Epidemiological and Molecular characterization of antigens extracted from hydatid cysts of camel, cattle and donkeys in Egypt.

O.A. Mahdy, Abdel- Maogood S.Z, Abdel - Wahab A.M. and M. M. El-Bahy
Department of Parasitology Faculty of Veterinary Medicine, Cairo Univ. El-Giza, Egypt

dr.olfat.mahdy@cu.edu.eg

Summary

Hydatidosis is one of the most important parasitic zoonoses and remains a public health and economic problem all over the world. The hydatid cysts (HC) were collected from freshly slaughtered 270 (180 camels, 90 cattle) at Cairo abattoir and from 120 inspected donkeys at Giza zoo, Egypt. Cyst fluid was obtained from hepatic and pulmonary cysts for demonstration of protoscolices and hooklets. The incidence of infection of HC was 18.9%, 3.3% and 14.2% among examined camels, cattle and donkeys respectively, 5.7 %, 66.7% and 90.2% had in HC liver respectively, while the infection rate was 94.3%, 33.3% and 9.8% in the lung of examined camels, cattle and donkeys respectively. The rate of fertile cysts was found to be 42 (79.24%) in camel and 15 (29.4%) in donkeys. while, all cysts collected from the inspected cattle were found calcified. PCR amplification was used for identification of internal transcribed spacer gene 1 (ITS 1) of fertile HC obtained from camel and donkeys by using specific primer. The amplified DNA fragment was further analyzed by PCR mediated restriction fragment length polymorphism (PCRRFLP) using two restriction enzymes (MSP1 and RSA1). The PCR yielded similar amplified DNA band of the same molecular size marker at 1115 bp in different isolates of Hydatid. No band variation of ITS 1 gene could be detected by PCRRFLP by using two restriction enzymes. Amplification product of ITSI
after digestion with MSP1 showed at 661 bp and 406 bp, while those restricted with RSA1 enzyme appeared at 745 bp and 360 bp.

**Key words:** Hydatid cyst, molecular, antigens, PCR, camel, cattle, donkey, Egypt.

**Introduction**

Hydatidosis is a global animal and human health problem of increasing economic and public health importance [1]. It is a helminthic cyclo-zoonotic disease caused by the larval stage (metacestode) of the canid tapeworm *Echinococcus* that require at least one other vertebrate host for completion of its life cycle. The disease is endemic in many parts of the world [2]. It is one of the major zoonotic parasitic diseases in the Middle East and Arab North Africa from Morocco to Egypt [3]. The disease has great public health importance and economic impact in countries where livestock industry is an important segment of the agricultural sector and when livestock production is based mainly on extensive grazing system [4].

Regarding its molecular characterization *E. granulosus* poses a high degree of genetic diversity based on genome pattern, morphology and host specificity have allowed the differentiation of at least Ten different genotypes (G1- G10) among which G4 (horse strain) have been formerly characterized [5-8] and Camel strain (G6) have been formally in Eastern Africa [9] North Africa [10] and Tunisia [11].

The present study was conducted on slaughtered camel, cattle and scarified donkeys aiming to determine the prevalence of hydatid infection. PCR was used for amplification of DNA extracted from fertile HC for identification of ITS1 gene of camel and donkeys followed by further identification by PCR- RFLP using two digestive enzymes *MSP1* and *RSA1*. 
MATERIAL AND METHODS
This study was assessed and approved by Faculty of Veterinary Medicine, Cairo University Ethics Committee and therefore been performed in accordance with the ethical standards laid down in the 1964 Declaration of Helsinki.

Sampling:
The HC were collected from freshly slaughter animals 270 (180 camels, 90 cattle) at Cairo abattoirs and from inspected 120 donkeys on post- at the zoo of Giza Zoo, Egypt. The donkeys were brought to the zoo for feeding lions. They were examined for the presence of HC on post- slaughtered and post- mortem inspection during the period from December 2011 to November 2012. Intact HC, isolated from the infected animals, were put separately in the polythene bags containing ice and brought to Veterinary Medicine of Cairo University for further processing. Examination of all internal organs was also done by using palpation and incision for the detection of HC according to the technique recommended by [12].

