In vitro Synergism between Chloroquine and Antibiotics against *Orientia tsutsugamushi*

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**Background:** To investigate whether chloroquine enhances the effect of antibiotics against *Orientia tsutsugamushi*, the causative organism of scrub typhus, we compared the effect of antibiotics in combination with chloroquine with the effect of antibiotics alone in vitro.

**Materials and Methods:** The Boryong or AFSC-4 strain was inoculated into ECV304 cells, and incubated in medium containing doxycycline (4 µg/mL), rifampin (4 µg/mL), azithromycin (0.5 µg/mL), chloroquine (1 µg/mL), and each of these antibiotics in combination with chloroquine for 7 d. Immunofluorescence (IF) staining for *O. tsutsugamushi* was performed 4 hr and 7 d after inoculation of the bacteria, and IF-positive foci were enumerated.

**Results:** Chloroquine inhibited the growth of *O. tsutsugamushi* by 15.5%. In combination with chloroquine, the antimicrobial effects increased by 4.4% for doxycycline (a 92.9% reduction of bacterial numbers for doxycycline versus a 97.3% reduction for doxycycline plus chloroquine), 4.6% for rifampin (90.0% versus 94.6%), and 8.3% for azithromycin (86.9% versus 95.2%). The antimicrobial effect of the antibiotics alone was significantly different compared to the combined effect of antibiotics and chloroquine (Wilcoxon signed-rank test, \(P = 0.001\)).

**Conclusions:** The combined use of chloroquine with an antibiotic for the treatment of *O. tsutsugamushi* infections may be useful for increasing the efficacy of the antibiotics.

**Key Words:** Azithromycin; Chloroquine; Combination drug therapy; Doxycycline; Drug synergism; *Orientia tsutsugamushi*; Rifampin; Scrub typhus

**Introduction**

*Orientia tsutsugamushi* causes scrub typhus, followed by a chronic latent infection irrespective of appropriate antibiotic therapy [1-3]; the infection sometimes relapses, manifesting as myocarditis or pneumonia [4, 5]. Several clinical studies...
have revealed that treatment with doxycycline and chloramphenicol presents acceptable clinical responses, even though these antibiotics are bacteriostatic to *O. tsutsugamushi*. In 1996, it was reported that scrub typhus exhibits a delayed response to doxycycline treatment in northern Thailand [6]. To prevent this persistent infection and to treat those cases of scrub typhus that are relatively resistant to antibiotic therapy, treatment regimens with more potent therapeutic efficacy, ideally bactericidal treatments, should be investigated. This requires the investigation of new antibiotics. Azithromycin and rifampin are relatively new antibiotics for the treatment of scrub typhus. These antibiotics have demonstrated high therapeutic efficacies and no cases with relapse; rifampin, in particular, has a bactericidal effect on certain bacteria [7, 8]. A previous *in vitro* experiment, however, indicated that these antibiotics are also bacteriostatic against *O. tsutsugamushi* [9].

A combination antibiotic therapy may also be considered for this purpose, although an *in vitro* experiment revealed that this modality is not more effective than a single antibiotic [10]. Chloroquine has been used as an anti-malarial agent for a long time. In addition, it has been shown to increase the antimicrobial efficacy of doxycycline for the treatment of Q fever [11] and Whipple's disease [12]. This activity is thought to be mediated by the alkalization of phagosomes [13]. Chloroquine enters cells, accumulates within the acidic components such as endosomes, phagosomes, and Golgi vesicles, and increases the pH of these vesicles. *O. tsutsugamushi* uses phagosomes to enter phagocytes and then escapes from phagosomes to release into the cytosol during the early stages of an infection. When these vesicles are alkalized by NH$_4$Cl or bafilomycin A, an alkalizing agent for acidic vesicles, the escape from the endocytic pathway and the infectivity of the bacterium are markedly reduced [14]. Thus, it may be hypothesized that chloroquine inhibits replication of *O. tsutsugamushi* and increases the antimicrobial activity of anti-rickettsial antibiotics. To verify this hypothesis, we performed an *in vitro* study to evaluate the antimicrobial effect of doxycycline, rifampin, and azithromycin, both individually and in combination with chloroquine against *O. tsutsugamushi*.

**Materials and Methods**

1. **Infection of ECV304 cells with *O. tsutsugamushi***

The ECV304 cell line, which has many characteristics similar to endothelial cells [15], was used for this study. ECV304 cell monolayers were cultured on polyethylene terephthalate (PET) cover slips in a 6-well (9.08 cm$^2$) PET culture plate (CorePath, Daegu, Korea). A PET cover slip was estimated to contain 1.5 × 10$^6$ ECV304 cells. The cell cultures were maintained in M199 medium (pH 7.2) (Gibco BRL, Gaithersburg, MD, USA) supplemented with 10% heat-inactivated fetal bovine serum (Gibco BRL) at 37°C in a humidified atmosphere containing 6% CO$_2$. The culture medium was changed every 3 to 4 days.

