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A common oocyst surface antigen of *Cryptosporidium* recognized by monoclonal antibodies

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Abstract Two hybridoma clones, CMYL3 and CMYL30, were generated by immunizing Balb/c mice with excysted oocysts of Cryptosporidium muris. Both clones secreted monoclonal antibodies against an oocyst-wall antigen with apparent molecular mass of 250 kDa (called CM250) from C. muris and C. parvum. The epitope appeared to be periodate-sensitive, suggesting the involvement of a carbohydrate moiety. Immunofluorescence and confocal microscopy on purified oocysts and infected mouse tissues revealed staining confined to the oocyst wall of both Cryptosporidium species. Immunogold labeling further revealed the presence of the CM250 antigen in electron-dense vesicles and cytoplasm of developing macrogametocytes, and ultimately localized to the oocyst wall of mature oocysts. Both antibodies cross-reacted with C . serpentis oocysts but did not recognize the other enteropathogenic protozoans Giardia muris, Eimeria falciformis and E. nischulz. These antibodies may be valuable tools for the analysis of oocyst-wall formation in Cryptosporidium and characterization of the common antigen.

Introduction

Cryptosporidiosis is an opportunistic disease that can be fatal to immunocompromised patients through the induction of chronic diarrhea. The infection is transmitted

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by oocysts in a fecal-to-oral route, with parasite development occurring in the epithelial cells of the gastrointestinal tract. In the genus Cryptosporidium, six to eight species have been tentatively approved as valid, based on differences in host specificity, oocyst morphology and the site of infection (Fayer et al. 1997; O'Donoghue 1995). Recent phylogenetic analysis of small subunit rRNA sequences of several Cryptosporidium isolates revealed that at least four distinct species, C. parvum, C. baileyi, C. muris, and C. serpentis, are valid (Xiao et al. 1999). Although human infections are chiefly due to C. parvum, some reports have revealed that C. baileyi and C. muris are also capable of infecting humans (Ditrich et al. 1991; Katsumata et al. 2000), but these observations still have to be confirmed. Moreover, clinically, C. muris infection is less pathogenic than C. parvum (Taylor et al. 1999).

In addition to oocyst morphology and phylogenetic differences at the gene level, the different sites of infection support a valid distinction between C. parvum and C. muris. C. parvum infects the small intestine, preferentially in the ileum, while C. muris infects the stomach. The oocyst wall undoubtedly provides protection for sporozoites via a highly resistant barrier to external influences, such as the environment external to the host, as well as assaults from the host organism. The differential localizations of infections between C. parvum and C. muris raise questions about possible differences in oocyst-wall composition.

Monoclonal antibodies (mAbs) have proven useful, not only in the characterization of unique antigens, but also in the detection of oocysts from feces for diagnostic purposes (Anusz et al. 1990; McLauchlin et al. 1987; Sterling and Arrowood 1986), in neutralizing infectivity, and in providing relief from cryptosporidiosis symptoms in murine models (Langer and Riggs 1996; Perryman et al. 1990). Various parasitic protists are known to induce diarrhea. One of these, Giardia lamblia, is a human parasite with clinical symptoms similar to cryptosporidiosis (Roberts and Janovy 1996). The cyst of G. lamblia in human stool is about $8-12$ µm long, only

slightly larger than Cryptosporidium oocysts. In addition, many domestic animals are affected by Eimeria species (Roberts and Janovy 1996). This Apicomplexan parasite also induces diarrhea. Whether similar oocyst components exist between Eimeria and Cryptosporidium remains to be determined. Therefore, the identification and characterization of new parasite oocyst antigens could be useful for diagnosis and for understanding the molecular biology of the parasites.

Most antibodies developed to Cryptosporidium are against C. parvum. To date, no mAb against C. muris has been developed. In this study, we generated mAbs against a C. muris antigen that localizes to the oocyst wall of both immature and mature oocysts, as well as in cytoplasmic vesicles of developing macrogametocytes.

