International Journal for Parasitology xxx (2016) xxx-xxx

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International Journal for Parasitology



journal homepage: www.elsevier.com/locate/ijpara

Hemozoin is a product of heme detoxification in the gut of the most medically important species of the family Opisthorchiidae

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ARTICLE INFO

22 Article history:

- 23 Received 28 October 2015
- 24 Received in revised form 8 December 2015
- 25 Accepted 10 December 2015
- 26 Available online xxxx
- 27 Keywords:
- 28 Opisthorchis felineus
- 29 Opisthorchis viverrini
- 30 Clonorchis sinensis
- 31 Hemozoin
- 32 Lipid droplets
- Blood-feeding parasite

ABSTRACT

Many species of trematodes such as Schistosoma spp., Fasciola hepatica and Echinostoma trivolvis are blood-feeding parasites. Nevertheless, there is no consensus on the feeding habits of the family Opisthorchiidae (Opisthorchis felineus, Opisthorchis viverrini and Clonorchis sinensis). Previously, histological studies of O. felineus and C. sinensis revealed some dark stained material in their gut lumen. In this study we conducted a comprehensive analysis of the gut contents of three members of the family Opisthorchiidae (O. felineus, O. viverrini and C. sinensis). Using transmission electron microscopy, we demonstrated for the first known time the presence of disintegrating blood cells in the gut of O. felineus as well as electron-dense crystals in the gut of O. felineus and C. sinensis. Electron energy loss spectroscopy revealed iron atoms in these crystals, and mass spectrometry of the purified pigment demonstrated the presence of heme. Fourier-transform infrared spectroscopy identified the signature peaks of the common iron-carboxylate bond characteristic in crystals isolated from O. felineus and C. sinensis. Scanning electron microscopy showed layered ovoid crystals of various sizes from 50 nm to 2 µm. Morphological, chemical and paramagnetic properties of these crystals were similar to those of hemozoin from Schistosoma mansoni. Crystal formation occurs on the surface of lipid droplets in O. felineus and C. sinensis guts. Our results suggest that the diet of O. felineus and C. sinensis includes blood. Detoxification of the free heme produced during the digestion proceeds via formation of insoluble crystals that contain iron and heme dimers, i.e. crystals of hemozoin. Furthermore, we believe that biocrystallisation of hemozoin takes place on the surface of the lipid droplets, similar to S. mansoni. Hemozoin was not detected in the closely related species O. viverrini.

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59 1. Introduction

Infections caused by trematodes of the family Opisthorchiidae–
Opisthorchis felineus, Opisthorchis viverrini and Clonorchis
sinensis—remain a major public health problem in many parts of
Asia and Europe. It has been established that more than 40 million
people are infected and approximately 600 million are at risk of
being infected by these parasites; furthermore, there has been an

http://dx.doi.org/10.1016/j.ijpara.2015.12.003

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increase in the incidence of these diseases in non-endemic regions such as Israel (Yossepowitch et al., 2004; Keiser and Utzinger, 2009). Infection is acquired through ingestion of raw or undercooked freshwater Cyprinoid fish carrying infective metacercariae. Mature flukes inhabit the large intrahepatic and extrahepatic bile ducts, the gallbladder and, occasionally, pancreatic ducts. In humans, the infection that is caused by these liver flukes is characterised by long duration (decades) and development of severe complications such as cholangitis, cholecystitis, obstructive jaundice, liver abscesses and pancreatitis including cholangiocarcinoma, a bile duct cancer (Al'perovich and Brazhnikova, 1989; Sripa, 2003; Hong and Fang, 2012).

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The parasites of the family Opisthorchiidae coexist with the host for a long time. Thus, understanding of their physiology, and the feeding habits in particular, is essential to identify the mechanism through which complications develop in the definitive host. This, in turn, may also help improve disease diagnosis and treatment.

The peculiarities of the digestive tract structure in trematodes, e.g., the oral sucker, participating both in attachment and ingestion of food, as well as the well-developed muscle layer around the initial segments of the digestive tract, ensure that these parasites can feed only via active suction of liquid or semiliquid food (Smyth and Halton, 1983). To date, there is no consensus on the feeding habits of species of the Opisthorchiidae. It is believed that they ingest bile, mucin or suck tissue fluid from the naked connective tissue stroma and possibly blood (Hou, 1955; Sripa, 2003). A large number of parasitic species, including trematodes (*Schistosoma* spp., *Fasciola hepatica, Echinostoma trivolvis*, and others) consume the host's blood (Smyth and Halton, 1983; Oliveira et al., 2000; Pisciotta et al., 2005). According to Dalton et al. (2004), hematophagy is the predominant feeding strategy of most trematodes.

