ASSESSMENT OF NEUROINFLAMMATORY AND NEURODEGENERATIVE ALTERATIONS IN THE BRAIN IN A MOUSE MODEL OF VISCERAL LARVA MIGRANS

BY

DINA I. ELGENDY^{1*}, AHMAD A. OTHMAN¹, EMAN M. SAIED², AND HAGER S. ZOGHROBAN¹

Department of Medical Parasitology, Faculty of Medicine, Tanta University¹ and Department of Pathology, Faculty of Medicine, Kafrelsheikh University², Egypt (*Correspondence: dina.elgendy@med.tanta.edu.eg

- ORCID: https://orcid.org/0000-0002-0070-6113)

Abstract

Neurotoxocariasis is a parasitic zoonosis resulting from the intrusion of the larvae of dog roundworm *Toxocara canis* into the central nervous system (CNS), leading to severe CNS insults, and provoking different neurological manifestations ranging from headache to convulsions and paresis. This work assessed the probable occurrence of neuroinflammatory and neuro-degenerative alterations in the brain tissues of *T. canis* infected mice and evaluated the changes in brain parasite loads over the course of infection. The total larval loads in the brain, histopathological, immunohistochemical (MMP-3 & MMP-9) examination as well as determination of NF- κ B and Tau protein levels in the brain tissues of mice, expressions of all the investigated markers of neuroinflammation and neurodegeneration showed an ascending pattern throughout the course of the infection and were directly proportional to the increase in the brain larval loads. **Keywords:** Neurotoxocariasis, Neuroinflammation, Neurodegeneration, MMP-3, MMP-9, NF- κ B, Tau protein.

Introduction

The chief causative agent of visceral larva migrans is the common nematode parasite of dogs: Toxocara canis (T. canis). It affects a large scale of animal hosts as well as humuns (Castillo et al, 2000). Studies on seroprevalance rates of human toxocariasis reported that it is one of the most prevalent helminthes worldwide (Abdi et al, 2014; Ebrahimifard et al, 2015; Wang et al, 2020). Its prevalence rates ranged from 2-5% in urban areas to 35-42% in rural areas (Magnaval et al, 2001; Rubinsky-Elefant et al, 2010; Shokouhi et al, 2018). Human infection took place by ingestion of the embryonated T. canis eggs containing third-stage larvae in contaminated soil, foodstuffs or water and by ingestion of raw or improperly cooked tissues of various paratenic hosts containing infective viable larvae (Strube et al, 2013). Upon liberation of the infective larvae in the gastrointestinal tract, they penetrated the intestinal mucosa and were carried by the circulation to diverse parts of the body. At about the 7th day post infection (P.I.), they

entered the CNS (Abo-Shehada and Herbert, 1984). The larvae produced a variety of injurious substances including enzymes, excretory substances, and cuticular structures, which stimulated remarkable local immune responses that resulted in destruction of the affected tissues (Xinou *et al*, 2003).

Invasion of the CNS by *T. canis* larvae results in the neurotoxocariasis (Strube *et al*, 2013). In humans, this condition may be asymptomatic or it may lead to severe CNS insults such as eosinophilic meningitis, encephalitis, myelitis and cerebral vasculitis (Dousset *et al*, 2003; Sánchez *et al*, 2018; Morsy, 2020). Patients may suffer from different neurological manifestations including principally headaches, convulsions, paresis and behavioral disorders (Fan *et al*, 2015).

Furthermore, a correlation between *Toxo-cara* infection and cognitive decline, learning disabilities and memory disorders had been evidenced (Scheid *et al*, 2008; Walsh and Haseeb, 2012; Erickson *et al*, 2015).

Neurotoxocariasis is still considered as a poorly understood disorder. Moreover, the

underlying mechanisms by which *T. canis* larvae could invade and induce damage in the brain tissues of the affected host remain particularly deficient (Waindok and Strube, 2019). Besides, data were accumulating in the literature about probable link between neurotoxocariasis and a number of neurode-generative diseases such as idiopathic Parkinson's disease, Alzheimer's disease and depression (Fan *et al*, 2015; Janecek *et al*, 2017).

The present study aimed to explore the potential development of neuroinflammatory and neurodegenerative alterations in the brain tissues of *T. canis* infected mice and to assess the correlation between these changes and the brain parasite loads over the course of infection.

