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## Interaction of *Escherichia coli* K1 and K5 with *Acanthamoeba castellanii* Trophozoites and Cysts

#### Abdul Matin<sup>1</sup> and Suk-Yul Jung<sup>2,\*</sup>

<sup>1</sup>Institute of Biomedical & Genetic Engineering, PO Box 2891, Islamabad, Pakistan; <sup>2</sup>Department of Biomedical Laboratory Science, Molecular Diagnosis Research Institute, Namseoul University, Cheonan 331-707, Korea

**Abstract:** The existence of symbiotic relationships between *Acanthamoeba* and a variety of bacteria is well-documented. However, the ability of *Acanthamoeba* interacting with host bacterial pathogens has gained particular attention. Here, to understand the interactions of *Escherichia coli* K1 and *E. coli* K5 strains with *Acanthamoeba castellanii* trophozoites and cysts, association assay, invasion assay, survival assay, and the measurement of bacterial numbers from cysts were performed, and nonpathogenic *E. coli* K12 was also applied. The association ratio of *E. coli* K1 with *A. castellanii* was 4.3 cfu per amoeba for 1 hr but *E. coli* K5 with *A. castellanii* was 1 cfu per amoeba for 1 hr. By invasion and survival assays, *E. coli* K5 was recovered less than *E. coli* K1 but still alive inside *A. castellanii*. *E. coli* K1 and K5 survived and multiplied intracellularly in *A. castellanii*. The survival assay was performed under a favourable condition for 22 hr and 43 hr with the encystment of *A. castellanii*. Under the favourable condition for the transformation of trophozoites into cysts, *E. coli* K5 multiplied significantly. Moreover, the pathogenic potential of *E. coli* K1 from *A. castellanii* cysts exhibited no changes as compared with *E. coli* K1 from *A. castellanii* trophozoites. *E. coli* K5 was multiplied in *A. castellanii* trophozoites and survived in *A. castellanii* cysts. Therefore, this study suggests that *E. coli* K5 can use *A. castellanii* as a reservoir host or a vector for the bacterial transmission.

Key words: Acanthamoeba castellanii, Escherichia coli K1, Escherichia coli K5, association, invasion, survival

### INTRODUCTION

Acanthamoeba is a free-living protozoan pathogen that has been isolated from diverse environments, including air, soil, tap water, swimming pools, and is known to be one of the most ubiquitous protozoans [1-3]. With the wide environmental distribution, they must effectively interact with a range of microorganisms to ensure their survival. It is not surprising that *Acanthamoeba* have been shown to interact with viruses, bacteria, algae, yeast, and other protists [4-7]. However, it is the ability of *Acanthamoeba* interacting with host bacterial pathogens that has gained most attention. For example, several studies have shown that *Acanthamoeba* act as a reservoir for a variety of bacterial pathogens, including *Escherichia coli* K1 (causative agent of meningitis) [1,8,9], *Legionella pneumophila* (causative agent of Legionnaire's disease) [10], *Coxiella burnetii* (causative

\* Corresponding author (syjung@nsu.ac.kr)

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agent of Q fever) [11], *Pseudomonas aeruginosa* (causative agent of keratitis) [12], *Vibrio cholerae* (causative agent of cholera) [13], *Helicobacter pylori* (causative agent of gastric ulcers) [14], *Listeria monocytogenes* (causative agent of listeriosis) [15], *E. coli* O157 (causative agent of hemolytic uremic syndrome) [16], and *Mycobacterium avium* (causative agent of respiratory infections) [17] and may act as vectors to transmit these pathogens to susceptible hosts.

The endosymbiotic relationship of bacteria with amoeba may serve (i) to protect bacteria in hostile environments, (ii) the amoebic intracellular environment might assist bacteria to adapt to survival in mammalian phagocytic cells, suggesting that amoeba-bacteria are involved in complex interactions. Further complexity is attributed to the fact that bacteria are a favourable food source for *Acanthamoeba*. What determines amoeba to act as a host for bacteria and at other times to consume bacteria remains incompletely understood. The fate of bacteria inside *Acanthamoeba* may be most likely dependent on the virulence properties of bacteria and/or *Acanthamoeba*. Based on this, it is hypothesized that the ability of bacteria to resist amoebic killing may have led to their evolution to produce human diseases, i.e., evade human immune cells, such as

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phagocytes. Therefore, amoebae may be the missing link to bacteria that are distributed in the environment and those causing human infections.

