

Contents lists available at ScienceDirect

Science of the Total Environment



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

Short Communication

Transfer of antibiotic resistance plasmids in pure and activated sludge cultures in the presence of environmentally representative micro-contaminant concentrations



Sungpyo Kim^{a,*}, Zuwhan Yun^a, Un-Hwan Ha^b, Seokho Lee^c, Hongkeun Park^d, Eilhann E. Kwon^e, Yunchul Cho^f, Sungwook Choung^g, Junsik Oh^a, Carl Angelo Medriano^a, Kartik Chandran^d

^a Department of Environmental Engineering, Korea University, Sejong 339-700, Republic of Korea

^b Department of Biotechnology and Bioinformatics, Korea University, Sejong 339-700, Republic of Korea

^c Department of Statistics, Hankuk University of Foreign Studies, 81 Oedae-ro, Mohyeon-myeon, Cheoin-gu, Yongin 449-791, Republic of Korea

^d Department of Earth and Environmental Engineering, Columbia University, 500 West 120th Street, New York, NY 10027, USA

^e Department of Environment and Energy, Sejong University, 98 Gunja-Dong, Gwangjin-Gu, Seoul 143-747, Republic of Korea

^f Department of Environmental Engineering, Daejeon University, 62 Daehak-Ro, Dong-Gu, Daejeon 300-716, Republic of Korea

^g Division of Advanced Nuclear Engineering, Pohang University of Science and Technology (POSTECH), San 31, Hyoja-dong, Nam-gu, Pohang 790-784, Republic of Korea

HIGHLIGHTS

• Increased plasmid transfer in ppb levels of tetracycline and sulfamethoxazole.

· Significant increase in plasmid transfer on activated sludge with the tetracycline.

• All pB10 plasmid received bacteria were enterics in presence of tetracycline.

ARTICLE INFO

Article history: Received 11 April 2013 Received in revised form 29 August 2013 Accepted 30 August 2013 Available online 25 September 2013

Editor: Eddy Y. Zeng

Keywords: Antibiotic resistance gene Conjugation Tetracycline Sulfamethoxazole Activated sludge

ABSTRACT

The presence of antibiotics in the natural environment has been a growing issue. This presence could also account for the influence that affects microorganisms in such a way that they develop resistance against these antibiotics. The aim of this study was to evaluate whether the antibiotic resistant gene (ARG) plasmid transfer can be facilitated by the impact of 1) environmentally representative micro-contaminant concentrations in ppb (part per billion) levels and 2) donor-recipient microbial complexity (pure vs. mixed). For this purpose, the multidrug resistant plasmid, pB10, and *Escherichia coli* DH5 α were used as a model plasmid and a model donor, respectively. Based on conjugation experiments with pure (*Pseudomonas aeruginosa* PAKexoT) and mixed (activated sludge) cultures as recipients, increased relative plasmid transfer frequencies were observed at ppb (µg/L) levels of tetracycline and sulfamethoxazole micro-contaminant exposure. When sludge, a more complex community, was used as a recipient, the increases of the plasmid transfer rate were always statistically significant but not always in *P. aeruginosa*. The low concentration (10 ppb) of tetracycline exposure led to the pB10 transfer to enteric bacteria, which are clinically important pathogens.

© 2013 Elsevier B.V. All rights reserved.

1. Introduction

Over the past decade, the presence of antimicrobial compounds and their residues in the environment have attracted great attention because of their intrinsic bioactivity and their continuous input to the aquatic environment (Kummerer, 2009a). The commonly found antimicrobial compound concentrations in the environment are around ppt (part per trillion) or ppb (part per billion) levels (Behera et al., 2011; Hirsch et al., 1999; Watkinson et al., 2009). Continuous input of antimicrobial agents to the environment could result in increased antibiotic resistance (Levy, 2002). This might be one of the major reasons that an increasing number of antibiotic resistant genes (ARGs) are found in the environment (Martinez, 2008). Several researchers showed that the abundance of these ARGs in the environment has been increasing because of human activities (Aminov and Mackie, 2007; Knapp et al., 2010; Zhu et al., 2013). Furthermore, the ARGs associated with pathogens have also increased (Brusselaers et al., 2011). Although a number of studies have documented positive relationships between antibiotic and the presence/persistence of antibiotic resistance in the environment (Kim et al., 2007; Merlin et al., 2011; Shakibaie et al., 2009), it is still unclear whether antibiotic

^{*} Corresponding author. Tel.: +82 44 860 1457; fax: +82 44 860 1588. *E-mail address:* ub1905ub@korea.ac.kr (S. Kim).

^{0048-9697/\$ -} see front matter © 2013 Elsevier B.V. All rights reserved. http://dx.doi.org/10.1016/j.scitotenv.2013.08.100

concentrations at ppb (parts per billion) levels can reach an effective threshold concentration towards proliferating antibiotic resistance in the environment (Barr et al., 1986; Ohlsen et al., 2003). In addition, the traditional selective pressure theory about antibiotics for the proliferation of antibiotic resistance in the microbial community might not be appropriate in environmental conditions since most of the ppb level of antibiotics cannot efficiently inhibit the antibiotic sensitive microorganisms. It is therefore possible that there is an alternate mechanism such as horizontal gene transfer (HGT) for the dissemination of antibiotic resistance traits in environments with such low levels of antimicrobial compounds.