Material and Methods

Parasite: *T. canis* eggs, obtained by flotation technique from stools of puppies, were cultured in $0.1N H_2SO_4$ for embryonation and storage. Infections were done using eggs from culture not more than two months old (Kayes and Oaks, 1976). Prior to infection of mice, the eggs were washed for three times in physiological saline to eliminate the physiological saline/ H_2SO_4 solution. Eggs were then re-suspended in distilled water and their concentration was adjusted to be 1000 embryonated eggs per 0.25ml. 1000 embryonated eggs were administered to each mouse in infected group by intragastric inoculation (Chou *et al*, 2017).

Animals and experimental design: Laboratory bred male Swiss albino mice (6-8weeks old and 20-25g in weight) were used in this work. Mice were housed and fed according to the national and institutional guidelines for the care and use of laboratory animals. A total of 45 mice were divided into 2 groups: G1:15 mice, non-infected mice as a negative control group; and G2: 30 mice, *T. canis*infected mice. Collection of samples was done at the end of weeks 3, 7, & 15 P.I. Ten mice from the infected group and 5 mice from the negative control group were sacrificed at each time point P.I. and divided as follows: Five mice from infected group for brain larval counting, and 5 mice from every one of the 2 groups, brains were divided into two parts one for histopathological and immunohistochemical examinations and the second for biochemical assays.

Total larval loads in the brain: The infected animals were sacrificed, and the whole brain from each mouse was retrieved in a clean Petri dish. The two halves of the cerebrum and the cerebellum were detached and each half was squashed between two microscopic slides followed by its examination under the low power of a light microscope to estimate total brain larval counts (Kayes and Oaks, 1976).

Histopathological study: Brain tissue samples from the studied groups were fixed by immersion in 10% formalin at once, and then routine histological processing was done. They were embedded in paraffin followed by microtomy and staining with hematoxylin & eosin to observe histopathological changes in the brain of the infected mice group at different studied time points P.I.

Immunohistochemical staining for determination of matrix metalloproteinases (MMP-3 & MMP-9) expression was done according to the manufacturer's protocol using the Ultra-Vision Detection Kit (TP-015-HD, Lab Vision, USA), rabbit polyclonal anti-MMP-3 antibody (Cat. No. PA5-27936, Lab Vision, USA) in 1:100 dilution and rabbit polyclonal anti-MMP-9 (Cat. No. ab38898: abcam, USA).

Evaluation of MMP-3 & MMP-9 was by immunohistochemical staining (Hou *et al*, 2012). Positive immunostaining was distinguished as brown granular cytoplasmic staining. The immunostaining of both antibodies was scored as follows: percentage of each antibody stained area in five rand-om images per mouse were determined & analyzed by Image J software (Java image processing program inspired by NIH) and mean was calculated for each group.

Biochemical study: determination of NF- κB (nuclear factor kappa-light-chain-enhan-

cer of activated B cells) and Tau protein (a microtubule-associated protein, mainly expressed in the neuronal cells) levels in brain tissue homogenates was done using ELISA kits for quantitative detection of mouse NFκB (Catalog no. MBS2023542: MyBio-Source, USA) & Tau protein (Catalog no. LSF-6386: Life-Span Bioscience, USA) following the manufacturer's protocol.

Statistical analysis: Data were presented as M±SD. Kruskal-Wallis test was used to compare all the experimental groups followed by post-hoc test to detect significance between groups. Differences were considered significant when (P < 0.05). Analyses were done using Statistical Program of Social Sciences (SPSS Inc., Chicago, Illinois, USA), software for windows, version (20).

Ethics statement: The study protocol was approved and conducted according to the guidelines of the Laboratory Animal Centre for Research Ethics Committee, Faculty of Medicine, Tanta University (Approval code: 33557/12/19).

Results

Brain larval loads: T. canis larval counts in the brain tissues of the infected mice at different time points P.I. (Tab. 1). Larval counts showed an ascending pattern, denoting progressive accumulation of T. canis larvae in the brain. Almost all larvae showed signs

of viability on close observation. In sum, there was a significant increment of larval counts in brain as the infection proceeded.

Histopathological examination of brain sections of infected mice revealed the T. canis larvae that were not surrounded by any inflammatory cell infiltration. Similar observations were detected in all examined brain sections of the infected group at all studied time points P.I. (Fig. 1).

Immunohistochemical assessment of the MMP-3 & MMP-9 reactivity: The expressions of MMP-3 & MMP-9 (Fig. 2 & 3) in brain tissues of infected mice were significantly higher than expressions in negative control group. Expressions of both markers increased significantly with increasing infection duration. They were detected as early as the 3rd week P.I. and showed an ascending pattern throughout the course of infection (Tab. 2).