Materials and methods

Oocyst purification and antigen preparation

The oocysts of C. muris were obtained from the feces of C57BL6 female mice, which were orally infected with 2×10^6 oocysts per mouse, as described previously (Yu 1998). The oocysts of C. parvum were obtained by the same method as C. muris, but the mice were immunosuppressed by intramuscular injection of 0.1 mg/g body weight of methylprednisolone acetate (Depomedrol, Pharmacia and Upjohn Ltd, Hwaseong, Korea) once a week (Yu and Chai 1995). Mouse feces were examined by modified Ziehl–Neelsen staining to confirm oocyst shedding (Casemore et al. 1985) and collected in 2.5% potassium dichromate at 4° C. The oocysts were purified as described by Petry et al. (1995). For the antigen preparation, the oocysts were excysted in 0.04% taurocholic acid in phosphate-buffered saline (PBS) at 37° C for 1 h. After washing once in PBS, excysted oocysts were homogenized by sonication for 20 min at 4 $^{\circ}$ C.

Production of mAbs

Eight-week-old female $BALB/c$ mice were immunized intraperitoneally by the injection of 1.5×10^7 excysted and homogenized oocyst antigen emulsified in an equal volume of TiterMax (CytRx Corporation, Norcross, Ga.). Three additional intraperitoneal injections, each with 1×10^7 oocysts, were given at 3-week intervals. At 63 days, mice were bled from the tail and the sera obtained were tested for anti-Cryptosporidium antibody by indirect immunofluorescence. A final boosting $(5\times10^6$ oocysts) was given intravenously 4 days before fusion. Spleen cells were isolated and fused with mouse myeloma NSI as described by Lin et al. (1985). The cultured medium of hybridomas was screened by enzyme-linked immunosorbent assay (ELISA), according to the method of Tijssen (1985). The cloning of positive hybridomas was performed as described previously (Lin et al. 1985). Two hybridoma clones, CMYL3 and CYML30, were used in this study. The antibody CMYL3 was an IgM, whereas the antibody CMYL30 was an IgG1 subclass, as determined by a Sigma ImmunoType kit (Sigma, St. Louis, Mo.).

Immunofluorescence microscopy

Purified oocysts were smeared and air-dried on glass slides. For immunofluorescence, the slides were then incubated with primary antibody and subsequently with fluorochrome-conjugated secondary antibodies, as described below. Mouse stomach and small intestine were dissected from Cryptosporidium-infected mice and immediately frozen with liquid nitrogen in OCT compound (Miles, Elkhart, Ind.). Tissues were cryo-sectioned at a thickness of 4 μ m and stored at -70 °C until use. Upon use, tissue sections were fixed in 3.7% formaldehyde and permeabilized in cold acetone. After a PBS wash, the slides were incubated with cell supernatants or ascites diluted appropriately with3% bovine serum albumin in PBS, washed in PBS and incubated with a 1:400 dilution of fluorescein isothiocyanate (FITC) – conjugated goat anti-mouse IgG whole molecule antibody (Cappel, West Chester, Pa.) in PBS. Slides were washed, mounted in polyvinyl alcohol (Gelvatol), and observed with a Zeiss epifluorescence microscope or a Leica scanning-laser confocal microscope.

SDS-PAGE and Western blotting

SDS polyacrylamide gel electrophoresis (SDS-PAGE) was carried out according to Laemmli (1970), with a low concentration of bisacrylamide as described previously (Lin et al. 1985). Total protein extracts were prepared by directly dissolving 1×10^8 oocysts in 1 ml of gel sample buffer, as described by Bonnin et al. (1991) and extracts from about $7.5\times10^5\sim2\times10^6$ oocysts were loaded per lane. Oocyst walls were further purified from both C. parvum and C. muris oocysts according to Harris and Petry (1999) and extracts from $2.5-5\times10^5$ oocyst walls were loaded per lane. After electrophoresis, the proteins in the gel were electrophoretically transferred to PVDF membranes. The membranes were blocked in 5% skim milk in Tris-buffered saline (TBS, 50 mM Tris pH 7.5, 150 mM NaCl) overnight at 4 °C . CMYL3 and CMYL30 ascites were diluted 4000-fold and 700-fold, respectively, in 3% bovine serum albumin in TBS. Eachwas administered to respective blots for 1 h at room temperature in a humidified chamber. After three changes of TBS containing 0.2% Tween 20 over 45 min, the membranes were incubated with horseradish peroxidase-conjugated sheep antimouse immunoglobulin (Ig) diluted 2,000-fold in TBS. After washing as above, bound antibodies were detected with ECL Western blotting analysis system (Amersham, UK).