Microscopic Identification of HC
The suspected infected organs were collected from slaughtered and scarified animals for routine microscopic examination according to[13]. Cyst fluid was obtained from pulmonary and hepatic cysts for demonstration of protoscolices and hooklets. Protoscolices were isolated from the fertile cysts and then washed three times by phosphate buffer saline (PBS), pH 7.2 and preserved in 70% alcohol (v/v) for isolation of DNA [14].

DNA extraction:
DNA was extracted from germinal layer of fertile HC using Genei Ultrapure TM Mammalian Genomic DNA Purification Tissue Kit (Bangalore Genei). According to manufacturer’s instructions, 25μl of
the antigen were suspended in 500μl of CTAB buffer and transferred to a microfuge tube, incubated for 15 min. at 55°C in water bath, then the mixture was centrifuged at 12000 rpm for 5 min. and the supernatant transferred to a clean microfuge tubes. To each tube 250μl of chloroform: IsoAmyl alcohol (24:1) was added and the solution then mixed by inversion. The mixture was spanned at 13000 rpm for 1 min. Here the upper aqueous phase contained the DNA which is transferred to a clean microfuge tube. To each tube 50μl of 7.5 M Ammonium acetate was added followed by 500μl of ice cold absolute ethanol to precipitate the DNA. The precipitate was transferred into a microfuge tube containing 500μl of ice cold 70% ethanol and then centrifuged at 13000 rpm for 1 minute supernatant discarded and the remaining DNA pellet washed by adding 70% ethanol, then centrifuged at 13000 rpm for 1 min. and again the supernatant removed and the DNA re-suspended in DNase and incubated at 65°C for 20 min and stored at 4°C.

**PCR Assay**

Amplification of ITS1 gene was done by using of primers described by [15]. The primer was designed as forward 5' GTC GTA ACA AGG TTT CCG TA'3 and reverse 5'TCT AGA TGC GTT CGA A(G/A) TGT CGA TG'3, (Jena, Bioscience, Germany). A100-bp DNA was used as molecular size marker. The amplification reaction was carried in 25μl volume containing 500mM Kcl, 10 mM Tris- Hcl (PH9.0), 1% Triton x-100, 4 mM Mgcl, 100uM dNTPs each, 15-20ng of ITS1 primer, 25ng of DNA and 1.5 units of Tag DNA polymerase. For data analysis PCR assay was performed in thermal cycler (Teche TC – 512UK). The DNA was denaturated for 6 min. at 95°C. The mixture was then subjected to 30 cycles of denaturation at 94°C for 45 sec., annealing of primers at 55°C for 60 sec. and primer extensions at 72°C for 90 second. The final extension was held at 72°C for 1 min. PCR products were analyzed after
electrophoresis in 1.5% (W/v) agarose gel and visualized in ethidium bromide.

**PCR - Mediated RFLP**

PCR product were digested with MSP1 and RSA1 (10u) using buffer recommended by the manufacture (Jena Bioscience, Germany). Restriction fragments were separated by gel electrophoresis through 2%TBE agarose gel. PCR products were analyzed after electrophoresis in 1.5% (W/v) agarose gel and visualized in ethidium bromide.

**Results**

The data demonstrated in table (1) cleared that the total incidence of infection by HC was 18.9%, 3.3% and 14.2% among examined camels, cattle and donkeys respectively. The cyst was diagnosed in 5.7%, 66.7% and 90.2% of the examined liver respectively, while, it was 94.3%, 33.3% and 9.8% in the examined lung of camels, cattle and donkeys respectively, table (2). Moreover, the rate of fertile cysts was found to be 42 (79.24%) in camel and 15 (29.4%) in donkeys. while, all cysts collected from the inspected cattle were found calcified. table (3).

