A report on 20 unrecorded bacterial species of Korea isolated from soil in 2021

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As a subset study to discover indigenous prokaryotic species in Korea, we isolated 20 bacterial strains and assigned them to the phyla *Actinobacteria*, *Bacteroidota*, *Firmicutes*, and *Proteobacteria*. From the high 16S rRNA gene sequence similarity (\geq 98.7%) and formation of a robust phylogenetic clades, we determined that each strain belonged to independent, predefined bacterial species. There are no official reports of these 20 species in Korea; therefore, 7 strains of the *Actinobacteria*, 2 strain of the *Bacteroidota*, 3 strains of the *Firmicutes*, and 8 strains of the *Firmicutes* are described in Korea for the first time. Gram reaction, colony and cell morphology, basic biochemical characteristics, and isolation sources are also described in the species description section.

Keywords: 16S rRNA, bacterial diversity, unreported species

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INTRODUCTION

We isolated 20 unrecorded bacterial species from various soil samples collected in Korea and identified them as members of the phyla Actinobacteria, Bacteroidota, Firmicutes, and Proteobacteria. The present report focuses on the isolation and description of unrecorded species. At the time of writing, the phyla comprise Actinobacteria (including 6 classes), Bacteroidota (including 6 classes), Firmicutes (including 7 classes), and Proteobacteria (including 8 classes) based on the List of Prokaryotic names with Standing in Nomenclature (LPSN) (http://www.bacterio.net). The phylum Actinobacteria, one of the largest phyla within the domain *Bacteria*, is widely distributed in aquatic and terrestrial environments (Lawson, 2018). The phylum is comprised of Gram-stain-positive organisms with a high G + C content (Gao and Gupta, 2005); the phylum Bacteroidota is distributed in many different ecological niches (fresh water, ocean, soil, and gastrointestinal tract of animals) and known to degrade polymeric organic compounds (Thomas et al., 2011); the phylum Firmicutes is distributed in diverse environments and characterized as Gram-stain-positive, low G+C content and rod/coccus-shaped (Nahar, 2018); the phylum Pro*teobacteria* is one of the largest phyla within the domain *Bacteria* (Seong, 2019) and members of this phylum are the most versatile.

The present report focuses on the description of 20 bacterial species belonging to the phyla Actinobacteria, Bacteroidota, Firmicutes, and Proteobacteria that have not officially been reported in Korea.

MATERIALS AND METHODS

A total of 20 bacterial strains assigned to the phyla *Actinobacteria*, *Bacteroidota*, *Firmicutes*, and *Proteobacteria* were isolated from various soils, including agricultural soil, forest soil, and marine soil. Samples collected from each environment were independently processed, serially diluted, spread onto diverse culture agar media [R2A, Nutrient agar (NA), Tryptic Soy Agar (TSA), Luria-Bertani (LB) agar, PTYG, and Marine agar], and incubated at 28°C for 3–7 days (Table 1). All strains were purified as single colonies and stored as 15–17% glycerol suspension at – 80°C as well as lyophilized ampoules.

Colony morphology and cell size of the strains were observed by using transmission electron microscopy.

Electron micrograph of the strains are shown in Fig. 1. Gram reaction was performed according to the classic Gram procedure described by Doetsch (1981). Biochemical characteristics were tested by using API 20NE, API 32GN, and API ZYM galleries (bioMérieux) according to the manufacturer's instructions. Genomic DNA was extracted and the 16S rRNA gene was amplified by PCR with 27mf and 1492r universal bacterial primers (Weisburg et al., 1991). The 16S rRNA gene sequences of the related taxa were obtained from EzBioCloud server (Yoon et al., 2017). The 20 bacterial strains and related taxa (retrieved from the NCBI database) were aligned with SINA (v1.2.11) according to the SILVA seed alignment (http:// www.arb-silva.de; Pruesse et al., 2012). The evolutionary distances were calculated using a two-parameter model (Kimura, 1983). Phylogenetic trees were constructed using the neighbor-joining (Saitou and Nei, 1987) in the MEGA7 program (Kumar et al., 2016) with bootstrap values based on 1,000 replications (Felsenstein, 1985).

RESULTS AND DISCUSSION

The 20 strains were distributed into four phyla: Actinobacteria, Bacteroidota, Firmicutes, Proteobacteria (Table 1). Among these strains, EM1075, EM1106, EM1158 were coccus-shaped, WS2 was oval-shaped, and the other strains were rod-shaped (Fig. 1). Unrecorded bacteria were identified as 20 genera of Arthrobacter, Azospirillum, Bacillus, Barrientosiimonas, Corynebacterium, Flexivirga, Georgenia, Kocuria, Lysinibacillus, Mesorhizobium, Novosphingobium, Paenibacillus, Pandoraea, Paraburkholderia, Patulibacter, Pedobacter, Pontibacter, Pseudooceanicola, Sandaracinobacter, and Sphingoaurantiacus (Fig. 2). Here we report 20 unrecorded bacterial species in Korea belonging to 12 orders, which were isolated in Korea; 1 strain of Bogoriellales, 2 strains of Micrococcales, 2 strains of Dermatophilales, 1 strain of Mycobacteriales, 1 strain of Solirubrobacterales, 1 strain of Cytophagales, 1 strain of Sphingobacteriales, 3 strains of Bacillales, 3 strains of Burkholderiales, 3 strains of Sphingomonadales, 1 strain of Rhodospirillales, and 1 strain of Rhodobacterales.