Periodate treatment

To determine whether the antibody epitopes contain carbohydrate, Western blot analyses were performed on the antigen before and after periodate treatment, as described by Woodward et al. (1985). Briefly, after blocking with 5% skim milk in TBS, the nitrocellulose membrane was incubated with 10 mM sodium orthoperiodate in 50 mM sodium acetate buffer (pH 4.5) for 1 hin the dark at room temperature, and then exposed to 50 mM sodium borohydride in PBS for 30 min. Following three washes with TBS-Tween 20, the membranes were incubated in CMYL30 as described above. A control membrane was incubated in 50 mM sodium acetate buffer without sodium orthoperiodate.

Immunogold labeling

The stomach and small intestine of Cryptosporidium-infected mice were dissected out and fixed in 2% paraformaldehyde and 0.4% glutaraldehyde for approximately 2 h at room temperature. The fixed tissues were washed with 0.1 M phosphate buffer (pH 7.2), and dehydrated through an alcohol series from 30% to 95%. Dehydrated tissues were embedded in LR gold resin (London Resin Company, UK) and polymerized at -20 °C for 72 h under UV illumination. The ultrathin sectioning was performed at 90-nm thickness and sections were mounted onto nickel grids. The immunogold labeling procedure followed the method of Yu and Chai (1995). Briefly, tissue sections were incubated in PBS-milk-Tween (PMT, 3% skim milk and 0.01% Tween 20 in PBS) for 10 min and exposed to ascites solution of mAb CMYL3 or CMYL30 diluted with PMT for 2 h at room temperature. The sections were washed thoroughly with PBS-BSA-Tween (1% bovine serum albumin and 0.01% Tween 20 in PBS) and incubated with 5 nm gold conjugated goat anti-mouse IgG or IgM (Sigma) overnight at $4 °C$. Silver enhancement was done with a commercial kit (Amersham) followed by staining with uranyl acetate and lead citrate. The stained sections were examined under a transmission electron microscope, Jeol 1200 EXII.

Results

A common antigen CM250 was recognized by both CMYL3 and CMYL30

Although total protein profiles, analyzed by SDS-PAGE, of oocysts prepared from C. muris and C. parvum were very similar, species-enriched protein bands could be also detected (Fig. 1). Whether these species-enriched proteins may have roles in determining the different localization of infection for different species remains to be determined. In this study, we have generated two mAbs, CMYL3 (IgM) and CMYL30 $(IgG1)$, by using total proteins from C. muris oocysts as immunogens. Western blot analysis showed that both CMYL3 and CMYL30 antibodies cross-reacted with a protein band of 250 kDa (called CM250) from C. parvum oocysts (lane 1 in Fig. 2) and oocyst walls (lane 2 in Fig. 2). Repeat Western blot analysis with CMYL3 antibody on the same oocyst wall sample after a week's storage at -20 °C revealed a size-shift in the antigen molecule (lane 3 in Fig. 2), suggesting that this CM250 antigen was very vulnerable to proteolytic cleavages. The proteolysis was more apparent when Western blot analyses were performed on the total proteins prepared from both oocysts and oocyst walls

of C. muris (lanes 1 and 2 of Fig. 3). However, both antibodies recognized a protein band (CM250), which appeared to have the same molecular size as that from C. parvum. In addition, several bands with lower molecular sizes (suchas 235, 150, 110, 92, 72, and 45 kDa) were detected, presumably representing proteolytic fragments of CM250. The inclusion of a mixture of protease inhibitors (complete protease inhibitor cocktail tablet, cat. no. 1607498, Boehringer Mannheim, Indianapolis, Ind.) in the sample preparation only slightly inhibited proteolysis (data not shown). Therefore, based on these studies, we concluded that a common band of 250 kDa (CM250) in C. muris and C. parvum was recognized by both CMYL3 and CMYL30. The CM250 appeared to be found in the oocyst walls of both C . parvum and C . muris. The next question was whether carbohydrates were involved in the antigenic determinants of CM250 for both antibodies. The oocyst-wall proteins, after membrane transfer, were treated with periodate as described above (Materials and methods) and then used for Western blot analysis. The reactivity of CMYL30 to periodatetreated CM250 and/or its fragments (such as 235 , 110, 92, and 45 kDa) was drastically reduced (Fig. 4B), compared with that in the control blot without periodate treatment (Fig. 4A). Thus, the CM250 appeared to be a glycosylated antigen on the oocyst wall of Cryptosporidium.