HGT is an essential step for competitive bacterial survival in the environment and is also believed to be one of the major drivers for antibiotic gene transfer (Aminov, 2011; Shakibaie et al., 2009). ARG acquisition rate by HGT can possibly be affected by various environmental contaminants and conditions. For example, contaminants such as metals or antibiotics can damage the genomic DNA and result in the induction of the SOS response; SOS response could then promote the dissemination of HGT (Aminov, 2011; Hastings et al., 2004). In addition, the rate of HGT also depends on various environmental microorganism-associated factors such as plasmid donor and recipient species (De Gelder et al., 2005; Dionisio et al., 2002). In a previous study (Ohlsen et al., 2003), the transfer of conjugative gentamicin resistant (*aacA-aphD*) plasmids of Staphylococcus aureus (S. aureus) were investigated with different antibiotic concentrations. Although most antibiotics have no effect on the transfer of plasmid, approximately 3-fold increases were observed for gentamicin at 100 ppb in one mating pair [methicillin resistant S. aureus $(MA31) \times$ methicillin resistant S. aureus (MA20)].

Based on a combination of these previous observations, it is hypothesized that the increase of ARGs in environmental conditions could be related to the increased HGT rate induced by micro-contaminants among bacteria. It is also hypothesized that HGT can be affected by the environmental system's microbial complexity. These hypotheses are worthy of careful study because the micro-contaminants' effect on HGT among various bacteria under environmental matrices is scarce. Therefore, the aim of this study is to evaluate the impact of 1) threshold environmental micro-contaminant concentrations and 2) donorrecipient microbial complexity (pure vs. mixed) in the transfer of plasmid encoded antibiotic resistant genes.

2. Materials and methods

2.1. Bacterial strains

According to previous studies (Nikaido, 1998; Kelch and Lee, 1978) gram-negative bacteria have more significance in terms of medical research and are found to be more resistant to antibiotics than grampositive bacteria. Therefore, we assumed that Escherichia coli (E. coli) and its derived plasmid is a good model plasmid donor to the environment. Accordingly, E. coli DH5 α , containing the multidrug resistance plasmid pB10 was selected as the plasmid donor in this study. The complete 64,508 bp nucleotide sequence of the IncP-1 β plasmid pB10 was originally isolated from a wastewater treatment plant in Germany and mediates resistance against the antimicrobial agents amoxicillin, streptomycin, sulfamethoxazole, tetracycline and metallic mercury (Schluter et al., 2003). As a pure culture recipient, gentamicin resistant Pseudomonas aeruginosa PAKexoT was used in this study (Kaufman et al., 2000). In this study, P. aeruginosa was selected as a model environmental microorganism since it is found in various environmental conditions such as soil and water (Alonso et al., 1999). In evaluating the impact of donor-recipient microbial complexity in the transfer of pB10, activated sludge was used as a recipient in this study.

The donor and recipient cultures were grown separately in LB (lysogeny broth) medium, supplemented with appropriate antibiotics [donor: amoxicillin (50 µg/mL), tetracycline (20 µg/mL), streptomycin

(50 μ g/mL) and sulfamethoxazole (150 μ g/mL), recipient: gentamicin (50 μ g/mL)], and placed on a 150 rpm rotary shaker at 20 or 37 °C.

As a complex recipient, two-liter grab activated sludge samples were directly collected from an aeration basin in Cheongwon Wastewater Treatment Plant located at Osong in Chungbuk, Korea during June to July and decanted into sterile 1 L plastic bottles. Activated sludge samples were kept in an ice box, transported to the laboratory, and stored in a refrigerator. Samples were used as a recipient within 24 h after storing.

2.2. Plasmid transfer mating experiment

Each donor and recipient culture was grown in LB broth, with appropriate antibiotics in a 37 °C shaking incubator until they reached an optical density (O.D.) of 0.9 at 600 nm. Activated sludge was diluted with phosphate buffer to achieve an O.D. of 0.9. When the O.D. value was reached at 0.9, initial concentrations of potential recipients (P. aeruginosa PAKexoT or activated sludge) were enumerated by plate cultivation method for later transfer frequency calculation (T/R)transconjugant/(potential) recipient). After harvesting, each culture was centrifuged at 4,000*g for 15 minutes. The most of the supernatant was then discarded and the pellets containing the donor (E. coli DH5 α pB10) and recipient (*P. aeruginosa* PAKexoT or activated sludge) were re-suspended in the remaining supernatant and then mixed together, and inoculated on mating LB media plates containing one of the five stressors, antibiotics (amoxicillin, tetracycline, streptomycin, sulfamethoxazole) or metal (mercury), with concentrations from 0 to 1 µg/mL. After 16 h of incubation, the pellets (donor and recipient mixture) were re-suspended with 1 mL of LB broth and transferred to a tube and vortex-mixed for ten seconds. The donor-recipient mixtures were serially diluted and spread onto a transconjugant selecting LB media plates containing mixture of antibiotics depending on the recipient (*P. aeruginosa* PAKexoT or activated sludge). The recipients possessing pB10 plasmid were called transconjugants.