Most bacteria contain some sort of a polysaccharide layer outside the cell wall or outer membrane. In a general sense, this layer is called a capsule. Capsules are known to protect bacteria from engulfment by predatory phagocytes and from attack by antimicrobial agents. As related with previous studies, invasive E. coli K1 interacted with A. castellanii trophozoites more than non-invasive E. coli K12, which meant more association, invasion, and survival of E. coli K1 with A. castellanii [1,8]. In particular, a capsule-deletion mutant of E. coli K1 (lacking the neuDB genes cluster that is necessary for the production of cytoplasmic precursors to the exopolysaccharide capsule) exhibited significantly reduced association compared with the wild type strain and limited ability for invasion/uptake by and survival inside Acanthamoeba. The bacterial capsule would be a very important factor for the interactions with amoeba. However, the importance of a somatic antigen as other virulence factor in E. coli is not clearly understood. E. coli K5 is a streptomycin-resistant strain of the serotype O7:K1 [18]. E. coli K1 and E. coli K5 possess different somatic antigens but an identical capsular antigen.

Here, to understand *E. coli*-interactions with amoeba using *E. coli* strains possessing different somatic antigens, we compared and analyzed the *E. coli* K1 and *E. coli* K5 interactions with *A. castellanii* using association, invasion, and intracellular survival assays in *A. castellanii* trophozoites and cysts.

## MATERIALS AND METHODS

#### Culture of Acanthamoeba trophozoites and E. coli

All chemicals were purchased from Sigma Laboratories (Poole, Dorset, England), unless otherwise stated. A clinical isolate of *A. castellanii* belonging to T4 genotype, isolated from a keratitis patient (American Type Culture Collection, ATCC 50492) was used. *A. castellanii* isolate was grown without shaking in 15 ml of PYG medium [proteose peptone 0.75% (w/v), yeast extract 0.75% (w/v), and glucose 1.5% (w/v)] in T-75 tissue culture flasks at 30°C as previously described [19]. The media were refreshed 17-20 hr prior to experiments. This resulted in more than 95% amoebae in the trophozoite forms, which were subsequently used in experiments. A laboratory non-invasive *E. coli* strain HB101 (K12) was used as a non-pathogenic bacteria. The *E. coli* K1 strain E44 used in the present study is a rifampicin-resistant mutant of strain RS218 (serotype O18:K1). This strain is a clinical isolate derived from the cerebrospinal fluid (CSF) of a neonate with meningitis. In particular, *E. coli* K5 is a streptomycin-resistant strain of serotype O7:K1 and were cultured with streptomycin (25 µg/ml) at 37°C overnight before use.

#### E. coli association assay

To study E. coli interactions with live Acanthamoeba, an association assay was performed as was done in a previous report [9]. Briefly, Acanthamoeba were prepared in 24-well plates. The cells were washed once with PBS and incubated with E. coli strains  $(2 \times 10^6 \text{ cfu/well/0.5 ml of PBS})$ . The plates were incubated for 1 hr at room temperature. Following this incubation, amoebae were washed with PBS for 3 times to remove nonadherent bacteria and amoebae counted using a hemacytometer. Finally, amoebae were lysed by adding SDS (0.5% final conc.) to each well for 30 min, and the number of bacteria was enumerated by plating on nutrient agar plates. The percent bacterial association was calculated as follows: recovered E. coli (cfu)/total E. coli (cfu)  $\times$  100 = % E. coli associated with Acanthamoeba. In addition, the ratio of bacteria to amoebae was calculated as follows: recovered E. coli (cfu)/number of A can tham oeba = E. coli/A can tham oeba ratio.