Fig. 1 Protein profiles of Cryptosporidium oocyst antigens as separated on a 12.5% SDS-polyacrylamide gel. M Molecular weight marker proteins; lane 1 oocyst antigens of C. muris; lane 2 oocyst antigens of C. parvum

Fig. 2 Western blot detection of C. parvum antigens probed with A CMYL3 and B CMYL30. Antigens reacting to these antibodies were visualized by ECL method after labeling with sheep antimouse Ig conjugated to horseradish peroxidase. Lane 1 C. parvum whole oocyst antigens; lane 2 C. parvum oocyst wall antigens; lane 3 same sample as that in lane 1 except the sample was stored for a week at -20 °C. Both antibodies recognized the antigen CM250 with molecular mass of 250 kDa. Upon storage of the same sample as that in lane 1, the antibody reacted strongly with a 150 kDa band (indicated by b) and weakly with a 235 kDa band (indicated by a). Consequently, the reactivity to the CM250 band became greatly diminished (lane 3). The result suggested that smaller size antigens were derived from the proteolytic cleavage of CM250

Fig. 3 Western blot detection of C. muris antigens probed with mAbs A CMYL3 and B CMYL30. Antigens reacting to these antibodies were visualized by the ECL method after labeling with sheep anti-mouse Ig conjugated to horseradish peroxidase. Lane 1 C. muris whole oocyst antigens; lane 2 C. muris oocyst-wall antigens. Both antibodies clearly recognized the CM250 band from oocyst-wall antigens (lane 2). In addition to the CM250, its proteolytic fragments including a 235 kDa band (a) and a 45 kDa band (f) in the wall antigens were also recognized by both antibodies. The proteolytic fragments recognized by mAbs were more prominent in the oocyst antigens (lane 1). These included 235 kDa (a), 110 kDa (c), 87 kDa (d), and 70 kDa (e)

CM250 antigens were found in wall-forming vesicles of macrogametocytes and predominantly located in the wall of mature oocysts

CMYL3 and CMYL30 reacted with the oocyst walls of C. muris and C. parvum, as observed by indirect immunofluorescence (Fig. 5 A, D and Fig. 6 A, D, respectively). The cryosections of mouse stomach and small intestine infected with Cryptosporidium were immunofluorescently stained with these two antibodies. The C. muris oocysts (indicated by σ in Fig. 5B, E) recognized by both antibodies were almost completely packed in the stomach glands (indicated by sg in Fig. 5C, F). Similarly, both antibodies strongly reacted with the C. parvum oocysts in small intestinal epithelia of infected mice (Fig. 6B, E).

To further demonstrate that the CM250 was localized in the oocyst walls, confocal microscopy was performed on antibody-labeled oocysts. Figure 7 B–H shows an optically sectioned CMYL30 labeled C. muris oocyst from the bottom to the top of oocyst. A uniform staining on the oocyst wall was particularly obvious in

Fig. 4A, B Effects of periodate oxidation on the binding of mAb CMYL30 to Cryptosporidium oocyst antigens. Proteins of C. muris (lane 1) and C . parvum (lane 2) oocysts were separated by SDS-PAGE gel and electrophoretically transferred to duplicated membranes. A The membrane was incubated in 50 mM acetate buffer pH 4.5 only, washed before immunostaining, and used as a control membrane. B The other membrane was treated with 10 mM sodium orthoperiodate in 50 mM sodium acetate buffer pH 4.5, washed before immunostaining, and used as a periodatetreated membrane. The antibody reactivity to the CM250 antigen, as well as its proteolytic fragments (indicated by a, b, c, d, e , and f) was greatly reduced or absent

sections C–F. An infolded suture of the oocyst wall was also clearly labeled (arrow in Fig. 7E). This antibody did not recognize sporozoites or other residual bodies inside of the oocyst (Fig. 7). The CMYL3 antibody also specifically stained the oocyst wall of C. muris as observed by confocal microscopy (data not shown). Similar wallstaining patterns were observed with these two antibodies on the C. parvum oocysts by confocal microscopy (data not shown).

