Role of heme oxygenase-1, cytokines, and vascular endothelial growth factor in murine *Schistosoma mansoni*

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ABSTRACT

**Objectives:** Among tropical diseases, schistosomiasis caused by *Schistosoma mansoni* is the second major cause of morbidity and mortality worldwide. Inflammation was considered as an adverse event that contributes to the pathology associated with schistosomiasis. Heme oxygenase-1 (HO-1) and vascular endothelial growth factor (VEGF) have been implicated in the process of angiogenesis. The current study aimed to evaluate the effect of *S. mansoni* infection on HO-1 gene expression, IL-4, IL-12, and VEGF to address the role of these factors in the pathogenesis of schistosomiasis.

**Methods:** Thirty mice divided equally into three groups comprised a non-infected control group and two *S. mansoni*-infected groups. Infected animals were studied at 8 and 12 weeks post-infection. Serum IL-4, IL-12, and VEGF were measured. HO-1 mRNA was detected by RT-PCR of liver homogenates and HO activity was assessed as percentage of carboxy hemoglobin.

**Results:** *S. mansoni*-infected mice showed a progressive increase in serum IL-4 and VEGF and decrease in IL-12 levels. In addition, HO-1 expression and activity were increased in infected mice compared to control group with the maximum increase at egg deposition stage.

**Conclusion:** Our results suggested that the body response to acute stage of *S. mansoni* infection by elevating the expression of the stress gene HO-1 and that VEGF may serve as a new indicator of progression of *S. mansoni* associated angiogenesis which regulates granuloma and/or fibrosis development in the liver of infected mice. Understanding the role of HO-1 and VEGF in pathogenesis of *S. mansoni* may provide a new pharmacological target.

**Keywords:** HO-1 gene expression, IL-12, IL-4, *Schistosoma mansoni*, vascular endothelial growth factor

**Introduction**

Schistosomes are members of a medically important group of parasitic helminthes that contribute to severe morbidity and mortality among people in 74 tropical and subtropical developing nations.[1] *Schistosoma mansoni* infection is considered a worldwide problem that affects large geographic areas in several countries situated in Africa, Madagascar, the Middle East, parts of South America, and the Caribbean region.[2] In Egypt, the prevalence of *S. mansoni* in five endemic governorates in the lower Egypt ranged from 17.5% to 42.9% with an average of 36.4%.[3]

Heme oxygenase (HO) is the rate-limiting enzyme in the conversion of heme into biliverdin, carbon monoxide (CO), and free iron (Fe²⁺).[4] Three HO isoforms have been identified; they are inducible HO-1, also known as heat shock protein 32, constitutively expressed HO-2 and a related but less well characterized HO-3.[5] Under physiologic conditions, HO-1 expression is relatively low. Upregulation of HO-1 may be among the most critical cytoprotective mechanisms that are activated during cellular stress and inflammation.[6] HO-1 is thought to play a key role as an anti-inflammatory gene.[4,5] Moreover, HO-1 has been implicated in the process of angiogenesis.[6,7]

Endothelial cells play a fundamental role in the pathogenesis of schistosomiasis as well as in granuloma formation.[8] In addition, *in vitro* studies have revealed that soluble schistosomal egg antigens upregulate vascular endothelial growth factor (VEGF) and angiogenesis.[9] Since angiogenesis appears to be a key factor for new connective tissue synthesis, assay of VEGF levels in schistosomiasis seemed to be appropriate.
Cytokines play a crucial role in the evolution and regulation of schistosoma-induced immunopathology. T lymphocytes are a major source of cytokines. These cells bear antigen specific receptors on their cell surface to allow recognition of foreign pathogens. There are two main subsets of T lymphocytes, CD4 and CD8. T lymphocytes expressing CD4 are also known as helper T cells, and these are considered the most prolific cytokine producers. T helper (Th) cells can be further subdivided into Th1 and Th2, and the cytokines they produce are known as Th1-type cytokines and Th2-type cytokines. Th1-type cytokines tend to produce the pro-inflammatory responses responsible for killing intracellular parasites and for perpetuating autoimmune responses. Excessive pro-inflammatory responses can lead to uncontrolled tissue damage, so a counteract mechanism is required which include Th2-type cytokines. In excess, Th2 responses will counteract the Th1 mediated microbicidal action.\(^{[10]}\) The immune response to schistosomal antigens manifest a shifting from a moderate Th1 to a robust Th2-dominant response.\(^{[11]}\) Interleukin (IL)-4 plays a key role in the Th2-associated immune response, on the other hand, IL-12 induces Th1 response.\(^{[12]}\)