Description of Pandoraea sputorum EM0341

Cells are Gram-stain-negative and rod-shaped. Cell size is 0.6–0.8 µm. Colonies are circular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for gluconate, caprate, adipate, malate, citrate, and phenyl-acetate in API 20NE; but negative for reduction of nitrates to nitrite (NO₃⁻>NO₂⁻), reduction of nitrates to nitrite (NO₃⁻>NO₂⁻), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -galactosidase

(PNPG), D-glucose, L-arabinose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, and D-maltose. Propionate, caprate, valerate, citrate, L-histidine, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, acetate, L-alanine, 3-hydroxy-benzoate, L-serine are utilized; but does not utilize D-mannitol, D-glucose, salicin, D-melibiose, L-fucose, D-sorbitol, L-arabinose, 2-ketogluconate, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, inositol, D-sucrose, D-maltose, itaconate, suberate, malonate, lactate, 5-ketogluconate, and glycogen. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), leucine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase in API ZYM; but negative for esterase lipase (C8), lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, *N*-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain EM0341 (=NIBRBAC000509124) was isolated from a forest soil sample, Seoul, Korea.

Description of Sandaracinobacter sibiricus EM0359

Cells are Gram-stain-negative and rod-shaped. Cell size is 1.0-1.2 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on R2A. Positive for D-glucose, L-arabinose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, D-maltose, gluconate, adipate, malate, and citrate in API 20NE; but negative for nitrates to nitrite ($NO_3^- > NO_2^-$), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, β -galactosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -galactosidase (PNPG), caprate, and phenyl-acetate. Utilizes D-mannitol, D-glucose, salicin, L-fucose, D-sorbitol, L-arabinose, citrate, 3-hydroxy-butyrate, L-proline, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, inositol, D-sucrose, D-maltose, suberate, malonate, lactate, 5-ketogluconate, and glycogen; but does not utilize D-melibiose, propionate, caprate, valerate, L-histidine, 2-ketogluconate, 4-hydroxy-benzoate, itaconate, acetate, L-alanine, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, naphtol-AS-BI-phosphohydrolase, and β -glucuronidase in API ZYM; but negative for lipase (C14), cystine arylamidase, α -galactosidase, β -galactosidase, β -glucuronidase, β -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain EM0359 (=NIBRBAC000509125) was isolated from a forest soil sample, Seoul, Korea.

Description of Paenibacillus edaphicus EM0662

Cells are Gram-stain-negative and rod-shaped. Cell

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Phylum	Class	Order	Family	Genus	Strain ID	NIBR ID	Most closely related species	Similarity (%)	Isolation source	Medium	Incubation conditions
		Bogoriellales	Bogoriellaceae	Georgenia	EM1187	NIBRBAC000509132	Georgenia muralis	7.66	Agricultural soil	R2A	28°C, 3 d
		Micrococcales	Micrococcaceae	Arthrobacter Kocuria	NC19 EM1106	NIBRBAC000509136 NIBRBAC000509129	Arthrobacter celericrescens Kocuria flava	9.99 9.99	Agricultural soil Agricultural soil	NA R2A	28°C, 3d 28°C, 3d
Actinobacteria	Acuitoniyceua	Dermatophilales	Dermacoccaceae	Flexivirga Barrientosiimonas	NSM-1 EM1075	NIBRBAC000509137 NIBRBAC000509128	Flexivirga endophytica Barrientosiimonas humi	99.9 99.2	Agricultural soil Forest soil	PTYG R2A	28°C, 3 d 28°C, 3 d
		Mycobacteriales	Corynebacteriaceae	Corynebacterium	RB28	NIBRBAC000509138	Corynebacterium glutamicum	6'66	Agricultural soil	R2A	28°C, 3 d
	Thermoleophilia	Solirubrobacterales	Patulibacteraceae	Patulibacter	EM1152	NIBRBAC000509130	Patulibacter brassicae	6.66	Agricultural soil	R2A	28°C, 3 d
Desteroidete	Cytophagia	Cytophagales	Hymenobacteraceae	Pontibacter	JN2214	NIBRBAC000509135	Pontibacter mucosus	99.1	Agricultural soil	R2A	28°C, 7 d
Daviel oluota	Sphingobacteriia	Sphingobacteriales	Sphingobacteriaceae	Pedobacter	WS11	NIBRBAC000509143	Pedobacter caeni	6.66	Agricultural soil	R2A	28°C, 3 d
Firmicutes	Bacilli	Bacillales	Paenibacillaceae Planococcaceae Bacillaceae	Paenibacillus Lysinibacillus Fredinand cohnia	EM0662 EM0665 JN1309	NIBRBAC000509126 NIBRBAC000509127 NIBRBAC000509134	Paenibacillus edaphicus Lysinibacillus odysseyi Bacillus timonensis	98.7 9.66 98.9	Forest soil Agricultural soil Agricultural soil	R2A R2A NA	28°C, 3 d 28°C, 3 d 28°C, 3 d
	Betaproteobacteria	Burkholderiales	Burkholderiaceae	Pandoraea Mesorhizobium Paraburkholderia	EM0341 FL16 WR2	NIBRBAC000509124 NIBRBAC000509133 NIBRBAC000509141	Pandoraea sputorum Mesorhizobium composti Paraburkholderia graminis	100.0 99.9 99.7	Forest soil Agricultural soil Agricultural soil	R2A LB R2A	28°C, 3 d 28°C, 3 d 28°C, 3 d
Proteobacteria		Sphingomonadales	Sphingosinicellaceae	Sandaracinobacter Sphingoaurantiacus	EM0359 EM1158	NIBRBAC000509125 NIBRBAC000509131	Sandaracinobacter sibiricus Sphingoaurantiacus polygramulatus	99.9 99.3	Forest soil Agricultural soil	R2A R2A	28°C, 3 d 28°C, 3 d
	Alphaproteobacteria		Sphingomonadaceae	Novosphingobium	RM1	NIBRBAC000509139	Novosphingobium soli	99.2	Agricultural soil	R2A	28°C, 3 d
		Rhodospirillales	Azospirillaceae	Azospirillum	WD4	NIBRBAC000509140	Azospirillum thiophilum	99.2	Agricultural soil	R2A	28°C, 3 d
		Rhodobacterales	Rhodobacteraceae	Pseudooceanicola	WS2	NIBRBAC000509142	NIBRBAC000509142 Pseudooceanicola nitratireducens	9.66	Marine soil	Marine	28°C, 3 d