To further confirm the CM250 antigen as a wall antigen, and to follow the fate of the antigen during the life cycle of the parasites, immunoelectron microscopy was performed on Cryptosporidium-infected mouse tissues. Again, both CMYL3 and CMYL30 recognized an antigen (as revealed by 5-nm gold particles) predominantly located in the oocyst-wall structures of the parasites, C. muris (Fig. 8 A, B) and C. parvum (Fig. 8D, E). No gold particles were associated with the parasitophorous vacuole membrane (pvm in Fig. 8), which is of host origin. A broad band of the gold particles was occasionally seen at the wall region near the attachment zone to the host cell (az in Fig. 8E). Other developmental stages were not labeled by gold particles at all (data not shown), with the exception of the macrogametocyte (Fig. 8C, F).

In macrogametocytes, electron-dense vesicles near the periphery were heavily labeled by gold particles (arrowheads in Fig. 8C, F). These vesicles closely Fig. 5A–F Indirect immunofluorescence with mAb CMYL3 and CMYL30 on purified C. muris oocysts and on C. murisinfected mouse stomach. A CMYL3 labeled C. muris oocysts; B CMYL3 labeled C. muris oocysts (indicated by o) in mouse stomach gland; C phase-contrast image of B; D CMYL30 labeled C. muris oocysts; E CMYL30 labeled C. muris oocysts (indicated by o) in mouse stomach gland; F phase-contrast image of E. Bar 10 μ m, sg stomach gland

Fig. 6A–F Indirect immunofluorescence with mAb CMYL3 and CMYL30 on C. parvum. A CMYL3 labeled C. parvum oocysts; B CMYL3 labeled C. parvum oocysts (indicated by o) in small intestinal villi of mouse; C phase-contrast image of B; D CMYL30 labeled C. parvum oocysts; E CMYL30 labeled C. parvum oocysts (indicated by ρ) in small intestinal villi of mouse; F phase-contrast image of E. Bar $20 \mu m$

Fig. 7A–H Confocal micrographs of a C. muris oocyst showing the CM250 antigen localized in the oocyst wall. Purified C. muris oocysts after labeling with CMYL30 were observed and photographed using confocal microscope. A Differential interference-contrast image of a C. muris oocyst; B–H confocal micrographs depict a Z-series of the same oocyst in 0.5 - μ m sections. The photograph in B represents the bottom of the oocyst. The arrow in E indicates the infolded suture of the oocyst. Bar $5 \mu m$

resembled so-called wall-forming bodies. In the C. muris macrogametocyte, many gold particles were also seen clustered around the amylopectin granules in the cytoplasm (arrow in Fig. 8C). No gold particles were detected at the pellicles of macrogametocytes in the infected mouse stomach and intestine $(p$ in Fig. 8C, F). CMYL antibodies cross-reacted with oocysts of all Cryptosporidium species tested but not with oocysts of *Giardia* and *Eimeria*

When both CMYL antibodies were used to stain a stomach mucosal smear from a C. serpentis-infected

Fig. 8A–F Immunogold electron microscopy of ultrathin sections of Cryptosporidium-infected mouse tissues withmAb CMYL3 and CMYL30. C. muris-infected mouse stomach(A–C) and C. parvuminfected mouse small intestine (D–F) were fixed in 2% paraformaldehyde and 0.4% glutaraldehyde and embedded in LR gold. The 90-nm sections were prepared and processed for immunogold labeling. Five-nanometer gold-conjugated goat anti-mouse Ig was used as a secondary antibody. A Oocyst of C. muris labeled with CMYL3; **B** immature oocyst of *C. muris* labeled with CMYL30; numerous 5-nm gold particles are present on the oocyst wall (ow) but not on the parasitophorous vacuolar membrane (pvm) . C Macrogametocyte of C. muris labeled with CMYL30. The peripheral electron-dense vesicles (arrowheads) are heavily labeled with gold particles. Some of the electron-lucent amylopectin granules (am) are also labeled with gold particles. Gold particles are also found in the cytoplasm around amylopectin granules (indicated by *arrow*). Note that the pellicle (p) of the macrogametocyte is not labeled. D Oocyst of C. parvum labeled with CMYL3; E immature oocyst of C . parvum labeled with CMYL30; both antibodies uniformly labeled the oocyst wall and a broader layer of gold particles can be occasionally observed at the attachment zone (az). F Macrogametocyte of C. parvum labeled with CMYL30. The electron-dense wall-forming bodies are labeled with gold particles (arrowhead), whereas the pellicle is not labeled (p) . Bar 1 µm