When *P. aeruginosa* PAKexoT was used as the recipient, the transconjugant selecting LB media plates contained tetracycline (2 mg/L) and gentamicin (10 mg/L). When the activated sludge was used as the recipient, the transconjugant selecting LB media plates contained a mixture of amoxicillin (1 ppm), tetracycline (1 ppm), streptomycin (1 ppm), sulfamethoxazole (1 ppm), and gentamicin (10 ppm). Donor and transconjugant are distinguished because of gentamicin (10 ppm) addition in selecting plate. The concentrations of four antibiotics (amoxicillin, tetracycline, streptomycin and sulfamethoxazole) were determined by preliminary study to confirm no growth of bacteria in recipients (activated sludge) (data are not shown).

After overnight incubation, the grown transconjugant colonies were counted. The colony averages were calculated using the triplicate plates. The pB10 transfer rate in this study was calculated using Eq. (1):

nB10 -	Transaction	/Recipient ratio on s	lective plate wi	th sterssor mating history	(1)
	Transaction	/Recipient ratio on s	lective plate wi	th sterssor mating history	(1)

2.3. Statistical analysis

The effects of the types and concentrations of micro-contaminants as well as the incubation times of the transconjugant colony count were tested by analysis of variance (ANOVA) using the linear fixed-effect model and linear mixed-effect model. Relative frequency (rate) of the transconjugant count over the total number of recipients is used as the response variable in the model because the large count makes the relative frequency behave as a normal distribution, while the values of relative frequency are quite small and almost nearly zero. The type and concentrations of micro-contaminants and the incubation times were considered as fixed factors in the models. Triplicated experiments were considered as random effects in the linear mixed-effect model to compensate for unwanted experimental effects which may not be

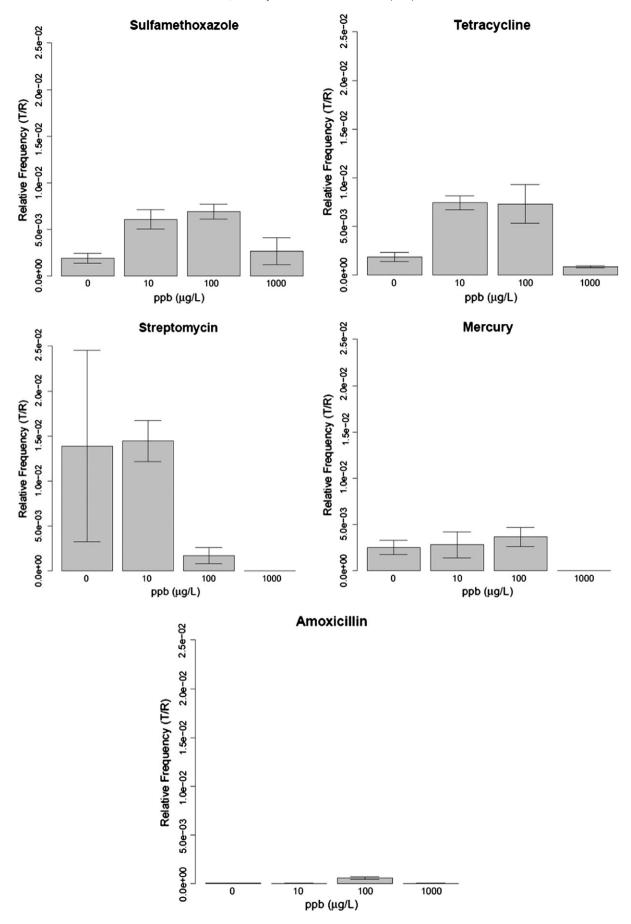


Fig. 1. Relative frequencies (*T/R*) of transconjugants (*P. aeruginosa* with pB10) as a function of various stressor concentrations [bar charts of averages are presented with standard error bars].

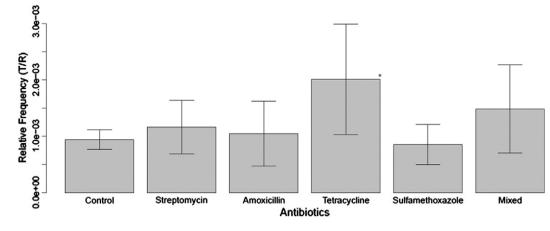


Fig. 2. Relative frequencies (*T*/*R*) of transconjugants (*P. aeruginosa* with pB10) with 2.5 ppb antibiotic concentration and antibiotic mixture [bar charts of averages are presented with standard error bars; statistical significance is denoted in the bar by * for *p*-value <0.05, ** for *p*-value <0.01, and *** *p*-value <0.001].

controllable in the experiments. Factor levels were compared using a post-hoc multiple-comparison procedure (Tukey's HSD). Statistical analyses were performed using R 2.15.1 with a significance level of 0.05.

