	Leucocytozoon sabrazesi	Chicken	Malaysia
LC550031	Leucocytozoon sabrazesi	Chicken	Myanma
MW600919	Leucocytozoon caulleryi	Gallus gallus	Thailand
MN540144	Plasmodium kentropyxi	Lizards	Brazil
MW296834	Haemoproteus sp.	Avian	Korea
JQ988310	Parahaemoproteus sp.	Coeligena torquata	Peru
MK721052	Leucocytozoon sp.	Emberiza godlewskii	China
MK061720	Haemoproteus sp.	Pachycephala hyperythra	Papua New Guinea
GU59370	Eimeria acervulina	Gallus gallus	United States

TABLE 1 The accession numbers used for construction of the phylogenetic tree, along with their species, hosts, and locations.

TABLE 2 Prevalence of Leucocytozoon spp. in relation to the age, sex, and season of the pigeons inspected in the present study.

Variables	No. of examined cases	No. of positive cases (%)	Pearson Chi-Square χ² (p value)
Age			19.063 (< 0.0001)*
< 2 months	86	0 (0)	
> 2 months	117	26 (22.22)	
Sex			8.836 (0.002)*
Male	21	7 (33.33)	
Female	182	19 (10.43)	
Season	36.593 (<0.05)*		
Winter	47	0 (0)	-
Spring	52	6 (1.92)	
Summer	51	19 (37.25)	
Autumn	53	1 (1.88)	

*Superscript indicates the significant difference at p < 0.05.

NY, United States). A significance level of $p \le 0.05$ was considered indicative of statistical significance (21).

3 Results

3.1 Occurrence of *Leucocytozoon* species and potential risk factors

Out of 203 inspected pigeons, 26 were infected with *Leucocytozoon* species, yielding an overall prevalence of 12.80%. Regarding infection frequency (Table 2), *Leucocytozoon* spp. showed significantly higher rates in the age group of >2 months (22.2%, $\chi^2 = 19.063$, p < 0.05), while the other age category remained unaffected, suggesting that the risk of infection rises with age. Likewise, the current investigation revealed that the occurrence percentage of *Leucocytozoon* spp. was 10.43% in females, whereas it

was 33.33% in males, indicating a statistically significant gender disparity in infection rates ($\chi^2 = 8.836$, p = 0.002) as depicted in Table 2. Additionally, the same table presents the prevalence of parasite infections relative to both age and sex groups in examined pigeons. It is intriguing that the proportion the occurrence rate of *Leucocytozoon* infection was notably higher in males (46.66%) compared to females (18.62%) within the same age category (>2 months), which demonstrated a statistically significant difference ($\chi^2 = 13.079$, p = 0.004). Conversely, no infections were observed in either male or female pigeons in the younger class (<2 months).

Investigation of seasonal dynamic of infection with *Leucoytozoon* species showed that the summer season exhibiting the highest prevalence of *Leucoytozoon* infections (37.25%), followed by spring (1.92%) and autumn (1.88%). No infection was observed during the winter (0%), indicating a notable difference in infection seasonality, as shown in Table 2. Additionally, the variations in infection rates between seasons were statistically significant (χ^2 = 36.593, p < 0.05).

3.2 Morphological description

Gametocytes of *L. sabrazesi* were detected in positive Giemsastained blood films measuring $7.75 \times 9.20 \,\mu$ m. Mature gametocytes appeared as a distinct parasite stage (Figure 1), occupying the entire cellular space and replacing the cell cytoplasm, occasionally forming elongated "horns." Macrogametocytes appeared darker, with small nuclei, dark blue cytoplasm, light red nuclei, small vacuoles, and magenta volutin cytoplasmic granules. In contrast, microgametocytes showed lighter blue staining, with extremely pale cytoplasm and pale pink nuclei (Figure 2). Additionally, an average of 3 ± 0.4 *Leucocytozoon* sp. infected cells per field were recorded.

3.3 Gross lesions and histopathological findings

During necropsy, the examined pigeons typically exhibited no noticeable gross lesions. However, microscopic examination documented the presence of *Leucocytozoon* spp. in different organs, including the liver, spleen, and pancreas.

Histopathological examination of liver tissue (Figure 3) revealed focal areas of coagulative necrosis with distortion of hepatocytes. The periportal areas exhibited heavy infection with multiple variable-sized megaloschizonts containing numerous basophilic schizonts. These schizonts were observed solitarily or in groups, often surrounded by



FIGURE 1 Mature macrogametocyte (A) and microgametocyte (B) of *Leucocytozoon sabrazesi* (arrows).