snake, clear cross-reactivity to the oocysts was observed (Fig. 9). Confocal microscopy revealed that the stained structures were the walls of C. serpentis oocysts (data not shown). It is known that *Giardia* infection in humans causes similar chronic diarrhea and produces slightly larger cysts than Cryptosporidium oocysts (Roberts and Janovy 1996). For diagnosis and detection of oocyst contamination in water supplies, it would be of interest to determine whether our CMYL antibodies can crossreact with Giardia cysts. Furthermore, it is known that Eimeria and Cryptosporidium belong to the same suborder *Eimeriina* and both can infect domestic animals (Roberts and Janovy 1996). It would also be of interest to determine whether our CMYL antibodies recognize oocysts from this protozoan. Therefore, we carried out immunofluorescence microscopy on purified cysts from Giardia muris, and purified oocysts from Eimeria nieschulzi and E. falciformis. There were no detectable stains on these cysts and oocysts (Fig. 10). These CMYL mAbs therefore specifically reacted with oocysts from Cryptosporidium only, and may be most useful in clinical diagnosis to distinguish different protozoan infections.

Discussion

Many kinds of mAbs against C, parvum have been reported (Anusz et al. 1990; Bonnin et al. 1991, 1995; McDonald et al. 1991). However, no mAbs generated against C. muris have been described to date. Although most cryptosporidiosis cases reported are from C. parvum infections, a recent study suggested that C. muris could be infective in humans as well (Katsumata et al. 2000). Our mAbs against C . muris have the potential to be as useful as those against C. parvum for clinical diagnosis of cryptosporidiosis, yet could be immediately useful for the analysis of oocyst-wall formation and composition. While mAbs against sporozoite antigens could be useful for inhibition of parasite invasion, and as markers to study host–parasite interaction, mAbs against oocyst-wall antigens are valuable tools for the diagnosis of Cryptosporidium in fecal material and for the determination of the extent of oocyst contamination

Fig. 9 Indirect immunofluorescence with mAb CMYL30 on a mucosal smear from a C. serpentis-infected snake stomach. Bar $10 \mu m$

Fig. 10 Indirect immunofluorescence with mAb CMYL30 on purified cysts of Giardia muris (A, B), and purified oocysts of Eimeria nieschulzi (C, D) , and E. falciformis (E, F) . A, C, E Phasecontrast images; B, D, F fluorescent images in A, C, and E, respectively. Bar 10 μ m

in water supplies (McLauchlin et al. 1987; Smith and Rose 1990; Sterling and Arrowood 1986).

In the present study, we describe two mAbs, CMYL30 (IgG1) and CMYL3 (IgM), generated against C. muris oocysts. Both recognize an oocyst-wall antigen with the molecular mass of 250 kDa (CM250) from both C. muris and C. parvum and also cross-react with C. serpentis. Occasionally, additional bands at 235, 150, 110, 92, 72, and 45 kDa in C. parvum can be also recognized by CMYL3. We believe that these smaller size antigens are likely derived from the proteolysis of the CM250 antigen, based on the fact that storage of the same protein sample has resulted in a shift in prominence from the 250- to the 150-kDa antigen (lane 3 of Fig. 2A). This interpretation is further supported by the detection of at least two types of proteases in sporozoites/oocysts (Nesterenko et al. 1995; Okhuysen et al.

1994), which have been proposed to play important roles in enhancing oocyst excystation during parasite infection (Forney et al. 1996).