Biochemical results: A significant increase in NF- κ B levels in brain tissue homogenates was detected in T. canis infected mice compared to the negative control group. Likewise, Tau protein levels were significantly upregulated in the infected group compared to the negative control group. Levels of both markers showed a significant elevation over the course of the infection reaching the highest levels at the 15th week P.I. (Tab. 3).

Table 1: Larval loads in brain of infected mice at different time points P.I. (n=5)					
Group	3 weeks P.I.	7 weeks P.I.	15 weeks P.I.		
G2	74.50 ± 5.15	106.50 ± 7.52	161.70 ± 5.46		
<i>P1</i> <0.001*, <i>P2</i> <0.001*, <i>P3</i> <0.001*					

* significante P<0.05; P1 compared between larval count at 3 & 7 weeks P.I., P2 compared between larval count at 3 & 15 weeks P.I., P3 compared between larval count at 7 & 15 weeks P.I.

Groups	Percentage of MMP-3 stained area	Percentage of MMP-9 stained area
Groups	Mean \pm SD	Mean \pm SD
G1	2.96 ± 0.25	3.06 ± 0.24
G2 (3 weeks P.I.)	18.02 ± 0.28	22.35 ± 0.57
G2 (7 weeks P.I.)	45.12 ± 2.03	50.45 ± 0.55
G2 (15 weeks P.I.)	81.71 ± 1.44	83.52 ± 0.71
P value (K W test)	<0.001*	<0.001*
P value (multiple	<i>P1</i> < 0.001*, <i>P2</i> < 0.001*, <i>P3</i> < 0.001*, <i>P4</i> <	<i>P1</i> < 0.001*, <i>P2</i> < 0.001*, <i>P3</i> < 0.001*
comparison)	$0.001*P_{5} < 0.001*P_{6} < 0.001*$	P4 < 0.001*, P5 < 0.001*, P6 < 0.001*

Table 2: Comparison of MMP-3 & MMP-9 expressions in brain sections of all studied groups (n=5)

* Significant P<0.05 Pl compared between G1 & G2 (3 weeks P.I.), P2 compared between G1 & G2 (7 weeks P.I.), P3 compared between G1 & G2 (15 weeks P.I.), P4 compared between G2 (3 weeks P.I.) & G2 (7 weeks P.I.), P5 compared between G2 (3 weeks P.I.) & G2 (15 weeks P.I.), P6 compared between G2 (7 weeks P.I.) & G2 (15 weeks P.I.)

Tuble 5. Comparison of 11 KB & Tau protein levels in orall disde nonogenates of an groups (in 5)		
Groups	NF-κB ng/ml	Tau protein ng/mL
G1	0.386 ± 0.019	0.69 ± 0.13
G2 (3 weeks P.I.)	3.798 ± 0.406	2.15 ± 0.26
G2 (7 weeks P.I.)	7.031 ± 0.275	6.09 ± 0.17
G2 (15 weeks P.I.)	10.987 ± 0.264	9.22 ± 0.51
F test	109.354	85.214
P value (multiple comparison)	$\begin{array}{l} P1 < 0.001^*, \ P2 < 0.001^*, \ P3 < 0.001^*, \\ P4 < 0.001^*, \ P5 < 0.001^*, \ P6 < 0.001^* \end{array}$	P1 < 0.001*, P2 < 0.001*, P3 < 0.001*, P4 < 0.001*, P5 < 0.001*, P6 < 0.001*

Table 3: Comparison of NF-κB & Tau protein levels in brain tissue homogenates of all groups (n=5)

Discussion

Some parasites possess a predilection to gain access to the CNS (Bencurova et al, 2011). T. canis larva is one of these parasites that invade the CNS, provoking a spectrum of manifestations in the affected humans and animals. Up till now, the mechanisms of invasion and injury of the brain parenchyma by T. canis larvae were not fully elucidated (Waindok and Strube, 2019). In this context, certain molecules may facilitate the larval movement and migration as well as the induction of pathological changes in the CNS. That's why, this study investigated the effect of T. canis infection on the expression of different markers including MMP-3, MMP-9, NF-kB and Tau protein which are highly involved in the pathogenesis of traumatic brain injury, neuroinflammation and neurodegenerative disorders (Liao et al, 2008; Sabbagh et al, 2013), suggesting that they may have a role in the pathogenesis of neurotoxocariasis.

In the present study, the brain larval loads showed an ascending pattern in the accumulation of larvae in the infected animal's brain tissues over the infection course. These results agreed with Lai *et al.* (2005) and Chou *et al.* (2017) who reported a steady increment of larval loads in the brain with time. This progressive accumulation of the larvae in brain could be due to the favorable environment for the larvae survive as it is an immune-privileged site where the larvae become protected from the host immune response (Holland and Hamilton, 2013).

