

AN INNOVATIVE REPURPOSING OF MEFLOQUINE; ASSESSMENT OF ITS THERAPEUTIC EFFICACY IN TREATING *CRYPTOSPORIDIUM PARVUM* INFECTION OF BOTH IMMUNOCOMPETENT AND IMMUNOCOMPROMIZED MICE

By

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Abstract

Cryptosporidium parvum is a protozoan parasite can affect humans, worldwide, causing asymptomatic infections or diarrheal disease, which may be life-threatening in immunocompromised and neonatal individuals. Mefloquine is one of the most promising anti-parasitic drugs. The present report aimed to study the *in-vivo* efficacy of Mefloquine when applied in immunocompetent and immunocompromised cryptosporidiosis-infected mice groups, each of the them was subclassified into the following groups :non-infected non-treated group (normal control), infected non-treated group (infected control), nitazoxanide treated group and Mefloquine treated group. One week post infection, treated groups received either Nitazoxanide (100mg/Kg daily for 5 days) or single dose of Mefloquine (400mg/ Kg). Two weeks post treatment, all mice were scarified. Stool samples and intestinal histopathological specimens were examined. For both drugs, immunocompetent groups showed better parasitological clearance than immunocompromized one. *Cryptosporidium* oocyst reduction rates with Nitazoxanide (NTZ) and Mefloquine were 53.3%, and 61.6% respectively in the immunocompetent groups. The corresponding rates in immunocompromized groups were 49.93% and 60.03%.for NTZ and Mefloquine respectively. A single dose of Mefloquine treatment (400mg/kg) resulted in higher oocyst reduction rates than the approved anti-cryptosporidiosis drug (Nitazoxanide with five days application regimen).The histopathological study supported the parasitological findings as mefloquine treated mice tissue showed mild to moderate inflammatory changes while that of other groups ranged from moderate to severe alterations in the mice tissue. These results showed that Mefloquine which is FDA approved, already marketed and commercially available on a global scale has an excellent anti parasitic activity against *C. parvum* infection with single dose application; which saves time, cost and efforts to search for additional or alternative drugs for treating cryptosporidiosis. More large scale studies are needed to illustrate its dose response relationship using multiple doses regimens, performance and limitations on immunocompromized population, synergistic effect with already approved drugs , mechanism of action on *Cryptosporidium* parasite and its possible role in chemoprophylaxis specially for high risk individuals.

Key words: *Cryptosporidium parvum*, Mefloquine, Nitazoxanide, *In-Vivo*,

Introduction

Cryptosporidium parvum is an internationally distributed protozoan parasite that is existed in both vertebrates and invertebrates (Lima *et al*, 2011).Infections are transmitted by fecal-oral route, or through contaminated food or water, and many major waterborne outbreaks have occurred (Yoder and Beach, 2010). Cryptosporidiosis represents a foremost health problem as it was a frequent cause of diarrhea in both immunocompetent

and immunodeficient individuals (Rossignol, 2010), all over the world (Yoder *et al*, 2012; Insulander *et al*, 2013). Nitazoxanide is the only approved drug for treatment of diarrhea caused by *Cryptosporidium* infection (Mainali *et al*, 2013). But, it showed imperfect efficacy in the most vulnerable patients (Amadi *et al*, 2002; Abubakar *et al*, 2007; Manjunatha *et al*, 2016). Besides, the dependence of a one drug as sole effective treatment for any microorganism is known