#### E. coli invasion assay

To determine the ability of bacteria to invade or be taken up by *Acanthamoeba*, an invasion assay was performed. Briefly, amoebae were grown to confluency in 24-well plates followed by the addition of  $2 \times 10^6$  of *E. coli* as described above. After 1 hr incubation, *Acanthamoeba* were washed with PBS for 3 times, followed by an addition of gentamicin (100 µg/ml, final concentration, for 45 min) to kill extracellular bacteria. Finally, amoebae were counted, and the intracellular bacteria were enumerated as described above. The percent bacterial invasion/uptake was calculated as follows; recovered *E. coli* (cfu)/ total *E. coli* (cfu) × 100 = % intracellular *E. coli*. In addition, the ratio of bacteria to amoebae was calculated as follows; recovered *E. coli* (cfu)/number of *Acanthamoeba*=*E. coli/Acanthamoeba* ratio.

#### Intracellular survival assay

To determine the long-term effects of *Acanthamoeba* and *E. coli* interactions, an intracellular survival assay was performed. Briefly, amoebae were incubated with *E. coli*, followed by an

addition of gentamicin for 45 min. After this incubation, *Acanthamoeba* were washed for 3 times with PBS and subsequently incubated in 0.5 ml of PBS for 24 hr at 30°C. Finally, amoebae and *E. coli* were enumerated as described above and intracellular bacteria calculated as follows: recovered *E. coli* (cfu)/total *E. coli* (cfu) × 100 = % intracellular *E. coli* after 24 hr in PBS. In addition, the ratio of bacteria to amoebae was calculated as follows: recovered *E. coli* (cfu)/number of *Acanthamoeba*=*E. coli*/ *Acanthamoeba* ratio after 24 hr in PBS.

#### Encystment assay

An encystment assay was performed as previously described [20]. In the invasion assay, 24-well plates were added with gentamicin and then washed with PBS mentioned above. All mixture from a 24-well plate was transferred into a 1.5 ml eppendorf tube. After the tube was centrifuged, the supernatant was removed, and 50  $\mu$ l PBS was added into the tube. Then, the 50  $\mu$ l mixture was inoculated on 3% non-nutrient agar, that is, purified agar (Oxoid, Basingstoke, Hampshire, England) plates. The plates were incubated at 30°C for up to 48 hr. The encystment of *Acanthamoeba* was periodically observed under a light microscope.

#### Bacterial survival from Acanthamoeba cysts induction

After non-nutrient agar surfaces were soaked with 10 ml of PBS by shaking, cysts on the non-nutrient agar plates were gently scraped-off the agar surface using a cell scraper. Cysts were collected by centrifugation at 3,500 rpm for 10 min, washed 3 times with PBS, transferred into a 1.5 ml eppendorf tube with 700 µl of PBS and used in subsequent survival assays. One hundred µl of 700 µl mixture was added into a 24-well plate for a 0 hr survival assay, followed by the addition of SDS (0.5% final concentration) to solubilize any remaining trophozoites as described above. Two 100 µl of 600 µl mixture was added into a 24-well plate containing 900 µl of PYG media for a 43 hr survival assay at 30°C. Two 200 µl of 400 µl mixture was added into a 24-well plate containing PYG and gentamicin (100 µg/ml) for 45 min at 30°C. At the experimental time point, SDS (0.5%) was added to solubilize any remaining trophozoites as described above. Finally, amoebae and E. coli were enumerated as described above and intracellular bacteria calculated as follows: recovered E. coli (cfu)/total E. coli (cfu)  $\times$  100 = % intracellular E. coli after 22 hr and 43 hr in PYG. In addition, the ratio of bacteria to amoebae was calculated as follows: recovered E. coli (cfu)/number of Acanthamoeba=E. coli/Acanthamoeba ratio after 22 hr and 43 hr in PYG.