The Boryong and AFSC-4 strains of *O. tsutsugamushi* were used in this study. The former strain is the most prevalent genotype in Korea and is susceptible to doxycycline. The minimal inhibitory concentrations for this strain, which were measured by a flow cytometric method, ranged from 0.05-0.1 µg/mL for doxycycline, 0.025-0.05 µg/mL for rifampin, and 0.1-0.05 µg/mL for azithromycin [16, 17]. The latter strain was isolated from Thailand and reported to be insensitive to doxycycline [18]. Stock solutions of each strain were inoculated onto monolayers of ECV304 cells seeded in 75-cm$^2$ culture flasks. When the infected ECV304 cells presented maximum cytotoxic effects, the degree of infection was assessed by immunofluorescence (IF) staining. The cells were then disrupted with glass beads and centrifuged at 350 × g for 5 min. The resultant supernatant was divided equally into 9 aliquots, inoculated onto ECV304 cell monolayers grown on the PET cover slips, and incubated for 4 hr. At the end of the initial incubation period, the inocula were replaced with fresh medium containing one of the following treatments: 4 µg/mL of doxycycline (Sigma Co., St. Louis, MO, USA), 4 µg/mL of rifampin (Sigma), 0.5 µg/mL of azithromycin (Sigma), 1 µg/mL of chloroquine (Sigma), doxycycline plus chloroquine, rifampin plus chloroquine, or azithromycin plus chloroquine. The concentrations used were the expected serum concentrations that can be achieved after administration of conventional doses of the antibiotics, as employed in a previous study [19]. The medium was replaced with fresh medium containing the same antibiotic or its respective combination with chloroquine on day 4 post-inoculation with *O. tsutsugamushi* (pi0). Uninfected ECV304 cells and cultures containing no antibiotic or chloroquine were used as controls. On day 8 pi0, the ECV304 cells were washed 2 times with phosphate buffered saline (PBS), and IF staining for *O. tsutsugamushi* was performed. The above experiments were performed in triplicate in Boryong-infected cells and in duplicate in AFSC-4-infected cells.

2. **Immunofluorescence staining**

The ECV304 cells infected with each strain of *O. tsutsugamushi* were fixed with acetone and treated with PS15, a
### Results

#### Statistical analysis

Continuous variables were expressed as the median (IQR). A Wilcoxon signed-rank test was used for comparisons of 2 paired values, and Mann-Whitney U tests were used for comparisons of independent values. All P values for comparisons of 2 independent values were two-tailed and statistical significance was defined as P < 0.05.