The aim of the current study was to evaluate the effect of schistosomiasis caused by murine \(S. \text{mansoni}\) on HO-1 gene expression, IL-4, IL-12, and VEGF to address the role of these factors in granuloma formation.

**Material and Methods**

**Animals and experimental design**

\(S. \text{mansoni}\) cercariae (Egyptian strain) were obtained from infected \(Biomphalaria alexandrina\) snails which were purchased from Theodor Bilharz Research Institute (TBRI), Giza, Egypt and used for mice infection using tail immersion technique by 100 cercariae/mouse.\(^{[13]}\) The study includes thirty males, Swiss albino mice aged 8–12 weeks old and weighting 20–25 g. Mice were housed in an animal house at Faculty of Pharmacy, Suez Canal University and were allowed to free access of food and water. Experimental animals were kept and used in accordance with the Guide for the Care and Use of Laboratory Animals (National Research Council, 2011). All experimental protocols were approved by the Ethics Committee at the Faculty of Pharmacy, Suez Canal University (Ismailia, Egypt) (code # 201906RA1). The study was continued for 6 months at the time from September 2019 to February 2020.

Mice were divided into three groups of ten mice each. The three groups comprised non-infected mice (control group) and two \(S. \text{mansoni}\)-infected groups (100 cercariae/mouse). Infected animals were studied at 8th (acute phase) and 12th (chronic phase) weeks post-infection.\(^{[14]}\) At the end of the study, serum was separated for determination of IL-4, IL-12, and VEGF. Mice were sacrificed and part of the liver was removed, rapidly frozen in liquid nitrogen and used for detection of HO-1 mRNA and HO-1 activity and another part was fixed in 10% formalin for histopathological examination and immunohistochemical staining of VEGF.

**Cytokines and VEGF determination**

IL-4, IL-12, and VEGF were measured in the serum of control and infected mice groups by enzyme-linked immunosorbtant assay (ELISA) technique according to the manufacturer’s instructions.

**RT-PCR for detection of HO-1 mRNA**

Total RNA was isolated from liver tissues using RNA extraction kit (Qiagen, Germany). Concentration of the extracted RNA was measured by NanoDrop ND-1000 (NanoDrop Tech., Wilmington, USA). Specific primers for mice HO-1 cDNA fragment: primer 1: 5'-AAC ACA AAG ACC AGA GTG CCT CAC-3’, primer 2: 5'-CAA GAG AAG AGA GCC AGG CAA GAT-3’ and G3PDH primers sense: 5'-CTG CCA TTT GCA GTG GCA AAG TGG-3’, antisense: 5'-TTG TCA TGG ATG ACC TTG GCC AGG-3’ were used.\(^{[15]}\) PCR was performed using one step RT-PCR kit (Qiagen, Germany). Cycling conditions for amplifying RT products were as follows: 95°C, 1 min; 57°C, 1 min; 72°C, 1 min, for 28 cycles, and then extended at 72°C for another 5 min.\(^{[15]}\) After amplification, PCR products were electrophoresed on 1.2% agarose gel, stained with ethidium bromide and visualized under UV light. The PCR products were then quantitated by using a quantitation kit (Promega Corporation, Madison, WI, USA). The results were expressed as μg/mg wet tissue.