Table 1. The taxonomic affiliations of isolated strains belonging to the phyla Actinobacteria, Bacteroidota, Firmicutes, and Proteobacteria.

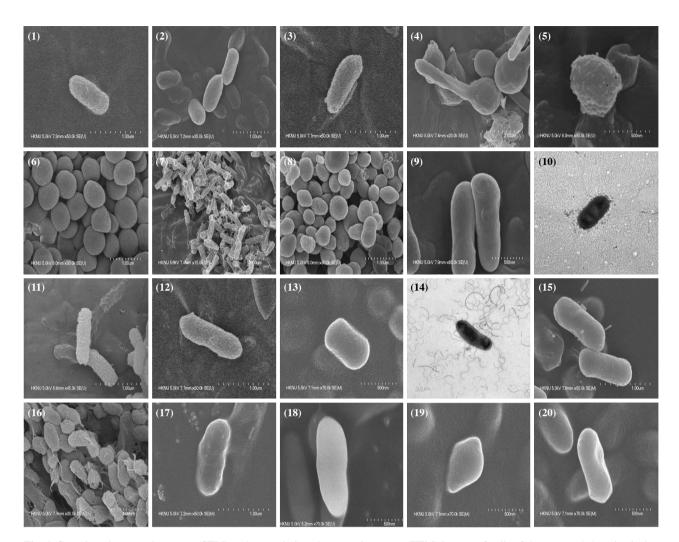


Fig. 1. Scanning electron microscopy (SEM) and transmission electron microscopy (TEM) images of cells of the unrecorded strains isolated in this study. Images 10 and 14 are TEM images, the rest are SEM images. The strains were cultured at their optimal growth conditions. (1) EM0341, (2) EM0359, (3) EM0662, (4) EM0665, (5) EM1075, (6) EM1106, (7) EM1152, (8) EM1158, (9) EM1187, (10) FL16, (11) JN1309, (12) JN2214, (13) NC19, (14) NSM-1, (15) RB28, (16) RM1, (17) WD4, (18) WR2, (19) WS2, (20) WS11.

size is 1.2-1.4 µm. Colonies are irregular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of nitrates to nitrite (NO3⁻ >NO₂⁻), β -Glucosidase (esculin hydrolysis), β -Galactosidase (PNPG), D-glucose, L-arabinose, D-mannose, *N*-acetyl-D-glucosamine, and D-maltose in API 20NE; but negative for indole production, glucose acidification, arginine dihydrolase, urease, protease (gelatin hydrolysis), D-mannitol, gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes D-glucose, D-melibiose, L-fucose, D-sorbitol, L-arabinose, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, D-sucrose, D-maltose, suberate, acetate, lactate, and glycogen; but does not utilize D-mannitol, salicin, propionate, caprate, valerate, citrate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, inositol, itaconate, malonate,

L-alanine, 5-ketogluconate, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), acid phosphatase, naphtol-AS-BI-phosphohydrolase, β -galactosidase, α -glucosidase, and β -glucosidase in API ZYM; but negative for lipase (C14), leucine arylamidase, valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -glucuronidase, *N*-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain EM0662 (=NIBRBAC000509126) was isolated from a forest soil sample, Seoul, Korea.