to carry an inevitable risk of electing resistant strains and failure of treatment. Hence, development of new effective drug for cryptosporidiosis represents a pressing need. Drug repurposing (also termed re-profiling, re-tasking, therapeutic switching or drug repositioning) is the process of developing new indications for existing, failed or abandoned drugs or advanced clinical candidates (Sekhon, 2013). Drug repurposing was a useful strategy to accelerate drug development process due to lower costs, little risk and decreased time to market due to availability of preclinical data (Padhy and Gupta, 2011). This enables not only pharmaceutical companies but also public-sector researchers to engage in drug discovery and development efforts (O'Connor and Roth, 2005), and hence might result in treatment options for diseases almost exclusively addressed by public sector researchers, such as the neglected tropical diseases. Mefloquine is a 4-quinolinemethanol synthetic antimalarial analogue of quinine developed in 1971 (Kumar *et al.*, 2009). Years later, it proved to be more than antimalarial drug and gave excellent antiparasitic activities including antischistosomal one (Keiser *et al.*, 2010; Ingram *et al.*, 2012; Basra *et al.*, 2013; Xiao *et al.*, 2013; Abou-Shady *et al.*, 2016). Besides, antischistosomal activity, Keiser *et al.* (2010) reviewed the effectiveness of mefloquine against food-borne trematodes and highlighted that mefloquine was not active against *Fasciola hepatica* or *Clonorchis sinensis*, but was active against juvenile and adult *Opisthorchis viverrini* in a hamster infection (Keiser *et al.*, 2009a). But, mefloquine failed to be effective against *O. viverrini* in clinical trials (Soukhathammavong *et al.*, 2011). Also, mefloquine yielded significant results against *C. sinensis* infection in a rat model and *Paragonimus westermani* in a dog model, but was not better than praziquantel (Xiao *et al.*, 2010). Mefloquine proved activity against *Babesia* in man and dogs (El-Bahnasawy *et al.*, 2011; Munkhjargal *et al.*, 2012), *Echinococcus multilocularis* (Küster *et al.*,

2015) and adult *Fasciola gigantica* (Shalaby *et al.*, 2016).

The aim of this study was to explore the efficacy of mefloquine against cryptosporidiosis infection in a controlled study including both the immuno-competent and the immuno-compromized mice.

Materials and Methods

This study was carried out on laboratory bred Swiss albino female mice (n=80) weighing about 20gm. They were categorized into the following eight groups of ten mice each: GI: Immunocompetent, infected non treated mice, GII: Immunocompetent, infected animals treated with nitazoxanide orally 100mg/kg (Li *et al.*, 2003) daily for five consequent days, one week post infection, GIII: Immunocompetent-infected mice treated with mefloquine orally 400mg/Kg single dose (Keiser *et al.*, 2009b), a week post infection. GIV: Immunocompetent, non-infected non treated, GV: Immunocompromized, infected non treated mice, GVI: Immunocompromized, infected animals treated with nitazoxanide orally (0.04-mg/mouse/day) daily for five consequent days, a week post infection, GVII: Immunocompromized, infected mice treated with mefloquine orally 400mg/Kg single dose, a week post infection and GVIII: Immunocompromized, non-infected non-treated. The animals were sacrificed two weeks post treatment.

Immunosuppression was performed by giving the animals synthetic corticosteroids (dexamethasone) orally at a dose of 0.25 mg /g/day for 14 successive days prior to inoculation with *Cryptosporidium* oocysts (Rehg *et al.*, 1988). *Cryptosporidium* oocysts were obtained from naturally infected calves (slaughter houses) by collection of scrapings of the ileal mucous membrane and fecal content (Anderson, 1985). The samples were examined for confirmation of the presence of oocysts by modified Ziehl-Neelsen staining method (Henriksen and Pohlen, 1981). The infective samples were preserved by mixing with an equal volume of 2.5% Potassium dichromate (K₂Cr₂O₇), after Campbell and

Current (1983). Infective inoculum was prepared (Reese *et al*, 1982) and the number of oocysts in the concentrated stock inoculum was counted to determine how much the fluid volume was the inoculum per mouse.

All mice in the studied groups except the control groups were infected orally with the prepared inoculum of *Cryptosporidium* oocysts; this occurred on day 15 of dexamethasone in immunosuppressed groups (Moon *et al*, 1982). The animals were deprived of water overnight, and were then inoculated intraesophageally with the prepared inoculums using a tuberculin syringe connected to a polythene tube. The amount given to each mouse was adjusted to contain approximately 1000 oocysts/mouse (Gaafar, 2007). Two weeks post treatment, all mice were sacrificed. Stool samples and intestinal histopathology were examined. Mefloquine hydrochloride (Lariam 250mg tablets) was obtained from ROCH Co. Nitazoxanide (Nanazoxid100mg) was provided by Medizen Pharmaceutical Industries for Utopia Pharmaceuticals.