FIGURE 2

Leucocytozoon sabrazesi macrogametocytes (A–C) and microgametocytes (D,E) in pigeon's cells. (A,E) gametocytes in elongated form and (C,D) gametocytes in round form in pigeon's leukocyte cells (C,D). The black arrow indicates host cell nuclei, the red arrow indicates nuclei of parasites, the yellow arrow indicates vacuoles, and the violet arrow indicates volutin granules. Scale bar = 10 µm.

well-defined intact or depleted capsules. Additionally, there was notable infiltration of lymphocytes and macrophages around the schizonts' periphery, contributing to hepatic cell necrosis near the portal areas. Other notable findings included portal vein congestion, blood sinusoid widening, fatty degeneration, and multiple thromboses in small and medium-sized hepatic vessels. Furthermore, congestion of the hepatic artery with thickening of its wall accompanied by amorphous eosinophilic infiltration was observed (Figure 3).

In the case of splenic infection (Figure 4), fewer megaloschizonts were observed in the interstitial tissue, accompanied by reticular hyperplasia. Intracellular hemosiderosis was noted as evidence for the destruction of erythrocytes. Moreover, lymphoid depletion and



FIGURE 3

(A) Histopathological lesions of *Leucocytozoon* infection in a pigeon's liver illustrate multiple necrosis foci (5-Point Star).
(B) Numerous developing megaloschizont aggregations in the portal area (Left & Right Arrows).
(C) Dilated blood sinusoids (Isosceles Triangle).
(D) Fatty degeneration (Arrow).
(E) Blood thrombosis (Oval).
(F) Eosinophilic infiltration (Double arrow), H&E stain.



FIGURE 4

(A) Histopathological lesions of *Leucocytozoon* infection in the spleen of a pigeon illustrating necrotic areas with lymphocytic depletion (Arrows).
(B) Few megaloschizonts distribution (5-Point Star).
(C) Heamosedrosis (Left & down Arrows).
(D) Lymphocytic cell infiltration and eosinophilic structures with vascular congestion (Oval).
(E,F) Bands of fibrous tissue (Isosceles Triangle), H&E stain.



FIGURE 5

(A,B) Histopathological lesions of *Leucocytozoon* infection in a pigeon's pancreas illustrate obvious necrosis areas with mild fibrous tissue proliferation (5-Point Star & Double Arrow). (B) Lymphocytic infiltration (Oval). (C) Numerous megaloschizonts (Isosceles Triangles). (D) Zymogen granules depletion (Arrows). (E) Apoptotic acinar cells (Elbow Arrow Connector). (F) Disorganized acinar cells (Curved Right Arrow), H&E stain.

spleen atrophy with numerous eosinophilic structures (due to destructed megaloschizonts) were observed adjacent to the congested splenic artery. Bands of fibrous tissue were detected among splenic tissue.

Regarding histopathological lesions in the pancreas (Figure 5), acute pancreatic necrosis was observed, represented by zymogen granules depletion and shrinkage in the exocrine cells with degeneration in islets of Langerhans, resulting from megaloschizonts distribution. Bands of fibrous tissue were observed to separate the pancreatic ductular system, accompanied by multifocal mononuclear cell infiltrations consisting of lymphocytes and eosinophils. These features resulted in an apparent loss of acinar arrangement. Additionally, some acinar cells exhibited apoptotic vacuoles as a result of parasitism.

3.4 Molecular confirmation of *Leucocytozoon* by molecular methods

In the present investigation, a single, homogenous electrophoretic band of 248 bp was yielded by polymerase chain reaction, resulting from the amplification of the mitochondrial DNA genome (*cytb* gene) within the nuclear ribosomal gene complex. Sequence analysis, depicted in Figure 6, revealed that all DNA sequences were identical to the *cytb* gene of *L. sabrazesi*. The obtained sequence was then deposited in GenBank with the accession number ON399180. BLASTN sequence analysis revealed 100% nucleotide sequence homology with the reference isolates. Additionally, nucleotide sequence homologies of 98.16, 95.85, and 93.55% were reported with sequences of *L. sabrazesi* isolates from chicken in Thailand (MW316432; MW316434) and Malaysia (AB299369), respectively (Figure 6).