Description of Lysinibacillus odysseyi EM0665

Cells are Gram-stain-positive and rod-shaped. Cell size is $4.6-4.8 \mu m$. Colonies are circular, smooth, entire, and

curia flava EM1106 (OL305676) Nocuria Itava E Imi 106 (UCJJ095 re) Kocuria Itava E Imi 106 (UCJJ095 re) Kocuria oceani FXLB 095² (JF346427) Kocuria oceani DXB 02047⁷ (VST756) Cocuria polaris CMS 766⁷ (JSUH01000031) Cocuria salina Hrv1ho¹ (LT674162) Arthrobacter Indicionaris NRFC 12137⁷ (BAECO Arthrobacter Indicationaris NRFC 12137⁷ (BAECO Arthrobacter Celericirescens NC19 (OK602676) Arthrobacter Celericirescens NC19 (OK602676) Arthrohacter celericrescens NEALLSA2 (MH063435) Amrobacter celencrescens NEAU-SA2 (MH053435) Arthrobacter cupress i CGMCC 1,10783 (1)21/076144) Arthrobacter silvisoli NEAU-SA1¹ (QRCU01000009) Georgenia subflava Y32⁷ (KP123395) - Georgenia wutainyi Z294¹ (MN020372)) Georgenia muralis EM1187 (OL305683) Georgenia muralis EM1147 (OL305683) Georgenia muralis SDM 14148¹ (RRRA01000001) – Georgenia adeguensis 2C6-43¹ (HO246163) Georgenia soli DSM 21803⁶ (PDJI010000004) Barrientosiimonas knmi 3⁶ (JC46171) Barrientosiimonas marna MSW-42⁶ (HC792770) Barrientosiimonas humi EM1075 (OL305687) Deveto kene o DDM 1106¹ (C400410000006) ŀ Demetria terzagena DSM 11295¹ (AOXMO1000002) Branchilibius hedensis Mer 29717¹ (AB520822) Flexivirga endophytica NSM-1 (01.305668) Flexivirga endophytica (NM 7505¹ (KX151661) Flexivirga alba ST13² (AB539735) Flexiving alba ST13' (ABS9735) Flexiving Laten BO-16' (MH094629) Flexiving a crase R1' (KR025522) Corynebacterium glutamicum ATCC 13032' (BA00065) Corynebacterium glutamicum ATCC 13032' (BA00065) Corynebacterium cullicuta JC14' (CP015522) Corynebacterium cullicuta JC14' (CP015523) Corynebacterium cullicuta JC14' (CP015524) Corynebacterium cullicuta JC14' (CP015534) Corynebacterium cruditactis J216¹ (CP015622) Corynebacterium calinae DSN 2014¹⁷ (CP015622) Corynebacterium seanareeee N24 (MK395364) Corynebacterium desare DSN 2014¹⁷ (CP00320) Patulibacter brassicae SD (K151458) Patulibacter medicamentivorans I11 (AGUD0100008) Patulibacter americanus DSM 1005¹ (K1912613) Patulibacter americanus DSM 1005¹ (K1912710) Patulibacter americanus DSM 1005¹ (K191271 andcohnia humi LMG 22167^T (AJ627210) ndcohnia aciditolerans YN-1^T (MG589508 Fredinandcohnia onubensis 0911MAR22V3^T (NSEB01000017) If Predinandochnia onubensis 0911MAR22292¹ (NESB01) "Fredinandochnia saidurans NUUC7312 (KX004715) Azospirillum thiophilum DSM 21624¹ (AEL0100007) Azospirillum thiophilum DSM 21624¹ (AEL0100007) Azospirillum nasamy M2T282¹ (GQ24693) Azospirillum nasamy M2T28¹ (GQ24693) Azospirillum naseminy M2T38² (GQ24693) Azospirillum naseminy M2T48² (GQ24693) Azospirillum naseminy M2T48² (KV1415) Mesorhizobium plantfarium IMG 1182¹ (V1415) Mesorhizobium sandmizum CNPH 5310¹ (GM201000020) rhizobium tamadayense DSM 28320^T (RQXT01000019) hizobium composti FL16 (OL305692) Pseudocceanicola nitratireducens DSM 29619⁺ (FOLX)
Sphingoaurantiacus polygranulatus EM1158 (OL305686)
Sphingoaurantiacus polygranulatus MC 3718⁺ (KP274054)
Sphingoaurantiacus capsulatus YL133⁺ (KF274054)
Sandarakinorhabdus rubra MO-4⁺ (LC491550)
Sandarakinorhabdus rubra MO-4⁺ (LC491550) 1000011) Sandarahitinitadus tuna MC4 (L4491580) daracinobacter neustonicus PANC 2813 "(KR552095) daracinobacter sibiricus EM0359 (0L305689) daracinobacter sibiricus RB16-17" (Y10678) Sphirogošinitella microcystinivorans DSM 19791" (RBWX01000007) 2) Sphingosinicella merocystinivorans DSM 19/91 (RBW Sphingosinicella xenopepiditylica 3-2W4/ (AY950663) Sphingosinicella soli KSL-125¹ (DO087403) Sphingomonas tabacisoli X1-8¹ (MF370621) Sphingomonas changbaiensis NBRC 104936^{*} (BBWU010 Vovosphingobium naphthalenivorans NBRC 102051^{*} (BC^{*}) (BCTX01000153) Novosphingobium panipatense SM16^T (jgj.1118302) Novosphingobium soli CC-TPE-1^T (FJ425737) Novosphingobium soli CC-TPE-1^T (FJ425737) Novosphingobium kindniclasticum LE142 (FJ425737) Novosphingobium aindaniclasticum LE142 (ATHLD1000125) Novosphingobium sain RMI (OL305658) Novosphingobium barchaimii LL02¹ (KO130454) Pandoraea sputorum EM0341 (OL305650) Pandoraea sputorum EM0341 (OL305650) Pandoraes sputorum EM0341 (c)L36589) Pandoraes aputorum EM0341 (c)CP010431 Pandoraes commovens LMG 31010⁻ (LR58843) Pandorae aquancia LMG 31011⁻ (LR58849) Pandoraes aquancia LMG 31117⁻ (LR58849) Pandoraes aquancia LMG 31117⁻ (LR58849) Pandoraes outaindroans DSM 2507⁻ (CP01123) Panaburkholderia phyofinmas Ps-IN⁻ (CP00103) Panaburkholderia parminis WR2 (C)L368639) Panaburkholderia graminis KR2 (C)L368639) Panaburkholderia graminis CAD1M⁻ (U)46839) Panaburkholderia graminis CAD1M⁻ (U)46839) Panaburkholderia graminis CAD1M⁻ (U)46868) pacter uncouss P30⁻ (KP081651) nhlbacter mucosus JN2214 (MT347086) nhlbacter mucosus PB3' (KP6)16351 Pontibacter chinhatensis LP51' (jgl. 110242) Pontibacter instensis NIL-005' (SI987494) ntibacter kortensis X1-41' (DO888330) ontibacter mangrovi HB172049' (MX253282) Prevolutionum 31463' (MT441864), Pedobacter Ioliolum 31455 (M1441604) Pedobacter duraquae WB2.1-25^T (AM491368) Pedobacter gandavensis LMG 31462^T (MT44 Pedobacter steynii DSM 19110^T (jgi.1085793) **Pedobacter caeni WS11 (OL305697)** obacter caeni LMG 22862^T (AJ786798)