Assessment of infection and the drug effect: 1- Stool examination: Fresh fecal pellets from each mouse in the study groups were collected separately. Each sample was suspended in 10% formalin and homogenized. Then, 1mg was prepared as a fecal smear and stained by the modified Ziehl-Neelsen staining method. The stained fecal smear was examined microscopically and

number of oocysts was counted in 10 high-power fields (Mohamed *et al*, 2015).

2- Histopathological examination: The terminal 2 cm of the ileum was submitted to routine histopathological processing at the Department of Pathology, TBRI, where they were fixed in 10% neutral buffered formalin, dehydrated in ascending grades of ethanol, followed by immersion in xylene, and then impregnated in paraffin. Two 5-mm thick sections were taken from each block. One section was stained with hematoxylin and eosin for evaluation, and then examined by light microscopy for assessing the histopathological changes (Tzipori *et al*, 1981).

Statistical analysis: Data were collected, tabulated, and statistically analyzed using SPSS program version 11. ANOVA test was used for detection of variation between the groups. Post hoc Bonferoni test was used for pair wise comparisons between groups. Unpaired Student *t* test was used to assess if the host immune status influenced performance of drugs by comparing mean oocyst count of immunocompetent and immunocompromized groups. The percentage infection reduction rate was assessed using the formula: (mean value of infected untreated group - mean value of infected- treated group) x100/mean value of infected- untreated group (Abdel Sal-am *et al*, 2008).

A *p*-value equal to or less than 0.05 was considered significant and less than 0.01 was considered highly significant.

Results

Table 1: Comparison between means of oocyst number in infected non-treated control, NTZ & Mefloquine treated immunocompetent mice groups.

Oocyst count	infected control (N=10)	NTZ (N=10)	Mefloquine (N = 9)	<i>P</i> value	<i>F</i>
Mean +SD	33517.1+5320.1	15636.3+6300.4	12856.8+ 3061.7	<0.001	46,385
Range	24750–43333	1366.3–27000	8222.2–19200		

There was a highly significant difference between infected control mean oocysts count and that of NTZ and mefloquine treated immunocompetent mice, but mefloquine gave an observably lower mean oocyst count that reflected on mean oocysts count reduction rate of both groups; 53.3% for NTZ group and 61.6% for mefloquine treated one.

Pairwise comparison between groups using *Bonferroni* post hoc test showed no significant difference between mean *Cryptosporidium* oocysts count of NTZ and mefloquine treated immunocompetent mice *P* value= 0.750, denoting that the mefloquine efficacy was comparable to the NTZ with the upper hand for the former drug.

Table 2: Comparison between means of oocyst number in infected non treated control, NTZ and Mefloquine treated immunocompromized mice.

Oocyst count	Infected control (N=10)	NTZ(N=10)	Mefloquine (N=10)	P value	F
Mean \pm SD	46973.95 \pm 11018	23511.99 \pm 8117.1	18767.6 \pm 5742.3	<0.001	31.067
Range	34888.9–72000	13555.6–39000	10888.9–25600		

There was a highly significant statistical difference between infected control group mean *Cryptosporidium* oocysts count and that of NTZ and mefloquine treated immunocompromized mice groups, though mefloquine ones showed an observably lower mean oocysts count which was reflected on the mean oocyst count reduction rate of both groups; 49.9% for NTZ group and 60% for

mefloquine treated one. Pairwise comparison between various groups using *Bonferoni* post hoc test showed that there was no statistically significant difference between the mean *Cryptosporidium* oocysts count of NTZ and mefloquine treated immunocompetent mice P value=0.679, denoting that mefloquine efficacy was comparable to that of NTZ with the upper hand for the earlier.

Table 3: Comparison between NTZ and mefloquine in treating *C. parvum* infection by host immunity.

Immune status	Immuno competent	Immunocompromized	t	P value
NTZ	15636.3 \pm 6300.4	23511.99 \pm 8117.1	2.4238	0.0261
Mefloquine	12856.8 \pm 3061.7	18767.6 \pm 5742.3	2.7510	0.0136

In NTZ and mefloquine treated mice, oocyst mean counts were substantially impacted by host immune status as they were significantly higher in the immunocompromized groups treated by both drugs.