98 Erythrocytes or red blood cells (RBCs) are the most common 99 type of blood cell. Hemoglobin constitutes approximately 90% of the total protein in RBCs, and its degradation by proteolytic 100 enzymes results in the release of large amounts of the 101 102 iron-containing prosthetic group, heme. Exogenous heme has to 103 be metabolised quickly due to its toxicity, so heme degradation is 104 one of the key physiological processes for blood-feeding organisms. 105 It has been established that free heme strongly catalyses the gener-106 ation of reactive oxygen species, which in turn lead to oxidation of 107 lipids, proteins and DNA (Kumar and Bandyopadhyay, 2005). Due to 108 its lipophilic nature, heme can also associate with phospholipid 109 membranes, disrupting their physical integrity (Jeney et al., 2002). Blood-feeding parasites have several strategies for detoxifi-110 cation of heme derived from the catabolism of host RBCs (Toh 111 112 et al., 2010). The most prevalent pathway involves formation of 113 an iron-containing black or dark brown crystalline material, which 114 was initially found in the digestive vacuole of the malaria parasite 115 and therefore bears the name "malaria pigment" or hemozoin 116 (Pagola et al., 2000). Blood-feeding trematodes such as Schistosoma 117 spp. and E. trivolvis use a similar pathway for heme detoxification (Oliveira et al., 2000; Pisciotta et al., 2005). Hemozoin consists of 118 centrosymmetric dimers of heme that are linked via coordinate 119 iron-carboxylate bonds between Fe³⁺ of one heme and propionate 120 121 side chains of the other; these dimers form crystals through hydrogen bonds (Slater et al., 1991; Pagola et al., 2000). In contrast to 122 123 monomeric heme, hemozoin crystals are insoluble in water, some 124 detergents or bicarbonate buffer and have paramagnetic properties 125 (Fitch and Kanjananggulpan, 1987; Butykai et al., 2013). Hemozoin 126 formation is specific to parasites; therefore, it is an attractive target 127 for the development of new diagnostic and treatment methods for 128 opisthorchiasis and clonorchiasis. For instance, detection of hemo-129 zoin as heme hyper-accumulated crystals in RBCs by laser desorption mass spectrometry (MS) is one of the most sensitive and 130 specific methods for diagnosis of malaria (Demirev et al., 2002). 131

132 Our comparative histopathological studies of the hepatobiliary system during experimental opisthorchiasis led us to hypothesise 133 134 that formation of hemozoin is likely a part of the nutrition process of O. felineus but not of O. viverrini (Lvova et al., 2012). We showed 135 136 that starting from week 3 after the infection by O. felineus and until 137 the end of the experiment (6 months after the infection); there 138 were brown-black pigments in the liver tissue of the hamsters 139 and in the gut lumen of O. felineus, but not in the gut of O. viverrini. 140 However, experiments with differential histological staining failed 141 to identify the nature of these pigments (Lvova et al., 2012). Similar 142 results were obtained with C. sinensis where the dark granules and 143 blood cells were found in the gut lumen of this liver fluke (Chu et al.,

1982). It is logical to assume that O. felineus and C. sinensis consume 144 blood and the lumen of their gut contains hemozoin. Thus, a com-145 prehensive analysis of the gut contents in some members of the 146 family Opisthorchiidae (O. felineus, O. viverrini and C. sinensis) was 147 the main aim of our study. To this end we used a complex analytical 148 approach combining electron microscopy, spectroscopy, MS and 149 Fourier-transformed infrared (FTIR) spectroscopy to reveal the nat-150 ure and the origin of granules observed in the gut of O. felineus and 151 C. sinensis. 152

2. Materials and methods

2.1. Ethics statement

Animal experiments were approved by the Committee on the Ethics of Animal Experiments of the Institute of Cytology and Genetics, The Siberian Branch of the Russian Academy of Sciences, Russia (permit number 7 of 19 December 2011). All of the procedures were in compliance with The Code of Ethics of the World Medical Association (Declaration of Helsinki) for animal experiments http://ec.europa.eu/environment/chemicals/lab_animals/ legislation_en.htm.

2.2. Parasites

Metacercariae of O. viverrini, O. felineus and C. sinensis were 164 obtained from naturally infected fishes (family Cyprinidae) cap-165 tured from fresh water reservoirs in the endemic areas of Khon 166 Kaen province, northeastern Thailand; Novosibirsk, Russia and 167 the Far East, Russia, respectively. The fish meat was digested by 168 pepsin – HCl overnight at 37 °C followed by filtration. After being 169 washed several times with normal saline, the metacercariae were 170 collected and identified under a light microscope. Three-month 171 old hamsters (Mesocricetus auratus) were infected per os with 50 172 viable active metacercariae. Adult worms of O. viverrini, O. felineus 173 and C. sinensis were obtained from bile ducts of euthanised ham-174 sters within 3 months after the infection. The fresh worms were 175 washed several times in sterile normal saline containing penicillin 176 (100 U/mL) and streptomycin (100 μ g/mL) to remove any debris or 177 residual blood and to prevent bacterial contamination. After thor-178 ough washing, the viable worms were used to prepare specimens 179 for electron microscopy and obtain crystals. 180

2.3. Transmission electron microscopy (TEM)

For TEM, we used 3-5 adult worms of each species (O. felineus, 182 C. sinensis and O. viverrini). They were dissected into small parts 183 that were fixed with 2.5% glutaraldehyde solution in 0.1 sodium 184 cacodylate buffer (pH 7.2) for 2.5 h. After that, the samples were 185 washed and incubated for 1 h in a 1% osmium tetroxide solution 186 in the same buffer. The samples were washed with water and 187 placed in 1% uranyl acetate aqueous solution for 12 h at 4 °C. The 188 samples were dehydrated through a graded series of ethanol and 189 acetone, and embedded in Agar 100 Resin (Agar Scientific Ltd., Uni-190 ted Kingdom). Ultrathin sections were double stained with uranyl 191 acetate and lead citrate according to Reynolds (1963) and exam-192 ined using a JEOL 100 SX electron microscope. 193

2.4. Electron energy loss spectroscopy (EELS) and electron spectroscopic imaging (ESI)

For parallel EELS and ESI, 70–80 nm thick unstained sections196were used. The latter were mounted on unfilmed copper grids197and then coated with carbon. The resulting samples were analysed198at 120 kV in an EFTEM Libra 120 microscope with an in-column199

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