Fig. 2. A neighbor-joining phylogenetic tree based on 16S rRNA gene sequences showing the relationship between the strains isolated in this study and their relatives. Bootstrap values (>70%) are shown at the nodes. Bar: 0.050 substitutions per nucleotide position.

white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of glucose acidification, arginine dihydrolase, and β -Glucosidase (esculin hydrolysis) in API 20NE; but negative for nitrates to nitrite (NO₃⁻ >NO₂⁻), reduction of nitrates to nitrogen, indole production, urease, protease (gelatin hydrolysis), β -galactosidase (PNPG), D-glucose, L-arabinose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, D-maltose, gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes 3-hydroxy-butyrate, acetate, and L-alanine; but does not utilize D-mannitol, D-glucose, salicin, D-melibiose, L-fucose, D-sorbitol, L-arabinose, propionate, caprate, valerate, citrate, L-histidine, 2-ketogluconate, 4-hydroxy-benzoate, L-proline, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, inositol, D-sucrose, D-maltose, itaconate, suberate, malonate, lactate, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase Lipase (C8), naphtol-AS-BI-phosphohydrolase, and β -glucosidase in API ZYM; but negative for lipase (C14), leucine arylamidase, valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain EM0665 (= NI-BRBAC000509127) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of Barrientosiimonas humi EM1075

Cells are Gram-stain-positive and coccus-shaped. Cell size is 0.55-0.57 µm. Colonies are circular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), D-glucose, D-mannitol, N-acetyl-D-glucosamine, D-maltose, gluconate, adipate, malate, and phenyl-acetate; but negative for nitrates to nitrite ($NO_3 > NO_2$), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, β -Galactosidase (PNPG), L-arabinose, D-mannose, caprate, and citrate. Utilizes D-mannitol, D-glucose, D-melibiose, propionate, valerate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, N-acetyl-D-glucosamine, D-sucrose, D-maltose, suberate, malonate, acetate, lactate, L-alanine, glycogen, and 3-hydroxy-benzoate; but does not utilize salicin, L-fucose, D-sorbitol, L-arabinose, caprate, citrate, L-histidine, 2-ketogluconate, L-proline, L-rhamnose, D-ribose, inositol, itaconate, 5-ketogluconate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, naphtol-AS-BI-phosphohydrolase, α -galactosidase, α -glucosidase, β -glucosidase, and α -mannosidase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, β -galactosidase, β -glucuronidase, *N*-acetyl- β -glucosaminidase, and α -fucosidase. Strain EM1075 (=NIBRBAC000509128) was isolated from a forest soil sample, Seoul, Korea.