**Assay of HO activity**

Carboxy hemoglobin (COHb) percentage of the arterial blood was performed gasometrically by Blood Gas Analyzer (Bayer Rapid lab 865, USA) according to the method described by Hampson \textit{et al.} \(^{[16]}\)

**Histology and immunohistochemistry**

The part of liver tissue fixed in 10% formalin was used. Representative tissue biopsies were selected, processed, and embedded in paraffin blocks. Five tissue sections (4 μm thick) were cut; each was 250 μm away from the preceding section to avoid assessment of the same granuloma. Tissue sections were deparaffinized and stained for Hematoxylin and Eosin (H&E) and Masson Trichrome.\(^{[17]}\) For each mouse, the hepatic histopathological findings were recorded and the degree of portal fibrosis was assessed.

Paraffin sections (5 μm thick) were cut on positively charged slides, dewaxed in xylene, and hydrated in descending grades of ethanol. The endogenous peroxidase activity was quenched by incubation in 100% methanol with 3% hydrogen peroxide for 20 min. Antigen retrieval was performed by subjecting the sections in citrate buffer (pH 7.0) for 15 min in microwave at 700 W. Sections were incubated at room temperature in
a humid chamber with primary mouse monoclonal antibody against VEGF (DakoCytomation, Clone VG 1, Code No. M7273) for 30 min. The antibodies were diluted 1:50 in phosphate-buffered saline (PBS). After rinsing in PBS, the sections were incubated at room temperature for 20 min with EnVision™ system (ChemMate Dako). After a further wash in PBS, the slides were incubated with DAB solution (0.01% hydrogen peroxide in 0.05% diaminobenzidine-tetrahydrochloride). Sections were counter stained with Meyer’s hematoxylin and dehydrated in ethanol prior to mounting. Liver sections with the primary antibody replaced with PBS served as negative controls, while colonic cancer sections served as VEGF-positive controls. The liver sections were examined using a Zeiss light microscope (Oberkochen, Germany). VEGF expression sites were examined intralobular in the periportal areas in hepatocytes, Kupffer cells and endothelial cells lining sinusoids and granuloma. Immunohistochemical staining of sections was evaluated without knowledge of data.

Statistical analysis

Data are expressed as mean ± SD. Significance was assessed by Student’s t-test. P < 0.05 was considered statistically significant. Data were processed using the SPSS 17.0 software package (SPSS, Chicago, IL, USA).

Results

HO-1 mRNA expression and HO activity in the studied groups

As seen in Figure 1a, HO-1 mRNA expression was increased in liver homogenates of *S. mansoni*-infected mice compared to normal mice. The increase in HO-1 expression levels were detected in the 8th week post-infection then significantly decreased in the 12th week post-infection but still significantly higher than normal mice [Figure 1b].

HO-1 activity as detected by percentage of COHb was increased in *S. mansoni*-infected mice compared to non-infected control group. This activity is declined at 12th week post-infection compared to the activity at 8th week post-infection [Table 1].

IL-4, IL-12, and VEGF serum levels in the studied groups

Significant elevation of serum IL-4 was observed in *S. mansoni*-infected mice at 8th and 12th weeks post-infection compared to non-infected control mice. However, IL-12 levels were significantly decreased in 8th and 12th weeks post-infection in infected mice compared to non-infected control mice. In addition, VEGF serum levels were significantly increased in

![Figure 1](image-url): (a) Expression of HO-1 mRNA in the studied groups. M; 100 bp marker, lanes 1, 2; *S. mansoni*-infected mice 8 weeks post-infection, lanes 3, 4; *S. mansoni*-infected mice 12 weeks post-infection, lanes 5, 6, 7, 8; non-infected control mice. (b) HO-1 mRNA expression (µg/mg wet tissue) in the studied groups. Values are represented as mean ± SD (n=30). *Significantly different from normal mice at P<0.05. #Significantly different from *S. mansoni*-infected mice 8 weeks post-infection at P<0.05.
S. mansoni-infected mice at 8th and 12th weeks post-infection compared to non-infected control mice [Table 1].