The absence of inflammatory cell infiltration in the brains of *T. canis* infected mice that had been detected in the present work is confirmative to the findings of Liao *et al.* (2008), Eid *et al.* (2015) and Chou *et al.* (2017) on *T. canis*-infected outbred mouse strains. But, Springer *et al.* (2019) detected perivascular cuffs with eosinophils and neutrophils in C57Bl/6 mice infected with *T. canis.* This could be explained by the difference between the immune response of inbred and outbred mice to *T. canis* infection.

This work showed that the infection with T. canis resulted in a significant upregulation of MMP-3 expression in the brain tis-sues of the infected mice. In addition, its expression increased progressively over the course of the infection and it was directly proportional to the larval counts in the brain tissues. To date and to the best of our knowledge, no data are available in the literature on the expression of MMP-3 in the brains of T. canisinfected mice. Interestingly, it was found that its expression increased in response to other parasitic infections such as Plasmodium falciparum infection (D'Alessandro et al, 2013). MMP-3 is able to break down numerous extracellular matrix proteins, trigger growth factors and degrade cell adhesion particles, chemokines, cytokines and a mixture of receptors. Also, it activates MMP-9 promoter (Ogata et al, 1992; Arza et al, 2000; McCawley and Matrisian, 2001). It could disrupt the blood-brain barrier, promote demyelination and apoptosis, and induce additional inflammatory responses.

Therefore, the over-expression of MMP-3 supports neuroinflammation and apoptosis and leads to exacerbation of numerous neurodegenerative diseases (Kim *et al*, 2010; Cauwe and Opdenakker, 2010).

In the current work, the expression of MMP-9 in the brain tissues of the infected animals significantly increased in comparison to the negative control animals and it

was proportional to the larval recovery rate in the brain tissues. These results were similar to those of Lai *et al.* (2005) who reported that activity of MMP-9 increased in the lungs, liver, muscles, and brain tissues during *T. canis* larval migration. Besides, Shyu *et al.* (2019) demonstrated that MMP-9 had a major job in granuloma formation during the progression of ocular toxocariasis via reinforcement of inflammatory cell infiltration and degradation of extracellular matrix proteins.

MMP-9 owns a noteworthy role in the pathogenesis of neuroinflammatory diseases (Khuth et al, 2001; Yong et al, 2001; Rosenberg, 2002). A consistent increment in its levels in the cerebrospinal fluid is considered as an indicator of neuroinflammation over course of various diseases influencing the CNS, for example, multiple sclerosis as well as bacterial and viral meningitis (Giraudon et al, 1998; Leppert et al, 1998; Sporer et al, 1998; Matsuura et al, 2000; Yushchenko et al, 2000). Also, a similar perception was recognized on investigating the cerebrospinal fluid of animals affected with infectious or inflammatory conditions of the CNS (Gijbels et al, 1993; Azeh et al, 1998; Paul et al, 1998). Besides, the critical role of MMP-9 in synaptic plasticity, learning, and memory were brought forward by many authors (Szklarczyk et al, 2002; Meighan et al, 2006; Nagy et al, 2006; Okulski et al, 2007; Tian et al, 2007).

Indeed, the excessive expression of the MMPs, especially MMP-3 & MMP-9, was implicated in several conditions of acute brain insult including hemorrhagic, ischemic and traumatic lesions (Kim *et al*, 2010; Heo *et al*, 1999; Wang *et al*, 2000). Moreover, a strong evidence about their contribution in the pathology of neurodegenerative diseases such as multiple sclerosis and Alzheimer's disease is well established (Hartung and Kieseier, 2000; Lorenzo *et al*, 2003). Additionally, in the last decade, several studies supported the role of MMPs in the pathogenesis of some neurological disorders caused

by the parasitic infections either protozoal (Clark *et al*, 2010; Thibeaux *et al*, 2014; Jacintho *et al*, 2018) or helminthic (Verma *et al*, 2011; Wei *et al*, 2011; Bruschi *et al*, 2014).