## RESULTS

*E. coli* K5 exhibited significantly lower association with *A. castellanii* as compared to the invasive *E. coli* K1

To determine the ability of *E. coli* K5 association with *A. castellanii*, the association assay was performed. Our findings revealed that the invasive *E. coli* K5 exhibited significantly lower





association with *A. castellanii* as compared to the invasive *E. coli* K1 for 1 hr (Fig. 1) (P < 0.05). Otherwise, the association of *E. coli* K5 was very similar to the non-invasive *E. coli* K12. Next, to determine the ratio of *E. coli* with *A. castellanii*, amoebae were enumerated prior to bacterial counting. Our results revealed that the ratio of *E. coli* K1 with *A. castellanii* was 4.3 cfu per amoeba (0.9 cfu per amoeba for K12) for 1 hr. In particular, the ratio of *E. coli* K5 with *A. castellanii* was 1 cfu per amoeba for 1 hr. *E. coli* K5 exhibited significantly lower association with *A. castellanii* as compared to *E. coli* K1, suggesting that the somatic antigen but not a capsular antigen might be an important factor that interacted with *A. castellanii* trophozoites.

## *E. coli* K5 exhibited significantly decreased invasion into or being uptaken by *A. castellanii* as compared to the invasive *E. coli* K1

Next, to determine the *E. coli* intracellular of amoebae, the invasion assay was performed. We observed a higher recovery (0.008 bacterial colony per amoeba) of *E. coli* K1 intracellular of *A. castellanii* as compared to K12 (no bacterial colony per amoeba) for 1 hr (Fig. 2) (P < 0.05). Interestingly, *E. coli* K5 was recovered less (0.00011 bacterial colonies per amoeba) than *E. coli* K1 but still alive inside *A. castellanii* for 1 hr (P < 0.05).

# *E. coli* K5 survived intracellularly in *A. castellanii* while non-invasive *E. coli* K12 were killed

To determine the fate of *E. coli* in long-term interactions with *A. castellanii*, the intracellular survival assay was performed by incubating *E. coli* with *Acanthamoeba* in PBS for 24 hr. With regard to the clinical isolates of *Acanthamoeba*, our findings revealed that once intracellular, *E. coli* K1 remained viable and multiplying, while K12 were killed (Fig. 3) (P < 0.05). Interestingly, *E. coli* K5 survived and multiplied a little intracellularly in *A. castellanii*. It is suggested that intracellularly invaded *E. coli* K5 multiply in the cytoplasm of *A. castellanii*.

## *E. coli* K5 survived inside the cyst form of *A. castellanii* as compared with *E. coli* K1

After *A. castellanii* was incubated with bacteria for 1 hr and then treated with gentamicin for 45 min, all *Acanthamoeba* trophozoites were inoculated onto non-nutrient agar plate to induce encystment of *A. castellanii*. The encystment was observed periodically up to 43 hr. The groups of *A. castellanii* cysts were grape-shaped as well as tightly linked (Fig. 4). Invasive *E. coli* K1 survived intracellularly and/or extracellularly in *A. castellanii* at 0 hr (0.052 bacterial colony per amoeba; 0.044 cfu per amoeba) (Fig. 5) (P<0.05). Otherwise, *E. coli* K12 and *E. coli* K5 didn't survive intracellularly and/or extracellularly in *A. castellanii* at 0 hr. To determine whether a very few number of *E.* 



Fig. 2. E. coli K5 exhibits significantly lower invasion and/or uptake by A. castellanii as compared to the invasive E. coli K1. To determine the ability of E. coli to invade and/or be taken up by A. castellanii, the invasion assay was performed. 'A' represents bacterial association with A. castellanii, while 'B' represents cfu per amoeba. Results are representative of 3 independent experiments performed in triplicate. Asterisks indicate a significant difference, i.e., P<0.05, using a paired t-test, one-tail distribution.



**Fig. 3.** *E. coli* K5 survives a little intracellularly while K12 are killed under unfavourable conditions. To determine the ability of *E. coli* to survive intracellularly, the survival assay was performed. 'A' represents bacterial association with *A. castellanii*, while 'B' represents cfu per amoeba. Results are representative of 3 independent experiments performed in triplicate. Asterisks indicate a significant difference, i.e., *P* < 0.05, using a paired *t*-test, one-tail distribution.