Histological findings

Figure 2 shows histopathological picture of liver tissue from control mice [Figure 2a]. Sections from the liver 8th week post-infection Figure 2b] showed the presence of adult bilharzia worm in hepatic central vein surrounded by mild inflammatory reaction. The rest of liver tissue showed the presence of living bilharzia ova in portal areas surrounded by collections of inflammatory cells (neutrophils, eosinophils, and macrophages), the size of the inflammatory reaction ranged from 1/2 – 1 high power field. Sections from the liver 12th week post-infection [Figure 2c] showed marked inflammatory reaction in hepatic lobules in absence of bilharzia ova forming fibrocellular granulomas, the reaction is also formed of neutrophils, macrophages, and eosinophils. There is also focal infiltration of inflammatory cells around blood vessels.

Figure 3 revealed Masson trichrome staining of mice liver’s. There is no fibrosis in portal areas or in hepatic lobules in non-infected control group [Figure 3a]. Fibrocellular granuloma around bilharzia ova was detected in liver of 8th week post-infection group [Figure 3b], while mild fibrosis in portal areas was detected in liver of 12th week post-infection group [Figure 3c].

Immunohistochemical aspects

As shown in Figure 4, the control non-infected mice were negative for VEGF monoclonal antibodies [Figure 4a]. Immunostaining for VEGF showed moderate staining in endothelial cells lining vessels and sinusoids 12 weeks post-infection [Figure 4c] in comparison to weaker staining 8 weeks post-infection [Figure 4b].

Discussion

Among tropical diseases, S. mansoni is the second major cause of morbidity and mortality worldwide.[19]

HO-1 is a cytoprotective enzyme, the expression of which is highly sensitive to induction by pro-oxidant stimuli including the substrate heme and nitric oxide.[20] This study showed that HO-1 mRNA expression together with HO activity was significantly higher in mice infected with S. mansoni compared to non-infected mice, also HO-1 activity was declined in chronic stage compared to acute stage of schistosomiasis. Our findings were in agreement with the results reported by Aziz et al.[21] The possible explanation of these results may be the fact that iron is used by schistosomes for development and reproduction and the use of iron was abundant in the stage of egg deposition. Roles of this abundant egg-associated iron use include early embryogenesis and stabilization of cross-linked proteins in eggshell formation.[22] In agreement with this explanation, Maines and Senft reported that schistosome adult worm catabolizes hemoglobin to heme which is the powerful inducer of HO-1 gene expression.[23]

VEGF is a major player in angiogenesis. Enhancement of VEGF expression is mediated by inflammatory cytokines.[24]

Table 1: Serum IL-4 (pg/ml), IL-12 (pg/ml), COHb (%), and VEGF in normal and Schistosoma-infected mice groups

| Groups       | Normal mice | Schistosoma-infected mice |
|--------------|-------------|---------------------------|
| IL-4 (pg/ml) | 71.25±14.8  | 196.7±32*                 |
| IL-12 (pg/ml)| 434.33±95.7 | 202±42.5*                 |
| COHb (%)     | 0.79±0.02   | 1.6±0.104*                |
| VEGF (pg/ml) | 36.8±8.48   | 70.12±19.63*              |

Data are represented as mean±SD and analyzed using Student’s t-test. *Significantly different from normal mice at P<0.05. IL-4: Interleukin-4, IL-12: Interleukin-12, COHb: Carboxyhemoglobin, VEGF: Vascular endothelial growth factor