NF- κ B is a common constituent in nearly all categories of cells in the brain (O'Neill and Kaltschmidt, 1997). Its activation and increased expression within astrocytes caused significantly amplify neuronal degeneration (Mattson and Camandola, 2001; Johnstone et al, 2013). Moreover, increased expression of NF-kB was reported in diverse acute and chronic neurodegenerative disorders such as ischemic brain injury (Zhou et al, 2018), Traumatic brain injury (Chen et al, 2017), Alzheimer's disease (Chen et al, 2005) and Parkinson's disease (Mogi et al, 2007). The present work showed that there was upregulation in the levels of NF- κ B in the brains of T. canis infected mice over the course of infection. These results suggest a role of NF-kB in the pathogenesis of neurotoxocariasis. Previous studies on the NF-kB signaling pathway showed that there is a link between NF-kB activation and the induction of MMP-9 gene expression. This could be a direct effect by binding to the NF-kB element in the MMP-9 promoter or an indirect effect by triggering other transcription factors that have the capacity to stimulate MMP-9 gene expression (Brown et al, 1995; Fujioka et al, 2004; Gordon et al, 2009).

The main protective barrier against the passage of various injurious elements including pathogens into the CNS is the bloodbrain barrier (BBB). The mechanisms by which the neurotropic parasites manipulate the BBB differ according to the extracellular or intracellular nature of these parasites. In contrast to intracellular parasites, the mechanisms of disruption or interaction with the BBB were largely unknown for extracellular parasites including *T. canis* (Masocha and Kristensson, 2012). However, metalloproteases were found to be crucial for crossing the BBB during the infection by the extracellular parasite *Balamuthia mandarillaris* (Matin *et al*, 2006). Also, other parasites reaching the CNS, various cytokines, chemokines, and adhesion molecules- all with close interaction with various MMPs were implicated in the interaction with BBB (Masocha and Kristensson, 2012). So, this study may help explain the cross-talk between Toxocara larvae and the BBB. This context suggested a mechanism for the process of CNS invasion and migration by T. canis larvae. It assumed that activation of MMP-9 & NF-KB results in an increase in BBB permeability and disruption of tight junction proteins, facilitating the pa-ssage and movement of the larvae through the tissues. The findings of Chiu and Lai (2013) who reported that BBB dysfunction in angiostrongyloidosis was mediated by MMP-9 via the NF-KB/MMP-9 signaling pathway are support for these explanations. However, whether T. canis larvae exclusively provokes the expression of host metalloproteases or produces its proper metalloproteases needs further research.

Tau protein is an essential physical component of neurons that has an important role in assembly and stabilization of microtubules. Alterations in the distribution of tau protein in neurons and its accretion in the form of insoluble fibrillary deposits are highly consistent with death of neuronal cell. So, it was considered as an indicator of neurodegeneration (Williams, 2006; Wang et al, 2012). The expression of tau protein was significantly elevated with time in brain tissues of T. canis infected mice compared to negative controls. This agreed with Liao et al. (2008) and gave evidence on the occurrence of progressive neurodegenerative changes in brain because of T. canis infection.

Conclusion

The results reported that *T. canis* infection induced the production of MMP-9 and its activator MMP-3 as well as NF-B and Tau protein in the brain tissues of infected mice. This could provide insights on the contribution of these molecules in the pathogenesis of neurotoxocariasis through the processes of larval invasion and migration within the CNS as well as the induction of inflammation and neurodegeneration. The molecules can also be useful as markers of inflammation during neurotoxocariasis. Also, correlation could be between NF- κ B/ MMP-9 signaling pathway and ability of *T. canis* larvae to invade and migrate within CNS. Thus, MMPs or their inhibitors have potential targets for dealing with diagnosis, treatment, and control of neurotoxocariasis and possibly of other CNS disorders as well.

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Explanation of figures

Fig. 1: A brain section of *T. canis*-infected mouse 7 weeks P.I. showed a cut section in *T. canis* larva (yellow arrow) without inflammatory cell infiltration around it (H&E \times 400).

Fig. 2: Brain sections of negative control and *T. canis*-infected mice showed immunohistochemical expression of MMP-3 within glial cells (Immunoperoxidase × 400). A- Control group showed negative MMP-3 expression, B- Infected group 3 weeks P.I. showed weak increase in

percentage of MMP-3 stained area, C- Infected group 7 weeks P.I. showed moderate increase in percentage of MMP-3 stained area, D-Infected group 15 weeks P.I. showed marked increase in percentage of MMP-3 stained area. Fig. 3: Brain sections of negative control and *T. canis*-infected mice showed immunohistochemical expression of MMP-9 within glial cells (Immunoperoxidase × 400): A- Control group showed negative MMP-9 expression, B- Infected group 3 weeks P.I. showed weak increase in percentage of MMP-9 stained area, C- Infected group 7 weeks P.I. showed moderate increase in percentage of MMP-9 stained area, D-Infected group 15 weeks P.I. showed marked increase in percentage of MMP-9 stained area.