4 Discussion

Leucocytozoonosis is a significant parasitic disease affecting avian hosts globally. This disease can lead to severe pathology and economic losses, including decreased meat yield and egg production. Previous literature has documented a limited number of reports on *Leucocytozoon* infection in various avian species in Africa (22, 23), New Zealand (24), and Turkey (25, 26). However, to our knowledge, no previous investigations have explored this parasite among pigeons in Egypt. Therefore, this study is the first to examine *Leucocytozoon* species' prevalence and phenotypic characteristics among Egyptian pigeons.

The findings showed that 12.80% of examined birds were infected with Leucocytozoon species, highlighting the significant challenge of haemoparasitic infections in the studied area. This finding is consistent with the observation of Gocok et al. (26), who noted an infection rate of 13% for Leucocytozoon spp. among Turkish pigeons in Ankara province. However, a substantial disparity in the incidence of Leucocytozoon infection was observed compared to those recorded worldwide. Notably, very low prevalence rates of 2, 2.16, and 6.4% were reported among domestic pigeons from Bangladesh, India, and Nigeria, respectively (27-29). On the contrary, other surveys have reported higher infection rates, such as 30% in Pink pigeons (Columba mayeri) from the island of Mauritius in the Indian Ocean (30), 20% in pigeons from the Mymensingh district in Bangladesh (6), and 25% in various bird species from Europe, Africa, and North America (12). The significant fluctuations observed among the prevalence rates could be attributed to various factors, including differences in geographical locations, climatic conditions, bird breeds, management practices, sample sizes, detection methods employed, the presence of vectors, and study design (31-34).



A phylogram of the maximum likelihood analysis of the *cytb* gene sequences of the *Leucocytozoon* spp. infecting poultry, depicting the robust association of the currently identified isolates of *Leucocytozoon sabrazesi* from pigeons marked by black circles (ON399180). The Maximum Likelihood Tree was implemented in MEGA software version 11 using *Eimeria acervulina* as outgroup. Bootstrap confident values were calculated with 500 repetitions.

The current investigation showed a significantly higher incidence of infection in adult birds than in young squabs, corroborating the findings of Garvin and Greiner (35). The same observation was documented in previous reports, revealing a significantly higher occurrence of avian haemoparasites in adults than grower birds in Bangladesh and Ethiopia (36, 37). However, our findings contradict previous reports from Kenya and Pakistan, which indicated that grower birds had a higher prevalence of avian haemoparasites than older birds (38, 39). Likewise, Van Oers et al. (40) and Castro et al. (41) observed a noticeably higher prevalence in young birds than in other age categories. On the other hand, other surveys concerned with blood parasites in various avian species have not found any significant association between the infection rate and host age (41-44). The positive correlation between the prevalence and age of screened birds could be accounted for higher mortality rates among young birds (2), declined immune response of adults (45), or increased exposure to the vectors (9).

The present results showed that the prevalence of *Leucoytozoon* spp. is influenced by sex, with a significantly higher infection rate in males (33.33%) than females (10.43%). This observation was supported by various previous reports (6, 46, 47). This observation is consistent with previous discussions by several authors who have suggested that higher testosterone levels and factors like stress during the courtship period play a significant role in immunosuppression in males, rendering them more susceptible to infection (48, 49). In contrast, Krone et al. (50) and Nath et al. (27) suggested that the

highest peak of *Leucocytozoon* spp. prevalence rate is in female birds as compared to male pigeons.

According to the present findings, the incidence of Leucocytozoon parasite infection in pigeons in winter was the lowest (0%), and the difference between the winter and other seasons was statistically significant. Similarly, Nath and Bhuiyan (51) in Bangladesh demonstrated that the incidence of Leucocytozoon infections in pigeons during the summer was 60.6%, significantly lower in other seasons. Additionally, other studies on weaver birds of South Africa (52) and rock pigeons in India by Gupta et al. (53) were in the same line and documented that the summer season had the highest peak as compared to other seasons. However, the present finding disagrees with Senlik et al. (54), who demonstrated that the highest infection rate was recorded in the autumn season (44%), while the lowest rate was observed in the spring season in Iran. Moreover, Lawal et al. (55) revealed that the highest infection rates of haemoparasites, including Leucocytozoon spp., occurred during the rainy season (39.3%), followed by the cold dry (12.5%) season and the hot, dry season (7.7%). The potential explanation for the higher prevalence of Leucocytozoon in the dry season could be attributed to the warm climatic conditions that support the abundance of vector, Simuliid blackflies, which are widely distributed throughout the surrounding environment during this time of year (56-58).