Most oocyst-wall antigens reported so far, including CM250 in the present study, are glycosylated with the exception of an antigen recognized by the 2D7 antibody reported by Ward and Cevallos (1998). Bonnin et al. (1995) describe a mAb, OW-IGO, that also recognizes a 250 kDa antigen in C. parvum, as well as multiple lower molecular mass antigens. However, degradation of this 250 kDa antigen was not observed. Thus, there was no suggestion that degradation was a possible reason for the appearance of multiple bands. Furthermore, whether the OW-IGO antibody also recognizes the same antigen in C. muris remains unknown. The antigens recognized by OW-IGO localized to the parasitophorous vacuole membrane as well as the oocyst wall (Bonnin et al. 1991). In contrast, the CM250 recognized by CMYL3 and CMYL30 was only found in the structures originating from the parasite. These differences strongly suggest that our antibodies recognize a different wall antigen from that recognized by the OW-IGO antibody. Several mAbs against C. parvum oocyst antigens, with a molecular mass of approximately 40 kDa, have been previously reported. Anusz et al. (1990) described mAb 18.280.2, which recognized a 40-kDa molecule located in the exterior surface of C. parvum oocysts. Additionally, mAb 181B5 reacted with 41- and 44-kDa molecules of C. parvum (McDonald et al. 1991; Nina et al. 1992). MAb OW-IGO also recognized a 40-kDa antigen of the C. parvum oocyst wall (Bonnin et al. 1991). Nina et al. (1992) suggested that the 40-kDa antigen of C. parvum described by Anusz et al. (1990) may correspond to the 41-kDa band recognized by mAb 181B5. Furthermore, these two mAbs, 18.280.2 and 181B5, exhibited crossreactivity to the C. muris oocyst wall. Thus, it appears that more than one type of glycoprotein exists on the Cryptosporidium oocyst wall. This is in contrast to the composition of Eimeria oocyst walls, in which only one glycoprotein has been described (Karim et al. 1996; Stotish et al. 1978). The purpose and function of these oocyst antigens are not known and should be evaluated further.

The origin of oocyst-wall proteins is not completely understood, although several papers have attempted to

	$CMYL3$ (IgM)			$CMYL30$ (IgG1)		
	$C.$ muris	C. parvum	C. serpentis	$C.$ muris	C. parvum	C. serpentis
Immunofluorescence Immunogold	Oocyst wall	Oocyst wall	Oocyst wall	Oocyst wall	Oocyst wall	Oocyst wall
Oocyst	Wall	Wall	n.d.	Wall	Wall	n.d.
Macrogametocyte	Vesicles and cytoplasm	Vesicles and cytoplasm	n.d.	Vesicles and cytoplasm	Vesicles and cytoplasm	n.d.
Meront			n.d.	-		n.d.
Sporozoite			n.d.			n.d.
Antigen size ^a	250 kDa	250 kDa	n.d.	250 kDa	250 kDa	n.d.

Table 1 Characteristics of monoclonal antibodies produced against Cryptosporidium muris. – Negative, n.d. not done

^aAs determined by Western blot analysis

address this question. Current and Reese (1986) reported that the Cryptosporidium wall is derived from two types of electron-dense (i.e., loose and tight electron-dense) wall-forming bodies present in the macrogametocytes. On the other hand, Bonnin et al. (1991) suggested that oocyst-wall antigens of C. parvum, recognized by mAb OW-IGO, are released from the macrogametocyte electron-lucent vesicles into the vacuolar space and incorporated into the oocyst wall. In the present study, the CM250 antigens recognized by CMYL3 and CMYL30 were detected in electron-dense vesicles in the cytoplasm around electron-lucent amylopectin granules of macrogametocytes. The electron-dense vesicles heavily labeled by these mAbs were located along the periphery of the macrogametocyte cytoplasm. Thus, there is high possibility that the component of the oocyst wall originates from these electron-dense vesicles of the macrogametocyte.

MAbs CMYL3 and CMYL30 against C. muris oocysts show cross-reactivity with the oocyst wall of C. parvum as well as C. serpentis but do not cross-react with Eimeria oocysts, even though both Eimeria and Cryptosporidium belong to the same suborder. Thus, the antigen CM250 is Cryptosporidium oocyst-specific. Cysts from Giardia are known to cause similar diarrhea symptoms in humans as cryptosporidiosis (Roberts and Janovy 1996). The present study also showed that our CMYL antibodies did not recognize Giardia cysts. In addition, these mAbs work well on indirect immunofluorescence, ELISA, and Western blot analysis. Table 1 summarizes such analyses. These antibodies, therefore, could be very useful for the diagnostic detection of Cryptosporidium oocysts from various sources and for the biological study of the oocyst wall of Cryptosporidium.

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