Microscopically, the higher incidence of viable motile protoscoleces (60.4%) was found in HC of camel origin, then (23.5%) in HC of donkey origin. The highest incidence of non motile protoscoleces (18.9%) were found in that of camel origin while it was 5.9% in that of donkey origin, table (4).

<table>
<thead>
<tr>
<th>Animals</th>
<th>No. Ex.</th>
<th>No. Inf.</th>
<th>%</th>
<th>No. cysts</th>
<th>Mean and average No. Cyst/animals</th>
</tr>
</thead>
<tbody>
<tr>
<td>Camels</td>
<td>180</td>
<td>34</td>
<td>18.9</td>
<td>53</td>
<td>(1-3)1.55</td>
</tr>
<tr>
<td>Cattle</td>
<td>90</td>
<td>3</td>
<td>3.3</td>
<td>6</td>
<td>(1-2)1</td>
</tr>
<tr>
<td>Donkeys</td>
<td>120</td>
<td>17</td>
<td>14.2</td>
<td>51</td>
<td>(2-3)3</td>
</tr>
<tr>
<td>Total</td>
<td>390</td>
<td>54</td>
<td>13.8</td>
<td>110</td>
<td></td>
</tr>
</tbody>
</table>
Table (2): Incidence of HC distributions in different site of infection in the examined animals

<table>
<thead>
<tr>
<th>Infected animals</th>
<th>No. cysts</th>
<th>Site of infection</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>No.</td>
<td>%</td>
</tr>
<tr>
<td></td>
<td>Lung</td>
<td>Liver</td>
</tr>
<tr>
<td>Camels</td>
<td>53</td>
<td>50</td>
</tr>
<tr>
<td>Cattle</td>
<td>6</td>
<td>2</td>
</tr>
<tr>
<td>Donkeys</td>
<td>51</td>
<td>5</td>
</tr>
<tr>
<td>Total</td>
<td>110</td>
<td>57</td>
</tr>
</tbody>
</table>

Table (3): Incidence of sterile and fertile HC in the examined animals.

<table>
<thead>
<tr>
<th>Animals</th>
<th>No. Ex.</th>
<th>No. Inf.</th>
<th>%</th>
<th>No. Cyst</th>
<th>Sterile HC</th>
<th>Fertile HC</th>
<th>Calcified HC</th>
<th>Calcified HC</th>
</tr>
</thead>
<tbody>
<tr>
<td>Camels</td>
<td>180</td>
<td>34</td>
<td>18.9</td>
<td>53</td>
<td>3</td>
<td>5.7</td>
<td>42</td>
<td>79.24</td>
</tr>
<tr>
<td>Cattle</td>
<td>90</td>
<td>3</td>
<td>3.3</td>
<td>6</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Donkey</td>
<td>120</td>
<td>17</td>
<td>14.2</td>
<td>51</td>
<td>17</td>
<td>33.3</td>
<td>15</td>
<td>29.4</td>
</tr>
<tr>
<td>Total</td>
<td>390</td>
<td>54</td>
<td>13.8</td>
<td>110</td>
<td>20</td>
<td>18.2</td>
<td>57</td>
<td>51.8</td>
</tr>
</tbody>
</table>

Table (4): Incidence of HC according to viability of Protoscolces

<table>
<thead>
<tr>
<th>Animals</th>
<th>No. Cyst</th>
<th>No. fertile cyst</th>
<th>Fertile cysts with</th>
<th>Motile Protoscolices</th>
<th>Non-motile Protoscolices</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>No.</td>
<td>%</td>
</tr>
<tr>
<td>Camels</td>
<td>53</td>
<td>42</td>
<td></td>
<td>32</td>
<td>60.4</td>
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<tr>
<td>Cattle</td>
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<td>0</td>
</tr>
<tr>
<td>Donkeys</td>
<td>51</td>
<td>15</td>
<td></td>
<td>12</td>
<td>23.5</td>
</tr>
<tr>
<td>Total</td>
<td>110</td>
<td>57</td>
<td></td>
<td>44</td>
<td>40</td>
</tr>
</tbody>
</table>