2.4. DNA extraction and PCR

Plasmid DNA from the pB10-containing *E. coli* DH5 α was isolated using a Nucleobond Kit PC100 on AX 100 columns (Macherey-Nagel) according to the manufacturer's supplied protocol. The concentration and purity of the DNA were evaluated by ultraviolet absorbance spectrophotometry at 260/280 nm. The presence of pB10 plasmid in either extract or grown DNA colony was regularly confirmed by PCR using highly specific primers (F5'-CAATACCGAAGAAAGCATGCG-3', R5'-AGATATGGGTATAGAACAGCCGTCC-3') (Bonot and Merlin, 2010). After any given mating experiment, 5–10 chosen colonies from the selective LB plates and *E. coli* DH5 α pB10 culture (positive control) were compared via colony PCR to confirm the presence of a pB10 plasmid in transconjugant (data not shown).

2.5. 16S rRNA gene analysis in colonies and phylogenetic tree construction

To characterize the transconjugant species isolated in sludge mating experiments, 16S rRNA gene based phylogenetic analysis was conducted. Colony PCR was conducted using universal primers 27f (5'-AGAGTTTGATCMTGGCTCAG-3') and 1492r (5'-GGYTACCTTGTTACG ACTT-3'), as previously described (Teyssier et al., 2003). PCR products of about 1,400 bp were sequenced in an Applied Biosystems Automatic Sequencer (Genome Express). 16S rRNA clone sequences were clustered into OTUs (operational taxonomic units) at a cut-off of 97% by an open-source software, MOTHUR (Schloss et al., 2009). Phylogenetic analysis was conducted using the neighbour-joining method with bootstrap parameters of 1000 replications and evolutionary distances were computed using Jukes–Cantor method by using MEGA version 5 (Tamura et al., 2011).

3. Results and discussion

3.1. Impact of the stressor concentration on the extent of pB10 transfer

High numbers of transconjugant colonies, 10^6 CFU/mL to 10^9 CFU/mL, were found on the final transconjugant selecting LB plates under *P. aeruginosa* PAKexoT mating experiments (data are not shown). Transfer frequencies (*T*/*R*) were calculated by getting the proportion of transconjugants with respect to the potential recipient count as presented in Figs. 1 and 2.

The transfer frequencies of pB10 from *E. coli* DH5 α to *P. aeruginosa* PAKexoT showed varying responses to the five micro contaminants at four different contaminant levels. For statistical analysis, three separate experiments with duplicate samples in each experiment have been conducted. In this test, increased frequencies were observed in 10 ppb

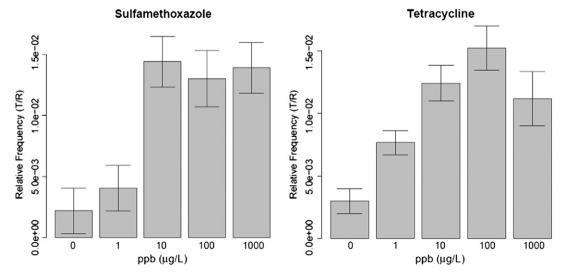


Fig. 3. Relative frequencies (*T*/*R*) of transconjugants (activated sludge with pB10) as a function of tetracycline or sulfamethoxazole concentrations [bar charts of averages are presented with standard error bars].

and 100 ppb of tetracycline and sulfamethoxazole (Tukey's HSD onesided test, *p*-value $< 10^{-4}$ for both 10 and 100 ppb levels) (Fig. 1). Mercury was not more considered after Fig. 1 experiment because mercury did not statistically significantly change for the pB10 transfer. In engineered environments such as wastewater treatment plants or natural environments, various antibiotics are usually found with mixed forms (Ding and He, 2010; Kummerer, 2009b). Therefore, we further tested the plasmid transfer frequencies with a lower microcontaminant concentration (2.5 ppb) of these mixed antibiotics to mimic real aquatic systems (Fig. 2). As shown in Fig. 2, tetracycline showed a significant increase in plasmid transfer frequencies (value) while other micro-contaminant doses did not show any statistically significant increases in the plasmid transfer frequencies. Based on this result, tetracycline and sulfamethoxazole are chosen for model antibiotics for following microbial complexity on extent of pB10 transfer.

Several previous studies also reported that plasmid transfer between bacteria can be enhanced by the presence of hundred ppb levels of some antibiotics. A previous study (Almasaudi et al., 1991) reported that the transfer of a pwG613 plasmid of *S. aureus* was enhanced 10-fold after the exposure to 500 ppb gentamicin. Another study (Ohlsen et al., 2003) also observed that the transfer frequencies of pSK41 type plasmid between *S. aureus* species can be increased by a threefold at 100 ppb gentamicin. However, the rates of plasmid transfer at even lower and more environmentally relevant antibiotic concentrations (10 ppb) have not been evaluated.