Description of Kocuria flava EM1106

Cells are Gram-stain-positive and coccus-shaped. Cell size is 0.5-0.7 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of nitrates to nitrite (NO3⁻ $>NO_2$), β -Galactosidase (PNPG), D-glucose, D-maltose, gluconate, adipate, malate, and phenyl-acetate; but negative for indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), L-arabinose, N-acetyl-D-glucosamine, caprate, and citrate. Utilizes D-mannitol, D-glucose, D-melibiose, valerate, 4-hydroxy-benzoate, L-proline, D-sucrose, D-maltose, suberate, 5-ketogluconate, 3-hydroxy-benzoate, and L-serine; but does not utilize salicin, L-fucose, L-arabinose. Propionate, caprate, citrate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, itaconate, malonate, and glycogen. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, naphtol-AS-BI-phosphohydrolase, α -galactosidase, β -galactosidase, and α -glucosidase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, β -glucuronidase, β -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain EM1106 (=NIBRBAC000509129) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of Patulibacter brassicae EM1152

Cells are Gram-stain-positive and rod-shaped. Cell size is 1.3-1.5 µm. Colonies are circular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of L-arabinose; but negative for nitrates to nitrite ($NO_3^- > NO_2^-$), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -Galactosidase (PNPG), D-glucose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, D-maltose, gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes L-arabinose, valerate, and acetate; but does not utilize D-mannitol, D-glucose, salicin, D-melibiose, L-fucose, D-sorbitol, propionate, caprate, citrate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, inositol, D-sucrose, D-maltose, itaconate, suberate, malonate, lactate, L-alanine, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, cystine arylamidase, acid phosphatase, and naphtol-AS-BI-phosphohydrolase; but negative for lipase (C14), valine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain EM1152 (=NIBRBAC000509130) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of *Sphingoaurantiacus polygranulatus* EM1158

Cells are Gram-stain-negative and coccus-shaped. Cell size is 0.5–0.7 µm. Colonies are circular, smooth, entire, and red pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of D-glucose, L-arabinose, and N-acetyl-D-glucosamine; but negative for nitrates to nitrite $(NO_3^- > NO_2^-)$, reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -Galactosidase (PNPG), D-mannose, D-mannitol, D-maltose, gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. D-Glucose, L-arabinose, 3-hydroxy-butyrate, L-proline, L-rhamnose, N-acetyl-D-glucosamine, and D-sucrose; but does not utilize D-mannitol, salicin, D-melibiose, L-fucose, D-sorbitol, propionate, caprate, valerate, citrate, L-histidine, 2-ketogluconate, 4-hydroxy-benzoate, D-ribose, inositol, D-maltose, itaconate, suberate, malonate, acetate, lactate, L-alanine, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, and N-acetyl- β -glucosaminidase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, α -mannosidase, and α -fucosidase. Strain EM1158 (=NIBRBAC000509131) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of Georgenia muralis EM1187

Cells are Gram-stain-positive and rod-shaped. Cell size is $1.5-1.7 \mu m$. Colonies are circular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of D-glucose, L-arabinose, D-mannose, D-mannitol, *N*-acetyl-D-glucosamine, and D-maltose; but negative for nitrates to nitrite (NO₃⁻ >NO₂⁻), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hy-

drolysis), β -Galactosidase (PNPG), gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes D-mannitol, D-glucose, salicin, L-arabinose, N-acetyl-D-glucosamine, D-sucrose, and D-maltose; but does not utilize D-melibiose, L-fucose, D-sorbitol, propionate, caprate, valerate, citrate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, L-rhamnose, D-ribose, inositol, itaconate, suberate, malonate, acetate, lactate, L-alanine, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for esterase (C4), esterase lipase (C8), leucine arylamidase, naphtol-AS-BI-phosphohydrolase, α -glucosidase, and N-acetyl- β -glucosaminidase; but negative for alkaline phosphatase, lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, α -galactosidase, β -galactosidase, β -glucuronidase, β -glucosidase, α -mannosidase, and α -fucosidase. Strain EM1187 (=NI-BRBAC000509132) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of Mesorhizobium composti FL16

Cells are Gram-stain-negative and rod-shaped. Cell size is 0.9-1.1 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on LB. Positive for reduction of nitrates to nitrite $(NO_3^- > NO_2^-)$, urease, β -glucosidase (esculin hydrolysis), D-glucose, L-arabinose, D-mannitol, N-acetyl-D-glucosamine, and D-maltose; but negative for indole production, glucose acidification, arginine dihydrolase, protease (gelatin hydrolysis), β -Galactosidase (PNPG), gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes D-mannitol, D-glucose, L-fucose, D-sorbitol, L-arabinose, propionate, valerate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, L-proline, N-acetyl-D-glucosamine, D-ribose, inositol, D-sucrose, D-maltose, acetate, lactate, and L-alanine; but does not utilize salicin, D-melibiose, caprate, citrate, 4-hydroxy-benzoate, L-rhamnose, itaconate, suberate, malonate, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, α -glucosidase, and β -glucosidase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain FL16 (=NIBRBAC000509133) was isolated from an agricultural soil sample, Chungcheongnam-do, Korea.