Fig. 4. Encystment of *A. castellanii*. Of the invasion assay, after gentamicin was treated to *A. castellanii* for 45 min, all *A. castellanii* were inoculated onto non-nutrient agar plate to induce encystment of *A. castellanii* up to 43 hr. It was observed that *A. castellanii* cysts formed grape-shaped group linked tightly. 'A' represents *A. castellanii* cysts observed with ×200 magnification power. Inset box was observed with ×400 magnification power under a light microscope in 'B'.

*coli* K5 may survive intracellularly in *A. castellanii*, cyst forms of *A. castellanii* were incubated under a favourable condition of PYG medium up to 43 hr, which would exhibit transformation of cysts into trophozoites and bacteria might come out of *A. castellanii* trophozoites. Under the favourable condition, no non-invasive *E. coli* K12 were come up but *E. coli* K5 multiplied significantly up to estimated 43 hr. In addition, the number of *A. castellanii* trophozoites was not significantly increased during 22 hr and 43 hr more than 0 hr (data not shown). Very

interestingly, *E. coli* K5 multiplied and survived in transformed *A. castellanii* trophozoites under favourable conditions, suggesting that *E. coli* K5 in *A. castellanii* cysts be alive and come out of trophozoites under favourable conditions.

Pathogenic potential of *E. coli* K1 from *A. castellanii* cysts exhibited no changes as compared with *E. coli* K1 from *A. castellanii* trophozoites

To determine whether invasive E. coli K1 from A. castellanii



**Fig. 5.** *E. coli* K5 survived extracelluarly in *A. castellanii* cysts. Encystment of *A. castellanii* was performed as mentioned above. For 22 hr and 43 hr survival assays under favourable conditions of PYG media, the cysts were treated with gentamicin for 45 min to remove extracellular *E. coli*. 'A' and 'B' represent the survival of *E. coli* from *A. castellanii* cysts at 0 hr with treatment of gentamicin. 'C' and 'D' represent the survival of *E. coli* from *A. castellanii* cysts at 22 hr under favourable conditions of PYG media. 'E' and 'F' represent the survival of *E. coli* from *A. castellanii* cysts at 43 hr under favourable conditions of PYG media. A, C, and E represent bacterial association with *A. castellanii*, while B, D, and F represent ratio of bacteria per amoeba. Results are representative of 3 independent experiments performed in triplicate. Asterisks indicate a significant difference, i.e., *P*<0.05, using a paired *t*-test, one-tail distribution.

trophozoites may have different pathogenic potential from invasive *E. coli* K1 from *A. castellanii* cysts, the association, invasion, and survival assays were performed as mentioned above.



**Fig. 6.** The pathogenic potential of *E. coli* K1 from *A. castellanii* cysts was not changed as compared with *E. coli* K1 from trophozoites. *E. coli* K1 from *A. castellanii* cysts and trophozoites was continuously assessed by association, invasion, and intracellular survival assays mentioned above. 'A' and 'B' represent association assays. 'B' and 'C' represent invasion assays. 'E' and 'F' represent intracellular survival association with *A. castellanii*, while B, D, and F represent ratio of bacteria per amoeba. Results are representative of 3 independent experiments performed in triplicate. Asterisks indicate a significant difference, i.e., P < 0.05, using a paired *t*-test, one-tail distribution.

The ability of *E. coli* K1 from *A. castellanii* cysts to interact with *A. castellanii* trophozoites exhibited little changes as compared with *E. coli* K1 from *A. castellanii* trophozoites to interact with *A. castellanii* trophozoites (Fig. 6). These results revealed that although the intracellularly survived *E. coli* K1 from *A. castellanii* cysts multiplied, the pathogenic potential of the *E. coli* K1 might have never changed.