Several studies reported that the presence of low level of antibiotics, sub-inhibitory concentrations of antibiotics or metal, can induce the genetic expression of microorganisms (Babic et al., 2010; Wang and Crowley, 2005; Yim et al., 2007) including those coding for the SOS response (Deneve et al., 2009; Mesak et al., 2008), virulence properties (Hacker et al., 1993), and biofilm formation (Rachid et al., 2000).

Antibiotics can damage DNA and result in the promoting the repair system of bacteria such as SOS response (Miller et al., 2004; Phillips et al., 1987). Based on the previous study (Beaber and Waldor, 2004),

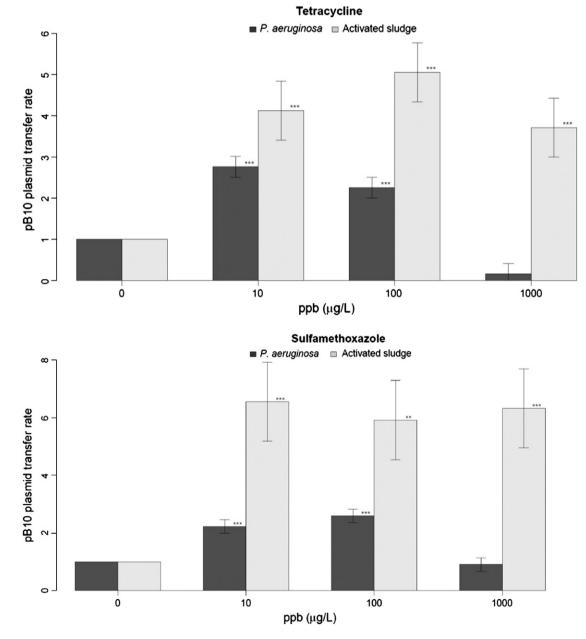


Fig. 4. Comparisons of pB10 plasmid transfer rates from *E. coli* to *P. aeruginosa* or activated sludge as a function of antibiotic concentrations [average rate in duplicate experiment; statistical significance is denoted by * for *p*-value <0.05, ** for *p*-value <0.01, and *** for *p*-value <0.001].

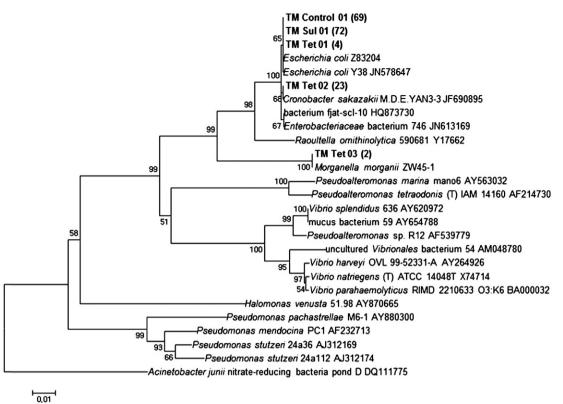


Fig. 5. Phylogenetic tree based on partial 16S rRNA gene to identify trans-conjugated microorganism species in each different antibiotic concentration in solid media [control (0 ppm; Control), tetracycline (10 ppb; Tet), sulfamethoxazole (10 ppb; Sul)]. The evolutionary distances were computed using the Jukes–Cantor method. Parentheses represent the number of clones analyzed in each experiment.

the transfer frequency of SXT, one of integrating conjugative elements (ICESs), between cell to cell can be enhanced by antibiotic-induced SOS response.

the reduced transconjugant concentration at 1000 ppb could be explained by the reduced recipient concentrations.

d 3.2. Impact of microbial complexity on extent of pB10 transfer

When activated sludge, representing more complex communities, was used as a recipient, the pB10 transfer from *E. coli* to activated sludge always increased when the microbial communities were exposed to a ppb level of tetracycline or sulfamethoxazole (ANOVA test, *p*-value $<10^{-4}$) (Fig. 3). The plasmid transfer rates of both *P. aeruginosa* and activated sludge were calculated using Eq. (1) and compared. For the activated sludge, the increases in the plasmid transfer rate were always statistically significant but not always for *P. aeruginosa* (Fig. 4).

As summarized in Fig. 5, with the absence of antibiotics in LB plates, all 69 of the pB10 transferred bacteria were phylogenetically related to *E. coli*. In the case of tetracycline amended plates, 4 out of 29 transconjugants were related to *E. coli*, whereas 2 transconjugants were closely related to *Morganella morganii*. However, most of the transconjugants (23/29) from tetracycline treated plates were related to *Cronobacter sakazakii*-like bacteria. In the case of sulfamethoxazole containing plates, *E. coli*-related bacteria were dominant which were similarly observed in the control plates.