The histopathological examination in this study was performed just to roughly assess our hypothesis regarding the mefloquine anticryptosporial activity as manifested by the degree of inflammatory changes in small intestine tissue and not to measure its effect on the intensity of infection. As seen in figures (1, 2, 3 & 4), the histopathological screening study results supported that of parasitological ones. The infected non treated samples showed severe villous atrophy and inflammatory cells infiltrations. There was a mild to moderate inflammatory reaction in the mefloquine treated mice samples while the intensity of inflammation ranged from moderate to marked in the corresponding NTZ samples as compared with the control samples.

Discussion

Cryptosporidiosis may become life threatening which can lead to death in some individuals specifically children and immunosuppressed patients. There was no completely satisfactory treatment for the cryptosporidial enteritis successfully developed (Castella-

nos-Gonzalez *et al*, 2016). Consequently, probing for an alternate effective therapy is a worthy research area. Mefloquine therapeutic efficacy has been proved for many other parasites (Xiao *et al*, 2010; Munkhjargal *et al*, 2012; Basra *et al*, 2013; Xiao *et al*, 2013; Küster *et al*, 2015; Shalaby *et al*, 2016; Abou-Shady *et al*, 2016), but up till now to our knowledge, there are no studies in the literature about repurposing it for treatment of *Cryptosporidium parvum* parasite infection. In this study, both NTZ and mefloquine treatment resulted in a partial parasitological clearance of subjected mice as manifested by the highly significant statistical difference between the mean *C. parvum* oocyst count in the treated groups as compared by the infected non treated one. These findings were valid for both immunocompetent and immunocompromized mice groups. In both NTZ and mefloquine treated mice the mean *Cryptosporidium* oocyst count was significantly higher in immunocompromized. Theoretically, this could be due to the logic profound replication of *Cryptosporidium* parasites happens as a result of dexamethazone induced immunosuppression so as to this enormous number exceeds the top limit of antiparasitic capabilities of both drugs and eventually mounting the parasite yield upon

treatment. On the molecular level, the dexamethazone inhibitory effect on CD4 cells numbers (Parimi *et al*, 1999) was substantially contributing to this finding as the ultimate control of *Cryptosporidium* parasite infection in adult mice is dependent on CD4 T cells (Aguirre *et al*, 1994; Schmidt *et al*, 2001). Consequently, in face of this dexamethazone induced CD4 declining numbers, it seems that even potent antiprotozoal agents as NTZ and mefloquine cannot affect the parasite replicating and infecting activities as if the immune system is working properly. This agreed with Amadi *et al*. (2009) reported the attenuating effect of the immunosuppressed status caused by HIV infection in children on NTZ therapeutic efficacy, even after applying a high dose prolonged treatment regimen. Also, Amadi *et al*. (2002); Abubakar *et al*. (2007) and Manjunatha *et al*. (2016) demonstrated the NTZ inadequacy in presence of immunocompromization status. The therapeutic effect of NTZ on *Cryptosporidium* infection in mice was previously reported since long time by others (Theodos *et al*, 1998; Blagburn *et al*, 1998; Gargala *et al*, 2000). Interestingly, mefloquine treated mice showed a higher *Cryptosporidium* oocyst reduction rates than the NTZ in both immunocompetent and immunocompromized mice, though the difference was statistically insignificant, but in view of the difference in dosage regimen between mefloquine (single dose) and NTZ (five consequent doses), we can effortlessly expect much better performance for mefloquine if used in a repeated dosage system.