In the current study, histopathological exploration revealed various lesions caused by *Leucocytozoon* spp. in the liver, spleen, and pancreas, including necrotic foci and loss of normal organization. Numerous megaloschizonts infiltrated the interstitial spaces, some displaying intact capsules with nuclei while others showed signs of degeneration. Surrounding the distribution of schizonts, a lymphocytic reaction with eosinophilic structures was observed, indicating a host defense response. Fatty degeneration was evident in the liver, indicating *Leucocytozoon* infection, pronounced hemosiderosis, lymphoid depletion in the spleen, and depletion of zymogen granules in the pancreas. Additionally, vascular congestion and thrombosis were diagnosed in some cases. Histopathology proved to be a definitive diagnostic tool for *Leucocytozoon* tissue reaction, with numerous megaloschizonts detected in highly vascularized organs such as the pancreas, lungs, pectoral muscles, liver, spleen, and heart. Our findings align with previously reported worldwide studies (59–61).

It should be noted that one of the most characteristic pathological lesions associated with *Leucocytozoon* infections includes fatty degeneration, which may sometimes be mistaken for fatty liver haemorrhagic syndrome. However, in cases of *Leucocytozoon* infection, distinct megaloschizonts are observed in various organs (62). Additionally, microscopic examination of spleens invaded by *Leucocytozoon* documented chronic inflammation characterized by aggregations of mononuclear cells and disorganized tissue with unclear boundaries between splenic pulps, resulting from merozoite invasion of erythrocytes (63). Furthermore, *Leucocytozoon* spp. colonization, blockage, and thrombosis lead to multiple focal areas of necrosis and ischemia, followed by cardio-respiratory failure and death (64).

Interestingly, our study marks the first molecular characterization of Leucocytozoon parasites in domestic pigeons from Egypt, shedding light on the presence of this parasite in avian hosts and expanding our understanding of blood parasites infecting Egyptian pigeons, addressing gaps in their phylogeny. Through sequencing analysis, we identified Leucocytozoon sabrazesi in pigeons. Furthermore, the isolate recovered in our study exhibited a 98.62% similarity to reference sequences (MW316431) previously identified in chickens. Similarly, research by Chawengkirttikul et al. (3) indicated a low diversity of L. sabrazesi populations in Thailand, with similarity values ranging from 89.5 to 100% with sequences from Malaysia and Myanmar. Additionally, genetic diversity studies of Leucocytozoon sp. based on cytb gene sequences have been conducted in various countries (7, 10, 23, 65–68), highlighting the *cytb* gene's utility as an effective marker for phylogenetic taxonomy on a large scale and as a valuable tool for epidemiological analysis of leucocytozoonosis. Another survey in Egypt (11) reported the presence of L. caulleryi in broiler chicken flocks for the first time, showing a 99.14% similarity to strains recovered from Asian isolates in India, Japan, Malaysia, South Korea, Taiwan, and Thailand.

5 Conclusion

The current investigation marks the inaugural molecular study of haemosporidian parasites in pigeons in Egypt. Notably, our study stands as the pioneering genetic characterization of *L. sabrazesi* infection among pigeons, marking a national and global milestone. Additionally, our study has uncovered a significant statistical association between the infection prevalence of the parasite and various potential epidemiological factors, such as the age and sex of screened birds, with notable seasonal fluctuations observed throughout the year. Further studies are suggested to explore potential vectors at the national level, aiming to identify optimal preventive and therapeutic strategies against leucocytozoonosis, thereby mitigating or eradicating its detrimental impact on the bird industry. Additional large-scale surveys about the occurrence of the parasite within the avifauna of other regions in Egypt could provide valuable insights into blood parasite–host relationships and distribution patterns.

Data availability statement

The original contributions presented in the study are included in the article/supplementary material, further inquiries can be directed to the corresponding author/s.

Ethics statement

The animal studies were approved by the Ethics Committee, Faculty of Veterinary Medicine, South Valley University, Egypt according to the ethical regulations and guidelines for using animals in research (under permit code No. 84). Written and oral informed consent was obtained from the owners for the participation of their animals in this study. The studies were conducted in accordance with the local legislation and institutional requirements. Written informed consent was obtained from the owners for the participation of their animals in this study.

Author contributions

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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