Description of Bacillus timonensis JN1309

Cells are Gram-stain-negative and rod-shaped. Cell

size is 1.1-1.3 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on NA. Positive for reduction of β -glucosidase (esculin hydrolysis), β -Galactosidase (PNPG), D-glucose, L-arabinose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, D-maltose, and gluconate: but negative for nitrates to nitrite $(NO_3^- > NO_2^-)$, reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, protease (gelatin hydrolysis), caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes D-mannitol, D-glucose, D-melibiose, L-arabinose, valerate, L-histidine, N-acetyl-D-glucosamine, D-sucrose, D-maltose, acetate, lactate, and glycogen; but does not utilize salicin, L-fucose, D-sorbitol, propionate, caprate, citrate, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, L-rhamnose, D-ribose, inositol, itaconate, suberate, malonate, L-alanine, 5-ketogluconate, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for esterase (C4), esterase lipase (C8), lipase (C14), naphtol-AS-BI-phosphohydrolase, β -galactosidase, α -glucosidase, β -glucosidase, and N-acetyl- β -glucosaminidase; but negative for alkaline phosphatase, leucine arylamidase, valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, α -galactosidase, β -glucuronidase, α -mannosidase, and α -fucosidase. Strain JN1309 (=NIBRBAC000509134) was isolated from an agricultural soil sample, Jeju Island, Korea.

Description of Pontibacter mucosus JN2214

Cells are Gram-stain-negative and rod-shaped. Cell size is 1.3-1.5 µm. Colonies are circular, smooth, entire, and pink pigmented after 7 days of incubation at 28°C on R2A. Positive for reduction of urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), D-glucose, L-arabinose, D-mannose, D-mannitol, and N-acetyl-D-glucosamine; but negative for nitrates to nitrite $(NO_3^->NO_2^-)$, reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, β -Galactosidase (PNPG), D-maltose, gluconate, caprate, adipate, malate, citrate, and phenyl-acetate. Utilizes D-mannitol, D-glucose, L-fucose, L-arabinose, L-rhamnose, and N-acetyl-D-glucosamine; but does not utilize salicin, D-melibiose, D-sorbitol, propionate, caprate, valerate, citrate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, D-ribose, inositol, D-sucrose, D-maltose, itaconate, suberate, malonate, acetate, lactate, L-alanine, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, cystine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, α -galactosidase, α -glucosidase, β -glucosidase, and *N*-acetyl- β -glucosaminidase;

but negative for lipase (C14), trypsin, α -chymotrypsin, β -galactosidase, β -glucuronidase, α -mannosidase, and α -fucosidase. Strain JN2214 (=NIBRBAC000509135) was isolated from an agricultural soil sample, Jeju Island, Korea.

Description of Arthrobacter celericrescens NC19

Cells are Gram-stain-positive and rod-shaped. Cell size is 0.65-0.85 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on NA. Positive for reduction of β -glucosidase (esculin hydrolysis), β -Galactosidase (PNPG), D-glucose, L-arabinose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, D-maltose, gluconate, malate, citrate, and phenyl-acetate; but negative for nitrates to nitrite ($NO_3 > NO_2$), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, protease (gelatin hydrolysis), caprate, and adipate. Utilizes D-mannitol, D-glucose, salicin, D-melibiose, D-sorbitol, L-arabinose, citrate, L-histidine, 2-ketogluconate, 4-hydroxy-benzoate, L-proline, N-acetyl-D-glucosamine, inositol, D-sucrose, D-maltose, lactate, L-alanine, 3-hydroxy-benzoate propionate, and L-serine; but does not utilize L-fucose, propionate, caprate, valerate, 3-hydroxy-butyrate, L-rhamnose, D-ribose, itaconate, suberate, malonate, acetate, 5-ketogluconate, and glycogen. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, cystine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, α -galactosidase, β -galactosidase, α -glucosidase, β -glucosidase, and α -mannosidase; but negative for lipase (C14), trypsin, α -chymotrypsin, β -glucuronidase, N-acetyl- β -glucosaminidase, and α -fucosidase. Strain NC19 (=NIBRBAC 000509136) was isolated from an agricultural soil sample, Chungcheongnam-do, Korea.

Description of Flexivirga endophytica NSM-1

Cells are Gram-satin-positive and rod-shaped. Cell size is 2.5–2.7 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on PTYG. Positive for reduction of urease, D-glucose, L-arabinose, D-mannose, gluconate, adipate, malate, and phenyl-acetate; but negative for nitrates to nitrite (NO₃⁻ >NO₂⁻), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -Galactosidase (PNPG), D-mannitol, *N*-acetyl-D-glucose, L-fucose, L-arabinose, propionate, valerate, 2-ketogluconate, 3-hydroxy-butyrate, L-rhamnose, D-ribose, itaconate, suberate, acetate, lactate, 5-ketogluconate, and 3-hydroxy-benzoate; but does not utilize D-mannitol, sal-

icin, D-melibiose, D-sorbitol, caprate, citrate, L-histidine, 4-hydroxy-benzoate, L-proline, N-acetyl-D-glucosamine, inositol, D-sucrose, D-maltose, malonate, L-alanine, glycogen, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, trypsin, acid phosphatase, and naphtol-AS-BI-phosphohydrolase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain NSM-1 (=NIBRBAC000509137) was isolated from an agricultural soil sample, Chungcheongnam-do, Korea.