In our findings, the standard deviation showed that plasmid transfer rate was more variable when a complex microbial community (activated sludge) was used as the recipient compared to a pure culture (*P. aeruginosa*), regardless of antibiotic concentrations (Fig. 4). This observed variability could be explained by the presence of the higher diversity of secondary plasmid donors in activated sludge. During *E. coli* and *P. aeruginosa* conjugation experiments, pB10 donors can be either *E. coli* (pB10) or transconjugant *P. aeruginosa* (pB10). However, when activated sludge culture was used as recipients, various types of microorganisms can become secondary pB10 donors after they receive the pB10 through primary conjugation. A previous study showed that

Cell to cell contact between donor and recipients might be secured by biofilm formation which is induced by the ppb concentrations of antibiotics that result in the enhanced HGT. More secretions of small proteins or peptides in the presence of certain antibiotics by the donor or recipients or both help the adhesion between bacteria or between bacteria and external surfaces (Kaplan et al., 2011) and these could result in the biofilm formation (Ong et al., 2009). Under these circumstances, plasmid transfer rate can be increased between two cells since bacterial cells in a biofilm stays in close contact since it helps the cell-to-cell gene transfer (Angles et al., 1993; Krol et al., 2011; Martiny et al., 2003).

In our experiment (Fig. 1), we also observed that relative frequencies of a transconjugant colony at a high micro-contaminant concentration (1,000 ppb) were always lower than those at 10 ppb micro-contaminant concentration (Tukey's HSD one-sided test, p-value 0.0029). A recent study (Yim et al., 2007; Linares et al., 2006) suggested that the effect of antibiotics on global bacterial transcription is concentration-dependent. Under the sub-inhibitory concentration, antibiotic can act as a signal molecule to the microorganisms or microbial community, and at higher concentrations; however, bacterial responses shift to the more stress-related ones. The loss of viability emerged at the highest antibiotic concentrations. Therefore, our observation could be explained by dual antibiotic activity: antibiotics help gene transfer at low concentrations (10–100 ppb) but inhibit or kill some of the donors and recipient cultures, as the concentration increases. To confirm this idea, both the donor (*E. coli* DH5 α) and the recipient (P. aeruginosa PAKexoT) were separately incubated with different concentrations of tetracycline or sulfamethoxazole to account for the survival rate. The experiment showed that about 10 fold of P. aeruginosa PAKexoT could be reduced under the 1,000 ppb concentration of tetracycline or sulfamethoxazole (data not shown). Therefore,

pB10 host diversity strongly influences the pB10 transfer rate to the complex microbial community (De Gelder et al., 2005). According to the De Gelder et al. (2005)), the *tra* gene regulation of the host, restriction-modification system for foreign DNA between donor and recipient, and non-random spatial distribution between cells in activated sludge, could result in the difference of pB10 transfer rate under host diversity. Another study also mentioned that the presence of efficient donors in heterogeneous bacterial populations can accelerate plasmid transfer and can spread by several orders of magnitude (Dionisio et al., 2002).

It can be noted that pB10 plasmids were transferred to the bacteria belonging to a potential clinically important family, the *Enterobacteriaceae* family. The high plasmid transfer frequency to the enteric bacteria implied that these microorganisms could become gene reservoirs and could play a key role in the recent rise of antibiotic resistance in an aquatic environment (Martinez, 2008). In this study, large amounts of the transconjugant colonies from tetracycline treatment were related to *C. sakazakii*-like bacteria (previously termed *Enterobacter sakazakii*) which has been identified as an opportunistic pathogen and has been linked with life-threatening infections among infants (Healy et al., 2010).

4. Conclusions

A number of studies have implicated the potential positive relationships between antibiotics and antibiotic resistant bacteria in the environment. However, little information exists regarding the threshold concentrations of antibiotics for enhancing ARGs in the environment through mechanisms such as horizontal gene transfer. Based on this study's findings, statistical increase in pB10 plasmid transfer via conjugation process was observed even in the presence of 10 ppb of antibiotics, especially tetracycline and sulfamethoxazole. It was also observed that the pB10 plasmid transfer rate to normal flora significantly can statistically increase with 10 to 100 ppb of tetracycline or sulfamethoxazole exposure rather than to P. aeruginosa. Although the exact mechanism for the increased pB10 transfer rate still needs further investigation, it is important to note that these results indicated that the spread of antibiotic resistance can be accelerated to the environmental microorganisms in the ppb level of antibiotics. One of the interesting findings of this study is that the low concentration (10 ppb) of tetracycline exposure led to the pB10 transfer to enteric bacteria, which are clinically important pathogens. Further study is needed to link antibiotic residue concentrations and the selection of other pathogens in mixed microbial communities such as activated sludge.

Acknowledgements

This research was equally supported by the Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Education, Science and Technology (No. 2012-0003505) and by Korea Ministry of Environment as "Global Top Project" (Project no.: GT-11-B-01-005-1).