Figure 2: Hematoxyline and Eosin staining section of liver of the studied groups. (a) Liver control mice (H and E, 400×), (b) Histological section of liver of S. mansoni-infected mice (8 weeks), I; (H and E, ×100), II; (H and E, ×400) showed the presence of adult bilharzias warm in hepatic central vein surrounded by mild inflammatory reaction. III; (H and E, ×100), IV; (H and E, ×400) showed the presence of living bilharzia ova in portal areas surrounded by collections of inflammatory cells forming granulomas (neutrophils, eosinophils, and macrophages). The inflammatory reaction is multiple affecting all portal areas and is variable in size. (c) Histological section of liver of S. mansoni-infected mice (12 weeks), I; (H and E, ×100), II; (H and E, ×400) showed marked inflammatory reaction in hepatic lobules forming fibrocellular granulomas, bilharzia ova are fewer in number than that after 8 weeks. The reaction is also formed of neutrophils macrophages and eosinophils. There is also focal infiltration of inflammatory cells around blood vessels.
In addition, HO-1 stimulates VEGF synthesis and cytokines expression.\[^7\] Therefore, the role of VEGF in the pathogenesis of *S. mansoni* infection was investigated in the present study. Serum level of VEGF was significantly raised in mice infected with *S. mansoni* compared to control group and this was agreed with the results of Tawfeek *et al.* and Shariati *et al.*\[^25,26\] Moreover, the level of VEGF was progressively increased in chronic stage at 12th week post-infection and the immunohistochemical staining in endothelial cells of liver biopsies from infected mice confirmed these results. Our findings suggested that VEGF may serve as a new indicator of progression of *S. mansoni* associated angiogenesis which regulates granuloma and/or fibrosis development in the liver of infected mice.\[^25\] In addition, understanding the role of VEGF in pathogenesis of *S. mansoni* may provide a new pharmacological target.

Schistosomiasis is caused by the host reaction to parasite eggs which when trapped in tissues induce fibrotic granulomatous lesions that eventually impair organ function. Schistosome eggs are responsible for the development of Th2 immune response seen in patently infected animals.\[^27\] Th2 immune response is characterized by the increased production of type 2 cytokines IL-4.\[^28\] Our results showed a significant progressive elevation of IL-4 in *S. mansoni* infected mice at 8th and 12th weeks post-infection compared to non-infected control mice and this was agreed with the results reported by Rolot and Dewals.\[^29\]

The role of Th2 response manifested by elevated IL-4 during schistosoma remain contradictory. IL-4 has been determined as necessary factor for resistance to *S. mansoni* super infection in mice.\[^30\] Other studies, instead has viewed Th2 response as deleterious mediating sever pathology and promoting worm survival and fecundity.\[^31\] This may explain the elevation of IL-4 in both acute and chronic phases of schistosomiasis.

On the other hand, IL-12 may have potential use in preventing or treating parasite-induced pathology resulting from Th2 cytokine production. In the current study, IL-12 levels were significantly decreased in acute and chronic infection with *S. mansoni* which may explain the progression of fibrosis that follows the inflammatory process during the chronic phase and this was agreed with Wynn *et al.*\[^32\] The previous report showed that IL-12 enhances protective immunity in mice engendered by immunization with recombinant *S. mansoni* fatty acid-binding protein as reduction of hepatic granuloma area was only observed when IL-12 was coadministrated with the vaccine.\[^33\]

**Conclusion**

Our results confirmed the role of cytokines in the evolution and regulation of schistosoma-induced immunopathology. Moreover, our data suggested that the body response to *S. mansoni* infection by elevating the expression of the stress gene HO-1 and that VEGF may serve as a new indicator of progression of *S. mansoni* associated angiogenesis which regulates granuloma and/or fibrosis development in the liver.
of infected mice. Understanding the role of HO-1 and VEGF in pathogenesis of *S. mansoni* may provide a new pharmacological target in therapeutic management of schistosomiasis.

**Authors’ Declaration Statements**

**Ethics approval**

All experimental protocols were approved by the Ethics Committee at the Faculty of Pharmacy, Suez Canal University (Ismailia, Egypt) (code # 201906RA1).

**Availability of data and material**

Not applicable

**Competing interests**

The authors declare that there are no conflicts of interest.

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**Authors’ Contributions**

All authors contributed in the design of the study. All authors participated in data collection and analysis, data interpretation, and manuscript writing.

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