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Heavy Metal Concentration and Identification of Microorganisms in Soil under Roadside Trees of Daejeon City, Korea

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Due to the rapid urbanization and scarcity of land, most of metropolitan city of Korea are subjected to many potential pollution sources, including vehicle exhaust and industrial emissions. Soil samples were collected under roadside trees, *Platanus occidentalis* and *Ginkgo biloba*, to study the current conditions of heavy metal contamination and microorganism in Daejeon city, one of the metropolitan city in Korea. The total concentrations of heavy metals in the samples were determined. The results indicate that soils of roadside trees in Daejeon city have elevated concentrations of Cd, Cu, Zn, and Pb. Especially, high metal concentrations in soil were found in old urban commercial district and heavy traffic areas, indicating that the major contamination sources in these soils are traffic emissions. Rhizosphere bacteria in soil of roadside trees in Daejeon area were characterized in that strain richness was lower than other soil samples. Bacterial community members were characterized by DNA sequence analysis of 16S rDNA gene fragments following colony's DNA isolation emerged from nutrient agar medium using dilution plate method after soil sampling, polymerase chain reaction amplification and sequencing. Phylogenetic analyses revealed that 16S rDNA sequences were classified β -Proteobacteria, Actinobacteria, Gram-positive bacteria group. Among three study sites, Daejeon station, Daedeok science town, and Daehwa industrial complex, Daehwa industrial complex's soil microbe's richness was lowest. The 16S rDNA sequence and fragment database constitutes a reference to which DNA profiles of roadside tree soil's bacterial communities can be compared.

INTRODUCTION

Heavy metals may come from many different sources in urbanized areas, including vehicle emissions, industrial discharge and other activities (Harrison *et al.*, 1981; Gibson and Farmer, 1986; Thornton, 1991; Li *et al.*, 2001). However, the extensive use of automobile is one of the most important sources of heavy metal contamination in the roadside environment (Garcia and Millan, 1994). These elements can give critical effect to the ecosystems and human health if they are present at high concentration levels.

In Korea, there has been an increase in scientific and public concern that heavy traffic are likely to result in a critical risks for the ecosystem and human health. Most studies (You and Son, 2003; Park, 1992) dealt with problems of heavy metal content in top soils and air, and mainly in Seoul, the capital city of Korea. However, in order to give a better picture of the heavy metal contamination, heavy metal content in soil should be also determined in other metropolitan city.

With the population of over 1.5 million and area of 539.59 km², Daejeon is one of the most densely populated areas in Korea. The rapid urbanization and increasing vehicle have put great pressure on the local environment. Due to the scarcity of land, most of roads

in Daejeon city are narrow and busy, and close to buildings, where the roadside trees are subject to potential pollution from various sources.

Although soils carry out important functions in ecological metabolism, soil ecology increasingly is degrading to the reasons such as population expansion and industrial activities according to the area. Plant interact with bacteria and fungi, some rhizosphere bacteria adhere tightly to soil fungal hyphae, whereas others are directly associated with the root surfaces (Bianciotto *et al.*, 2001). The study of microbial diversity in natural environments has advanced significantly after the introduction of SSU rDNA-based molecular techniques (Amann *et al.*, 1995). Since the knowledge on the diversity of road tree's soil microbial aggregates in city is lagging, an attempt to isolate and identify rhizosphere soil samples through molecular methods was made.

In this work the heavy metal concentrations and microorganism small-subunit (SSU) 16S rDNA gene sequence of the soil adjacent to roads of varying traffic flow were determined. The effect of the traffic flow to heavy metal content and identification of microorganism in soil under roadside trees, *Platanus occidentalis* and *Ginkgo biloba*, was studied.

MATERIALS AND METHODS

This study was proceeded in 2004 at the Daejeon city, southern part of Seoul, Korea. The total number of roadside trees in Daejeon city was 107,980 and 32.7% of the total trees was *Ginkgo biloba* L. and 15.9% was *Platanus occidentalis* L. (EPA of Daejeon city, 2004).

Four study sites were selected for this study. The

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Table 1. Characteristics of the study species.

Sampling Sites	Species	Mean D.B.H* (cm)	Mean Height (m)	Ages (yr)
Chungnam National University	<i>Platanus occidentalis</i>	38	10	35
	<i>Ginkgo biloba</i>	24	9	25
Daejeon Station	<i>Platanus occidentalis</i>	42	10	55
	<i>Ginkgo biloba</i>	34	8	50
Daedeok Science Town	<i>Platanus occidentalis</i>	22	10	25
	<i>Ginkgo biloba</i>	18	8	25
Daehwa Industrial Complex	<i>Platanus occidentalis</i>	33	11	35
	<i>Ginkgo biloba</i>	16	7	20

* D.B.H: Diameter of Breast Height.

most traffic and commercial areas are located in Daejeon station area. Many of commercial and chemical factory were in the Daehwa industrial complex. The new town and usually consists of residential and research areas was in Daedeok science town. The less traffic area was located in Chungnam National university site. The characteristics of the study species was in Table 1.