References

- Almasaudi SB, Day MJ, Russell AD. Effect of some antibiotics and biocides on plasmid transfer in *Staphylococcus aureus*. J Appl Bacteriol 1991;71:239–43.
- Alonso A, Rojo F, Martinez JL. Environmental and clinical isolates of *Pseudomonas* aeruginosa show pathogenic and biodegradative properties irrespective of their origin. Environ Microbiol 1999;1:421–30.
- Aminov RI. Horizontal gene exchange in environmental microbiota. FMICB 2011;2:401-4.
- Aminov RI, Mackie RI. Evolution and ecology of antibiotic resistance genes. FEMS Microbiol Lett 2007;271(2):147–61.
 Angles ML, Marshall KC, Goodman AE. Plasmid transfer between marine-bacteria in the
- Angles ML, Marshali KC, Goodman AE, Plashid transfer between manne-bacteria in the aqueous phase and biofilms in reactor microcosms. Appl Environ Microbiol 1993;59: 843–50.
- Babic F, Venturi V, Maravic-Vlahovicek G. Tobramycin at subinhibitory concentration inhibits the RhII/R quorum sensing system in a *Pseudomonas aeruginosa* environment isolate. BMC Infect Dis 2010;10:148.

- Barr V, Barr K, Millar MR, Lacey RW. Beta-lactam antibiotics increase the frequency of plasmid transfer in *Staphylococcus aureus*. J Antimicrob Chemother 1986;17:409–13.
- Behera SK, Kim HW, Oh J-E, Park H-S. Occurrence and removal of antibiotics, hormones and several other pharmaceuticals in wastewater treatment plants of the largest industrial city of Korea. Sci Total Environ 2011;409:4351–60.
- Beaber JW, Waldor MK. Identification of operators and promoters that control SXT conjugative transfer. J Bacteriol 2004;186(17):5945–9.
- Bonot S, Merlin C. Monitoring the dissemination of the broad-host-range plasmid pB10 in sediment microcosms by quantitative PCR. Appl Environ Microbiol 2010;76:378–82.
- Brusselaers N, Vogelaers D, Blot S. The rising problem of antimicrobial resistance in the intensive care unit. Ann Intensive Care 2011;1:47.
- De Gelder L, Vandecasteele FP, Brown CJ, Forney LJ, Top EM. Plasmid donor affects host range of promiscuous IncP-1beta plasmid pB10 in an activated-sludge microbial community. Appl Environ Microbiol 2005;71:5309–17.
- Deneve C, Bouttier S, Dupuy B, Barbut F, Collignon A, Janoir C. Effects of subinhibitory concentrations of antibiotics on colonization factor expression by moxifloxacinsusceptible and moxifloxacin-resistant *Clostridium* difficile strains. Antimicrob Agents Chemother 2009;53:5155–62.
- Ding C, He J. Effect of antibiotics in the environment on microbial populations. Appl Microbiol Biotechnol 2010;87:925–41.
- Dionisio F, Matic I, Radman M, Rodrigues OR, Taddei F. Plasmids spread very fast in heterogeneous bacterial communities. Genetics 2002;162:1525–32.
- Hacker J, Ott M, Hof H. Effects of low, subinhibitory concentrations of antibiotics on expression of a virulence gene cluster of pathogenic *Escherichia coli* by using a wild-type gene fusion. Int J Antimicrob Agents 1993;2:263–70.
- Hastings PJ, Rosenberg SM, Slack A. Antibiotic-induced lateral transfer of antibiotic resistance. Trends Microbiol 2004;12:401–4.
- Healy B, Cooney S, O'Brien S, Iversen C, Whyte P, Nally J, et al. Cronobacter (Enterobacter sakazakii): an opportunistic foodborne pathogen. Foodborne Pathog Dis 2010;7:339–50.
- Hirsch R, Ternes T, Haberer K, Kratz K-L. Occurrence of antibiotics in the aquatic environment. Sci Total Environ 1999;225:109–18.
- Kaplan JB, Jabbouri S, Sadovskaya I. Extracellular DNA-dependent biofilm formation by *Staphylococcus epidermidis* RP62A in response to subliminal inhibitory concentrations of antibiotics. Res Microbiol 2011;162:535–41.
- Kaufman MR, Jia J, Zeng L, Ha U, Chow M, Jin S. Pseudomonas aeruginosa mediated apoptosis requires the ADP-ribosylating activity of exoS. Microbiology 2000;146(Pt 10):2531-41.
- Kelch WJ, Lee JS. Antibiotic-resistance patterns of gram-negative bacteria isolated from environmental sources. Appl Environ Microbiol 1978;36(3):450–6.
- Kim S, Jensen JN, Aga DS, Weber AS. Tetracycline as a selector for resistant bacteria in activated sludge. Chemosphere 2007;66:1643–51.
- Knapp CW, Dolfing J, Ehlert PAI, Graham DW. Evidence of increasing antibiotic resistance gene abundances in archived soils since 1940. Environ Sci Technol 2010;44:580–7.
- Krol JE, Nguyen HD, Rogers LM, Beyenal H, Krone SM, Top EM. Increased transfer of a multidrug resistance plasmid in *Escherichia coli* biofilms at the air–liquid interface. Appl Environ Microbiol 2011;77:5079–88.
- Kummerer K. Antibiotics in the aquatic environment a review part II. Chemosphere 2009a;75:435–41.
- Kummerer K. Antibiotics in the aquatic environment a review part I. Chemosphere 2009b;75:417–34.
- Levy SB. The 2000 Garrod lecture. Factors impacting on the problem of antibiotic resistance. J Antimicrob Chemother 2002;49:25–30.
- Linares JF, Gustafsson I, Baquero F, Martinez JL. Antibiotics as intermicrobial signaling agents instead of weapons. Proc Natl Acad Sci U S A 2006;103:19484–9.
- Martinez JL Antibiotics and antibiotic resistance genes in natural environments. Science 2008;321:365–7.
- Martiny AC, Jorgensen TM, Albrechtsen HJ, Arvin E, Molin S. Long-term succession of structure and diversity of a biofilm formed in a model drinking water distribution system. Appl Environ Microbiol 2003;69:6899–907.
- Merlin C, Bonot S, Courtois S, Block JC. Persistence and dissemination of the multipleantibiotic-resistance plasmid pB10 in the microbial communities of wastewater sludge microcosms. Water Res 2011;45:2897–905.
- Mesak LR, Miao V, Davies J. Effects of subinhibitory concentrations of antibiotics on SOS and DNA repair gene expression in *Staphylococcus aureus*. Antimicrob Agents Chemother 2008;52:3394–7.
- Miller C, Thomsen LE, Gaggero C, Mosseri R, Ingmer H, Cohen SN. SOS response induction by beta-lactams and bacterial defense against antibiotic lethality. Science 2004;305: 1629–31.
- Nikaido H. Antibiotic resistance caused by gram-negative multidrug efflux pumps. Clin Infect Dis 1998;27:S32–41.
- Phillips I, Culebras E, Moreno F, Baquero F. Induction of the SOS response by new 4-quinolones. | Antimicrob Chemother 1987;20:631–8.
- Ohlsen K, Ternes T, Werner G, Wallner U, Löffler D, Ziebuhr W, et al. Impacts of antibiotics on conjugational resistance gene transfer in *Staphylococcus aureus* in sewage. Environ Microbiol 2003;5:711–6.
- Ong CL, Beatson SA, McEwan AG, Schembri MA. Conjugative plasmid transfer and adhesion dynamics in an *Escherichia coli* biofilm. Appl Environ Microbiol 2009;75:6783–91.
- Rachid S, Ohlsen K, Witte W, Hacker J, Ziebuhr W. Effect of subinhibitory antibiotic concentrations on polysaccharide intercellular adhesin expression in biofilm-forming *Staphylococcus epidermidis*. Antimicrob Agents Chemother 2000;44:3357–63.
- Schloss PD, Westcott SL, Ryabin T, Hall JR, Hartmann M, Hollister EB, et al. Introducing mothur: open-source, platform-independent, community-supported software for describing and comparing microbial communities. Appl Environ Microbiol 2009;75: 7537–41.