Identification of the genetic characters of hydatid cysts obtained from camel and donkeys after PCR amplification of ITS1 gene showed similar pattern of PCR product, all amplified DNA products have band of the same molecular size at 1115bp on agarose gel (Fig.1). Further more molecular analysis using PCR-RFLP for amplification to the product of ITS1 after digestion with MSP1 showed that all strains sharing in the MW at 661 bp and 406 bp, while those restricted with RSA1 enzyme appeared at 745 bp and 360 bp.
Fig.(1): Ethidium bromide stained agarose (2%) gel showing amplification product of ITS1 of HC germinal layer of E. granulosus by PCR .lane MWA: a 100 bp molecular size marker ; lane A Camel DNA & B Donkey DNA containing 1115 bp.

**Discussion**

*Echinococcus granulosus* is medically and economically one of the most important zoonoses. Hydatid cyst develops in the internal organs of human and herbivore intermediate hosts, mainly in the liver and lung [16]. In Egypt the prevalence of hydatidosis still a point under investigation, In the present study, the incidence of HC were 18.9 % and 3.3% in slaughtered camels and cattle from Cairo abattoirs. This result was in agreement with [17] and [18] who recorded 18.9% and 17.61% in slaughtered camels in Ismailia and Cairo abattoirs, Egypt. In the same time, this result considered to be lower than that recorded by[19] in Assiut governorate, as 7.67% in camels and no infection was recorded in the examined cattle and buffaloes. Presently, incidence of donkey HC infection was (14.2%) this finding considered to be higher than that result by [20]On the other hands, the low incidence of Cattle HC infection (3.3%) was in agreement with [21] and [19] where, both of them failed to diagnose HC infection in the examined slaughtered cattle.
The current study indicated that the rate of infection in camel was higher in Lung (94.3 %) than in liver (5.7 %). An observation in accordance with that noticed in Egypt camel, [18 &22] who found that 100%, 63.7%, and 94.6% in lung camel. The majority of infected donkeys (90.2%) in the present study harbored hydatid cysts in their liver. This result was in agreement with [8], who revealed that the majority of infected donkeys (70%) harbored HC in their livers at Beni-Suef, Egypt.

Based on the epidemiology and molecular studies, the fertility of cyst is one of the most important factors in the epidemiology of *E. granulosus*. The fertility of cyst varies depending on the hosts and geographical situations [23]. In the current study, fertility rate of HC in camel and donkeys have been found to be 42 (79.24%) in camel lung and 15 (29.4%) in donkey liver. The high rate of fertile cyst may indicate that the cause of infection in investigated animals might be due to camel and donkeys strain (G6 & G4). As such genotype is commonly recognized as a predominating species of *E. granulosus* in Mediterranean countries[24].

Molecular genetics study has been carried out to identify the genetic characters of HC obtained from the infected camel and donkeys. After PCR amplification of ITS1 gene, similar amplified DNA band of the same molecular size marker at 1115bp were recorded in the different isolates. No band variation of ITS1 gene could be detected by PCR-RFLP after using the two restriction enzymes, MSP1 and RSA1, This meaning absence of ITS1 variant which could not differentiated using these two restriction enzymes. This was in agreement with [25-26]. In the author's opinion and in agreement with [26] absence of variation in amplified ITS1 and indistinguishable genetic character in PCR-RFLP, meaning that the analyzed camel and donkey HC samples
are infected with *E. granulosus* of sheep strain. This can be accepted as all examined animals are from the same localities. Moreover more research are continue aiming to further identification of hydatid infection of camel and donkeys based on PCR amplification and sequence of mitochondrial genes.

**Acknowledgment**

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**References**