#### Immunofluorescence analysis

To clearly define O. tsutsugamushi, the host cells were stained with 0.003% Evans Blue in PBS. After the slides were rinsed, they were placed in a moist chamber, and finally rinsed 3 times with PBS. To further identify O. tsutsugamushi cells, the 56-kDa major outer membrane protein was defined as a positive focus on the 56-kDa major outer membrane protein. (Zeiss, Germany) with a confocal laser scanning system (Bio-Rad Laboratories, USA). For CD172a, a mouse monoclonal antibody that reacts with the linear epitope on the 56-kDa major outer membrane protein (Euroimmun, Germany) was used for confocal microscopic analysis. The host cells were stained with 0.003% Evans Blue in PBS. After the slides were rinsed, they were placed in a moist chamber, and finally rinsed 3 times with PBS. To further identify O. tsutsugamushi cells, the 56-kDa major outer membrane protein was defined as a positive focus on the 56-kDa major outer membrane protein. (Zeiss, Germany) with a confocal laser scanning system (Bio-Rad Laboratories, USA). For CD172a, a mouse monoclonal antibody that reacts with the linear epitope on the 56-kDa major outer membrane protein (Euroimmun, Germany) was used for confocal microscopic analysis. The host cells were stained with 0.003% Evans Blue in PBS. After the slides were rinsed, they were placed in a moist chamber, and finally rinsed 3 times with PBS. To further identify O. tsutsugamushi cells, the 56-kDa major outer membrane protein was defined as a positive focus on the 56-kDa major outer membrane protein. (Zeiss, Germany) with a confocal laser scanning system (Bio-Rad Laboratories, USA). For CD172a, a mouse monoclonal antibody that reacts with the linear epitope on the 56-kDa major outer membrane protein (Euroimmun, Germany) was used for confocal microscopic analysis. The host cells were stained with 0.003% Evans Blue in PBS. After the slides were rinsed, they were placed in a moist chamber, and finally rinsed 3 times with PBS. To further identify O. tsutsugamushi cells, the 56-kDa major outer membrane protein was defined as a positive focus on the 56-kDa major outer membrane protein. (Zeiss, Germany) with a confocal laser scanning system (Bio-Rad Laboratories, USA). For CD172a, a mouse monoclonal antibody that reacts with the linear epitope on the 56-kDa major outer membrane protein (Euroimmun, Germany) was used for confocal microscopic analysis. The host cells were stained with 0.003% Evans Blue in PBS. After the slides were rinsed, they were placed in a moist chamber, and finally rinsed 3 times with PBS. To further identify O. tsutsugamushi cells, the 56-kDa major outer membrane protein was defined as a positive focus on the 56-kDa major outer membrane protein. (Zeiss, Germany) with a confocal laser scanning system (Bio-Rad Laboratories, USA). For CD172a, a mouse monoclonal antibody that reacts with the linear epitope on the 56-kDa major outer membrane protein (Euroimmun, Germany) was used for confocal microscopic analysis. The host cells were stained with 0.003% Evans Blue in PBS. After the slides were rinsed, they were placed in a moist chamber, and finally rinsed 3 times with PBS.
more IF-positive foci in AFSC-4-infected cells (Mann–Whitney U test, $P = 0.08$) in medium containing neither antibiotics nor chloroquine. Because the inoculated size was somewhat variable among experiments and the AFSC-4 strain was greater in number than the Boryong strain, the observed values were converted to percentages of inhibition (calculated as the observed values divided by the corresponding bacterial numbers in cultures containing no antibiotic on day 8 piO). The percentage of bacteria (with respect to cultures containing no antibiotic) was significantly reduced by 15.5% (95% CI: 13.2 to 18.1%) after the addition of chloroquine (Wilcoxon signed-rank test, $P = 0.043$) (Fig. 1).

When each of the antibiotics was added to the PET disc, the median percentages of bacteria were reduced by 92.9%, 90.0%, and 86.9% in doxycycline-, rifampin-, and azithromycin-treated cells, respectively. When the antibiotics were combined with chloroquine, the median percentages of IF-positive foci decreased further by 4.4%, 4.6%, and 8.3% in doxycycline- (resulting in a 97.3% reduction by doxycycline and chloroquine), rifampin- (94.6%), and azithromycin- (95.2%) treated cells, respectively. The difference for two pooled groups (all antibiotics individually and all combined treatments) was statistically significant (Wilcoxon signed-rank test, $P = 0.001$). Every pair-wise comparison between the antibiotic and its respective combination with chloroquine was also significantly different (Wilcoxon signed-rank tests, $P$-values $= 0.043$) (Fig. 1).

The median percentage of bacterial cells after the treatment with antibiotics alone was 206.2% (95% CI: 167.6 to 251.1%) larger than after treatment with the antibiotic and chloroquine combination (9.9 [6.1-12.4] versus 4.8 [2.5-6.9]). The median number of bacteria in AFSC-4 strain-infected cells treated with the antibiotic and chloroquine combinations was nearly equal to that observed for the Boryong strain-infected cells treated with the antibiotics alone (7.0 [5.4-7.8] versus 7.6 [4.9-11.3]) (Mann–Whitney U test, $P = 0.56$).

**Discussion**

In the present study, chloroquine was observed to have a mild inhibitory effect and increased the effect of antibiotics against *O. tsutsugamushi*. The growth of *O. tsutsugamushi* was inhibited by chloroquine itself by 15.5%, consistent with previous results for *Coxiella burnetii* (6.1%) and less extreme than the inhibition observed for *Listeria monocytogenes* and *Legionella pneumophila* (45–80%) [19, 21, 22]. These organisms differ in their endocytic pathways: *O. tsutsugamushi*, *Rickettsia* species, and *L. monocytogenes* escape from phagosomes before fusion with the lysosomes and multiply in the cytosol; *L. pneumophila* inhibits phagolysosomal fusion and multiplies in the phagosomes; lastly, *C. burnetii* survives and multiplies in the phagolysosomes [13]. When phagosomes fuse with lysosomes, the pH of these vesicles becomes more acidic, from 6.5 in phagosomes to 4.5 in phagolysosomes [19]. Because chloroquine alkalizes these acidic vesicles and augments the antibacterial effects of the antibiotic, it is expected that chloroquine more potently suppresses *C. burnetii* than *O. tsutsugamushi*; however, the inhibition of *C. burnetii* growth by chloroquine was not as great when compared with *O. tsutsugamushi*. It is not certain whether this finding is due to differences in experimental designs or due to characteristics of the organisms.