In study sites, soil samples under roadside trees, *Ginkgo biloba* and *Platanus occidentalis*, were collected from 0–10 cm soil depth by using an Oakfield soil sampler during September, 2004. Soils were air-dried and sieved to pass through a 2 mm sieve. All soil samples were analyzed by methods detailed by Bickelhaupt and White (1982). Soil samples were digested in a concentrated HCl and the heavy metal concentration was determined by ICP–AES. Tests of significance were set at the 0.05 level unless otherwise stated. The SAS computer software system was used in this study (SAS Institute, 1985).

The soil samples collected and transferred were stored in an ice pack at 4 °C in the laboratory. One milliliter of the appropriate (10–5~10–7) dilutions of the soil samples was plated on nutrient agar's medium for the isolation of cultural microbes. Petri plates were then incubated, at 28 °C for 48 hours for bacteria. After the required incubation period, DNA was extracted and purified using the QIAamp DNA stool Mini Kit (QIAGEN Corp.) from colonies. The DNA of colonies emerged in

the plates was identified by the molecular methods using SSU rDNA polymerase chain reaction (PCR)–based fingerprinting.

Our study focuses on the cultured bacterial community in general thus we are using primers 5'–GAGTTTGATCCTGGCTCAG–3' (9F: positions 9 to 27nt, E. coli 16S rDNA numbering) and 5'–ACGGCTACCTTGTTACGACTT–3' (1542R: positions 1542 to 1525nt, E. coli 16rDNA numbering). The PCR was performed using a MyGenie 32 Thermal Block (Bioneer Corp. Korea) and the following cycles: initial denaturation at 94 °C for 5 min followed by 34 cycles of denaturation at 95 °C for 1 min, annealing at 60 °C for 1 min, and extension at 72 °C for 1 min 30s, and final extension at 72 °C for 7 min. PCR products were examined on a 1% agarose gel and the DNA was diluted for sequencing and assessing.

Amplification of DNA with the eubacterium primer 9F and 1542R produced almost the entire SSU gene fragments. PCR products were purified with QIAquick Gel extraction Kit (Qiagen, Corp.). DNA concentrations were quantified using a fluorometer. PCR products were sequenced by automatic sequencer ABIPRISM™ (Perkin–Elmer, USA at Eugeneteck Co., Korea). After making the sequencing, GENDOC program, Tree View (3.2) and CLUSTAL X (v1.81; Thompson *et al.*, 1994) were used for multiple alignment and neighbor-joining phylogeny (Saitou and Nei, 1987) using partial 16S rDNA gene sequences of species selected from gene bank of NCBI (<http://www.nlm.nih.gov>, Gehrig *et al.*, 1996).

RESULTS AND DISCUSSION

The cadmium, copper, lead and zinc concentrations in soil of *Ginkgo biloba* ranged from 1.12 to 1.92, 21.5 to 28.2, 17.2 to 28.2, and 50.0 to 429.5 mg kg⁻¹ at 0–10 cm soil depth in the four study sites, respectively (Table 2). Li *et al.* (2001) reported that Cd, Cu, Pb, and Zn concentration in urban soil of Hong Kong ranged from 0.02 to 5.89, 5.12 to 190, 5.27 to 404, and 38.7 to 43.5 mg kg⁻¹.

There was significant difference in the cadmium, copper, lead, and zinc concentrations among the four study sites of *Ginkgo biloba*. The percent cadmium, copper, lead, and zinc in soil was higher in heavy traffic area (Daejeon station) than in the light traffic area (Chungnam national university). The result could be from rubbing and wear of metal and car tire when they

Table 2. Soil cadmium, copper, lead, and zinc concentrations (mg kg⁻¹) at 0~10 cm soil depth under roadside trees, *Ginkgo biloba*, in Daejeon metropolitan city.

Heavy metal	C.N.U*	D.S	D.I.C	D.S.T
Cadmium	1.29±0.10b*	1.69±0.16a	1.55±0.28ab	1.65±0.27ab
Copper	27.14±4.02c	120.10±39.11a	76.48±24.35b	40.40±6.04bc
Lead	19.12±1.36b	25.50±2.65a	24.62±2.65a	24.96±2.86a
Zinc	64.20±6.17b	276.80±120.1a	109.14±26.83b	84.10±36.56b

* C.N.U: Chungnam National University, D.S: Daejeon Station, D.I.C: Daehwa Industrial Complex, D.S.T: Daedeok Science Town.

* Different letters indicate statistical differences among the treatments at the 5% level.

travel (Li *et al.*, 2001; You and Son, 2003; Park, 1992).