- Schluter A, Heur H, Szczepanowski R, Forney LJ, Thomas CM, Puhler A, et al. The 64 508 bp IncP-1beta antibiotic multiresistance plasmid pB10 isolated from wastewater treatment plant provides evidence for recombination between members of different branches of the IncP-1beta group. Microbiology 2003;149:3139–53.Shakibaie MR, Jalilzadeh KA, Yamakanamardi SM. Horizontal transfer of antibiotic resis-
- Shakibaie MR, Jalilzadeh KA, Yamakanamardi SM. Horizontal transfer of antibiotic resistance genes among gram negative bacteria in sewage and lake water and influence of some physico-chemical parameters of water on conjugation process. J Environ Biol 2009;30:45–9.
- Tamura K, Peterson D, Peterson N, Stechner G, Nei M, Kumar S. MEGA 5: molecular evolution genetics analysis using maximum likelihood, evolutionary distance, and maximum parsimony methods. Mol Biol Evol 2011;28:1–18.
- Teyssier C, Marchandin H, Simeon De Buochberg M, Ramuz M, Jumas-Bilak E. A typical 16S rRna gene copies in Ochrobactrum intermedium strains reveal a large genomic rearrangement by recombination between rrn copies. J Bacteriol 2003;185:2901–9.
- Wang A, Crowley DE. Global gene expression responses to cadmium toxicity in Escherichia coli. | Bacteriol 2005;187:3259–66.
- Watkinson AJ, Murby EJ, Kolpin DW, Costanzo SD. The occurrence of antibiotics in an urban watershed: from wastewater to drinking water. Sci Total Environ 2009;407:2711–23.
 Yim G, Wang HH, Davies J. Antibiotics as signalling molecules. Philos Trans R Soc Lond B
- Biol Sci 2007;362:1195–200. Zhu YG, Johnson TA, Su JQ, Qiao M, Guo GX, Stedtfeld RD, et al. Diverse and abundant resistance genes in Chinese swine farms. Proc Natl Acad Sci U S A 2013;110:3435–40.