Despite its rather low intrinsic antibacterial effect, chloroquine increased the antibacterial effect of various antibiotics against *O. tsutsugamushi*, as was previously shown in other organisms including *C. burnetii* [19, 21, 22]. The increases in the antimicrobial effect against *O. tsutsugamushi* after the addition of chloroquine were larger than the sum of the independent antibiotic activities, indicating that the combinations were synergistic. This synergistic effect suggests several therapeutic implications; for instance, the combination of an antibiotic and chloroquine is one option for the treatment of scrub typhus when it responds poorly to doxycycline therapy.
The AFSC-4 strain in cultures containing neither antibiotic nor chloroquine showed 126.7% and 123.3% larger bacterial populations than the Boryong strain at 4 hr and 7 days piO, respectively. Similar among-strain differences were also shown in other studies where the AFSC-4, C1, and C27 strains, which are reported to be insensitive to doxycycline, exhibited a 10–18% higher inoculum number than the Karp strain, an antibiotic-susceptible strain [6, 18]. These differences in the inoculum numbers may be due to higher in vitro infectivity of these strains, as shown in the AFSC-4 strain [23]. Thus, if this high in vitro infectivity is associated with delayed defervescence in the treatment of scrub typhus, a 206.2% difference between the bacterial numbers in cultures containing the antibiotic combined with chloroquine and those with the antibiotic alone indicates that this combination could be used to treat scrub typhus caused by doxycycline-insensitive strains. This assumption is supported by data indicating that the bacterial numbers in AFSC-4 strain-infected cells, treated with combinations of an antibiotic and chloroquine, were nearly equal to those in the Boryong strain-infected cells treated with the antibiotics alone. Another potential application of the antibiotic and chloroquine combination treatments may be the rapid resolution of scrub typhus. Because the amount of O. tsutsugamushi organisms in severe scrub typhus is larger than in mild scrub typhus [24], a rapid reduction in the amount of bacteria in critically ill patients may result in more favorable clinical outcomes. Lastly, these combinations may be useful in the management of chronic O. tsutsugamushi infections, if these conditions are identified in the near future. Q fever endocarditis is an example of a chronic intracellular bacterial infection that is managed with doxycycline plus chloroquine [11]. Therefore, by combining antibiotics with chloroquine, higher efficacies may be achieved against O. tsutsugamushi infections. Alternatively, the same efficacy may be achieved with reduced doses of antibiotics and thereby this regimen may reduce the incidence of adverse effects during long-term treatment.

Although there is a previous report that a traveler taking chloroquine as a malaria prophylaxis contracted scrub typhus and improved within 48 hours after taking doxycycline [25], the theoretical benefits of the antibiotic combined with chloroquine must be investigated in clinical trials. This case also suggests that previous clinical trials conducted in malaria endemic areas might include patients with scrub typhus who were taking chloroquine as a malaria chemoprophylaxis. Chloroquine at a dose of 300 mg per week, a typical regimen for the prophylaxis of malaria, results in a plasma concentration of 1.6 µg/mL or more [26]. Therefore, these studies may estimate better responses to an anti-rickettsial agent. For example, Sheehy et al. [27] evaluated the clinical efficacies of chloramphenicol and tetracycline in United States servicemen with scrub typhus who had been deployed during the Vietnam War and were likely administered chloroquine prophylaxis; therefore, this study may overestimate the antibiotic efficacies and underestimate relapse rates.

Some differences were observed between the 3 antibiotics with respect to their efficacy in combination with chloroquine. In particular, when azithromycin was combined with chloroquine, O. tsutsugamushi growth was more strongly inhibited than when chloroquine was combined with doxycycline or rifampin, which could be attributed to increased antimicrobial activity of azithromycin in the alkaline pH of the growth medium.

Because there were discordant values between the bacteria at 4 hr and those on day 8 piO in cultures lacking antibiotics, some variation may be due to counting error, particular when the bacteria are high in number (>300 per field). In this instance, IF-positive dots were confluent, so it may have been difficult to accurately enumerate IF positivity; however, when the number was small, the counting was rather accurate. Thus, the effects of various antibiotics must account for variations in the denominator (i.e., the bacterial numbers in cultures containing no antibiotic on day 8 piO), but the comparison of the antimicrobial effect, i.e., the numerator, between the antibiotics and their combinations with chloroquine is reliable.

**Conflicts of Interest**

No conflicts of interest.

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