#### DISCUSSION

The predatory role of Acanthamoeba in the control of bacterial populations and their ability to host bacteria and/or to act as a reservoir for bacterial pathogens suggests that these protists play important roles in the environment and contribute to human infections with bacteria. For the latter, a clear evidence is that the environmental bacterium, L. pneumophila parasitize Acanthamoeba as well as use it as a host or a vehicle to infect the human lung macrophages. The ability of L. pneumophila to survive intracellularly in Acanthamoeba has been suggested to be a key step in the evolution of this environmental bacterium to produce human infections. This concept is further strengthened with the finding that Acanthamoeba resembles human macrophages in many ways, particularly in their phagocytic activity and cell surface receptors [21] and that macrophages and Acanthamoeba exhibit parallel mechanisms in their interactions with L. pneumophila [22].

Here, to determine the differential interactions of E. coli K5 and E. coli K1 with A. castellanii, we analyzed association with, invasion into, and survival of these bacteria in A. castellanii. In previous studys, E. coli K1 of other capsular antigen compared with E. coli K12 were highly associated, invaded, and intracellularly survived in A. castellanii [1,8]. Interestingly, LPS chemical mutant of E. coli K1 exhibited less association, invasion, and survival in A. castellanii as compared with invasive E. coli K1. Beside FimH, both OmpA and LPS were crucial in E. coli K1 binding, association, and subsequent survival and replication within A. castellanii tested. This finding showed remarkable similarities with human macrophages. For example, recent studies have shown that E. coli K1 but not K12 are able to enter, survive, and replicate intracellularly in murine and human macrophages [23]. This property may be crucial for E. coli K1 survival in the bloodstream, a primary step in the development of meningitis.

As shown in this study, *E. coli* K5 exhibited significantly low association, invasion, and intracellular survival in *A. castellanii* trophozoites as compared with *E. coli* K1. It survived in *A. castellanii* cysts. Although *A. castellanii* cysts were transformed into trophozoites under PYG media, no *E. coli* K5 survived up to 0 hr which indicated the invasion assay. Thereafter, *E. coli* K5 survived in *A. castellanii* cysts and also recovered in transformed *A. castellanii* trophozoites under favourable conditions. Taken together, they implied that *A. castellanii* trophozoites more active in phagocytosis and movement than cysts would engulf *E. coli*  K5. In fact, *E. coli* K1 differs from *E. coli* K5 by the somatic antigen but not capsular antigen. *E. coli* K5 showed less association, invasion, and survival as compared with *E. coli* K1 with *A. castellanii* trophozoites and cysts. More importantly, the pathogenic potential of *E. coli* K1 was not changed as shown in Fig. 6.

Previous studies have shown that the attachment of L. pneumophila to the protozoan host, Hartmanella vermiformis is mediated by galactose/N-acetylgalactoseamine residues on the surface of bacteria that are recognized by a 170 kDa receptor lectin on the surface of protozoa [24,25]. In contrast, adherence of L. pneumophila to macrophages is dependent on heatlabile serum opsonins [26]. However, the mechanisms of subsequent survival and replication of L. pneumophila within protozoa and macrophages are remarkably similar [22,27-30]. The fact that Acanthamoeba resembles human macrophages in many ways, particularly in their phagocytic activity [9], and that macrophages and Acanthamoeba exhibit parallel mechanisms in their interactions with E. coli, suggest that Acanthamoeba may provide an alternative model to study E. coli pathogenesis as well as to understand bacterial immune evasion strategies. Future studies will establish this hypothesis and will precisely determine whether E. coli K5 survival and multiplication intracellularly of Acanthamoeba affects its virulence.

Overall, these studies suggest that *E. coli* K5 interactions with *A. castellanii* trophozoites are not significantly increased as compared with *E. coli* K1. Otherwise, *E. coli* K5 interactions were very similar with non-invasive *E. coli* K12. However, the interaction of *E. coli* K5 and *A. castellanii* cysts was obviously different from that of *E. coli* K12. Therefore, although the capsular serotypes were identical, these results suggested that the different somatic antigen in *E. coli* K5 would be a very important role in association, invasion, and survival of *E. coli* K5 in *A. castellanii* trophozoites and cysts.

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