The cadmium, copper, lead and zinc concentrations in soil of *Platanus occidentalis* ranged from 0.92 to 1.69, 16.0 to 288.0, 16.9 to 32.4, and 100.0 to 415.0 mg kg⁻¹ in the four study sites, respectively (Table 3). There was significant difference in the copper and zinc concentrations among the four study sites. The percent copper and zinc in soil were higher in heavy traffic area (Daejeon station and Daehwa industrial complex) than in the light traffic area (Chungnam national university). Li *et al.* (2001), You and Son (2003), and Park (1992) reported that high metal concentrations was found in heavy traffic areas in Hong Kong and Seoul, Korea. However, difference in the case of *Ginkgo biloba*, there was no significant difference in the cadmium and lead concentrations among the study sites because of probably soil variation.

Roadside soil samples were collected to examine microbe in the rhizosphere from study sites, Daejeon-station, Daehwa industrial complex, and Daedeok Science Town. We used nutrient agar medium to culture eubacterium after dilution (10⁻⁵~10⁻⁷). And then, the cell emerged in the plates cultured during 3 days used to identify species using SSU 16S rDNA partial sequences. Colony number appeared in the plates was low level compared to general soil such as, agricultural and forest soils, Daejeon station appeared 15, Daehwa industrial complex 2, Daedeok science town 5. DNAs isolated from colony were subjected to PCR with bacterium-specific 16S rDNA primers (9F, 1542R). The size of the DNAs obtained with primers 9F-1542R

were consistent with those an amplified portion of the 16S rDNA.

Amplified DNAs were purified on agarose gel and sequenced automatic sequencer ABIPRISM™ (Perkin-Elmer, USA at Eugenetech Co., Korea). Two to 8 16S rDNA clones of 3 roadside tree soil samples were partially sequenced at the 5' end of the 16S rDNA gene. From NCBI searching results, the sequences of DNA fragments obtained from colonies were synthesized to 1500bps and revealed to be similar to those of some eubacteria within comparison of 16S rDNA sequences (Table 4).

Phylogenetic analyses revealed that roadside soil 16S rDNA clones were classified as β -Proteobacteria, Actinobacteria, Gram-positive bacteria (George, 2001). DS-1, DIC-1, DS-3 were β -Proteobacteria lineage, DS-3, DS-4, DS-5 were Actinobacteria and DS-6, DS-7, DIC-2 were Gram-positive bacteria lineage (Figure 1).

Similarity value was 96~99% except that DS-8 was 80%. Roadside soil eubacteria of Daejeon station and Daedeok science town were general bacteria isolated from soils as streptomycetes strains of Actinobacteria lineage, these organisms are strict aerobes (Tortora *et al.*, 2002). Daehwa industrial complex represented only two colonies compare with other NA plates; these organisms are bacillus megaterium and agricultural soil bacterium. Daedeok science town's soil was identified as *Streptomyces* sp. The strains abundance varied between the three study sites. The Daejeon station area showed the highest microbial colony, followed by the Daedeok science town and Daehwa industrial complex

Table 3. Soil cadmium, copper, lead, and zinc concentrations (mg kg⁻¹) at 0~10 cm soil depth under roadside trees, *Platanus occidentalis*, in Daejeon metropolitan city.

Soil depth (cm)	C.N.U*	D.S	D.I.C	D.S.T
Cadmium	1.14±0.27a*	1.18±0.15a	1.43±0.18a	1.32±0.18a
Copper	22.40±5.49b*	116.20±93.37a	48.10±10.92ab	27.10±5.79b
Lead	22.02±4.55a*	24.36±4.35a	25.98±4.29a	21.98±1.14a
Zinc	103.20±4.31b*	231.20±114.71a	154.30±14.13ab	124.20±12.01b

* C.N.U: Chungnam National University, D.S: Daejeon Station, D.I.C: Daehwa Industrial Complex, D.S.T: Daedeok Science Town.

* Different letters indicate statistical differences among the treatments at the 5% level.

Table 4. Identification of the 16S rDNA sequencing from roadside tree soil samples.

No.	16S rDNA identification	Accession number	Sequence similarity (%)
DS-1	<i>Agricultural soil bacterium</i>	AJ252575	697/703 (99%)
DS-2	<i>Streptomyces lincolnensis</i>	X79854	770/797 (96%)
DS-3	<i>Agricultural soil bacterium</i>	AJ252575	680/693 (98%)
DS-4	<i>Streptomyces capoamus</i>	AB045877	775/793 (97%)
DS-5	<i>Rhodococcus wratislaviensis</i>	AY785739	681/757 (89%)
DS-6	<i>Bacillus benzoevorans</i>	AY167808	714/718 (99%)
DS-7	<i>Uncultured Bacillus sp.</i>	AY876909	362/389 (93%)
DS-8	<i>Streptomyces sp.</i>	AF131520	545/675 (80%)
DIC-1	<i>Agricultural soil bacterium</i>	AJ252575	701/703 (99%)
DIC-2	<i>Bacillus megaterium</i>	AY030338	751/768 (97%)
DS-1	<i>Streptomyces sp.</i>	AF131563	809/830 (97%)

* DS: Daejeon station, DIC: Daehwa industrial complex, DS: Daedeok science town.