Generally, the histopathological study results supported the parasitological data, in mefloquine treated groups it showed mild to moderate degree of intestinal tissue inflammation, while that of NTZ ranged from the moderate to marked inflammatory changes. Being both protozoa, both belong to Apicomplexa Phylum; *Sporozoa* Class and *Coccidia* subclass, it could be hypothesized that the mechanism of mefloquine action on *Cryptosporidium* parasite might be related to

that exerted by it on *Plasmodium* parasites. *Plasmodium* spp. could synthesize several species of sphingolipids (cell membrane component) and retrieve others from their host cells (Coppens, 2013). A pioneering study of the lipid content in *P. falciparum*-infected red blood cells reported a significant increase in the sphingomyelin content compared with uninfected red blood cells, probably owing to the parasite's activities (Lawrence and Cenedella, 1969). Sphingomyelins are used by the parasite to form and stabilize a tubulovesicular network (TVN) of membranes that extends from the membrane of the parasitophorous vacuole to the erythrocytic plasma membrane (Haldar, 1996; Lauer *et al*, 1997). Studies on the mefloquine mode of action reported that its activities through sphingomyelinase enzyme led to the excess in the ceramide production (Olliaro and Wells, 2009) and decreased infectivity *in-vitro* and in humans (Lauer *et al*, 2000; Grellepois *et al*, 2005; Pankova-Kholmyansky and Flescher, 2006). The excess ceramide was detrimental to the parasite (Pankova-Kholmyansky *et al*, 2003). Sphingolipid-enriched membrane micro-domains (SEMs) were glycosylphosphatidylinositol (GPI)-anchored proteins found in cell membrane and involved in various membrane functions. A primary function of SEMs was thought to regulate the protein interactions by their ability to selectively recruit or exclude proteins as well as their ability to cluster into larger platforms (Manes *et al*, 2003). Interestingly, in another study on *Cryptosporidium* parasite, SEMs were found to have a role in facilitating parasite attachment to host cellular membrane and hence affecting their infectivity. They used an *in vitro* model of human biliary cryptosporidiosis, demonstrated that *C. parvum* infection triggers the clustering of SEM components at infection sites. Thus, disruption of SEM components or inhibition of SEM platform formation decreases *Cryptosporidium* attachment to and entry of cultured cholangiocytes. Also, the authors reported that clustering of SEM components at

infection sites appeared to be involved in the aggregation of Gal/GalNAc-associated membrane receptors and *Cryptosporidium*-induced activation of the PI-3K/Cdc42/actin signalling pathway, processes were associated with parasite attachment to host cells and subsequent cellular entry. Thus, SEMs are required for *C. parvum* attachment to and entry of the host cells, likely via clustering of membrane-binding proteins and facilitating of *C. parvum*-induced actin remodelling at infection sites via activation of the PI-3K/Cdc42 signalling pathway (Nelson *et al*, 2006). Since mefloquine proved to be a potent sphingomyelinase enzyme activator (Pankova-Kholmyansky *et al*, 2003), theoretically it could affect SEMs in which sphingomyelin is a basic constituent (Manes *et al*, 2003) and consequently, hinder the ability of *Cryptosporidium* parasite to attach, infect and form the tubulovesicular network necessary for its biology.

Conclusion

Mefloquine is an effective, well characterized and FDA approved alternative treatment for *C. parvum* infection. It showed a higher activity than the NTZ which is the only FDA approved for treatment of *C. parvum* infection. More large scale studies are needed to illustrate its dose response relationship using multiple doses regimens, performance and limitations on immunocompromised population, synergistic effect with already approved drugs, mechanism of action on *Cryptosporidium* parasite and its possible role in chemoprophylaxis specially for high risk individuals. Clinical trials could be instantly and effortlessly designed, approved and executed as it is already marketed drug.

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Explanation of figures

Fig.1: *Cryptosporidium parvum* oocysts count in immunocompetent groups for each individual mouse.

Fig. 2: *Cryptosporidium parvum* oocysts count in immunocompromized groups as recorded for each individual mouse.

Fig.3: Mean oocyst reduction rate of NTZ and mefloquine immunocompetent and immunocompromized mice.

Fig. 4: Histopathological changes of small intestine of studied groups A& B: Non infected control mice tissue showing normal villi structure and length, no inflammatory reaction. C &D: Infected non treated mice tissue showing acute inflammatory cells infiltrations with sever villous atrophy E,F&G: Mefloquine treated mice tissue showing mild to moderate acute inflam-

matory cells infiltrations with villous widening H& I : NTZ treated mice tissue showing moderate to marked acute inflammatory cells infiltrations with villous widening



