

B047

Metagenomic Description of Lignocellulose-degrading Potentials in Soil near the Antarctic King Sejong Station Using Single Molecules Real-Time Sequencing

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B048

Comparative Analysis of Skin Microbiome between Young and Old Women

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B049

Distinct Soil Bacterial Distribution in Four Different Farming Management Soils

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B050

Microbial Community Structure and Composition in Rhizosphere and Endosphere of Tomato Plants Cultivated in Greenhouses

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B051

Characterization of Spatial Distribution of the Bacterial Community in the South Sea of Korea

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B052

Bacterial Community Structure of Surface Snow in Antarctica

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B053

Isolation of a BTEX-degrading Bacterium, *Rhodococcus* sp. B23, from a Freshwater and Optimization of Biodegradation Conditions

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B054

Metagenomic Analysis of Relationships between Carbon and Nitrogen Metabolisms during Tannery Wastewater Treatment Undergoing a Microbial Bioaugmentation

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B055

To Study the Effect of Change in NaCl Concentrations on Virulence of *Streptococcus parauberis*

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B056

Exploring the Influence of Environmental Factors on Rock-associated Bacterial Communities in Northern Victoria Land, Antarctica

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B057

Control of Soil Community Structure with Oxygen and Organic Substrates

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B058 9th ASME

Bacterial Community in Enrichment Culture of Cut Grass Residue Where Simultaneously Occurred both Cellulose Degradation and Nitrogen Fixation

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