Description of Corynebacterium glutamicum RB28

Cells are Gram-stain-positive and rod-shaped. Cell size is 1.1-1.3 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of nitrates to nitrite $(NO_3 > NO_2)$, glucose acidification, urease, D-glucose, D-mannose, D-maltose, gluconate, and citrate; but negative for indole production, arginine dihydrolase, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -Galactosidase (PNPG), L-arabinose, D-mannitol, N-acetyl-D-glucosamine, caprate, adipate, malate, and phenyl-acetate. Utilizes D-glucose, propionate, citrate, 4-hydroxy-benzoate, D-ribose, inositol, D-sucrose, D-maltose, malonate, acetate, lactate, and 3-hydroxy-benzoate; but does not utilize D-mannitol, salicin, D-melibiose, L-fucose, D-sorbitol, L-arabinose, caprate, valerate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, L-proline, L-rhamnose, N-acetyl-D-glucosamine, itaconate, suberate, L-alanine, 5-ketogluconate, glycogen, and L-serine. Enzymatic activity reaction was positive for esterase (C4), leucine arylamidase, acid phosphatase, and naphtol-AS-BI-phosphohydrolase; but negative for alkaline phosphatase, esterase lipase (C8), lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain RB28 (=NIBRBAC000509138) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of Novosphingobium soli RM1

Cells are Gram-satin-negative and rod-shaped. Cell size is 0.8–1.0 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of β -glucosidase (esculin hydrolysis), D-glucose, D-maltose, and malate; but negative for nitrates to nitrite (NO₃⁻>NO₂⁻), reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, protease (gelatin hydrolysis), β -Galactosidase (PNPG), L-arabinose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, gluconate, caprate, adipate, citrate, and phenyl-acetate. Utilizes D-glucose, valerate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, L-rhamnose, D-sucrose, D-maltose, and acetate; but does not utilize D-mannitol, salicin, D-melibiose, L-fucose, D-sorbitol, L-arabinose, propionate, caprate, citrate, L-histidine, 2-ketogluconate, N-acetyl-D-glucosamine, D-ribose, inositol, itaconate, suberate, malonate, lactate, L-alanine, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, α -glucosidase, and β -glucosidase; but negative for lipase (C14), cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain RM1 (= NIBRBAC000509139) was isolated from an agricultural soil sample, Chungcheongnam-do, Korea.

Description of Azospirillum thiophilum WD4

Cells are Gram-stain-negative and rod-shaped. Cell size is 1.4-1.6 µm. Colonies are irregular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of nitrates to nitrite $(NO_3^- > NO_2^-)$, urease, β -glucosidase (esculin hydrolysis), β -Galactosidase (PNPG), D-glucose, L-arabinose, D-mannitol, gluconate, caprate, malate, and phenyl-acetate; but negative for indole production, glucose acidification, arginine dihydrolase, protease (gelatin hydrolysis), D-mannose, N-acetyl-D-glucosamine, D-maltose, adipate, and citrate. Utilizes D-mannitol, D-glucose, D-sorbitol, L-arabinose, propionate, caprate, valerate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, D-ribose, malonate, acetate, lactate, and L-alanine; but does not utilize salicin, D-melibiose, L-fucose, citrate, L-rhamnose, N-acetyl-D-glucosamine, inositol, D-sucrose, D-maltose, itaconate, suberate, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, β -galactosidase, and β -glucosidase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -glucuronidase, α -glucosidase, N-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain WD4 (=NIBRBAC000509140) was isolated from an agricultural soil sample, Chungcheongnam-do, Korea.

Description of Paraburkholderia graminis WR2

Cells are Gram-stain-negative and rod-shaped. Cell

size is 1.4-1.6 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of nitrates to nitrite $(NO_3 > NO_2)$, β -Galactosidase (PNPG), D-glucose, D-mannose, D-mannitol, N-acetyl-D-glucosamine, gluconate, caprate, malate, and citrate; but negative for indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), L-arabinose, D-maltose, adipate, and phenyl-acetate. Utilizes D-mannitol, D-glucose, L-fucose, caprate, citrate, 2-ketogluconate, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, inositol, D-sucrose, and 5-ketogluconate; but does not utilize salicin, D-melibiose, D-sorbitol, L-arabinose, propionate, valerate, L-histidine, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, D-maltose, itaconate, suberate, malonate, acetate, lactate, L-alanine, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, acid phosphatase, naphtol-AS-BI-phosphohydrolase, α -galactosidase, β -galactosidase, and N-acetyl- β -glucosaminidase; but negative for lipase (C14), cystine arylamidase, trypsin, α -chymotrypsin, β -glucuronidase, α -glucosidase, β -glucosidase, α -mannosidase, and α -fucosidase. Strain WR2 (=NI-BRBAC000509141) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

Description of Pseudooceanicola nitratireducens WS2

Cells are Gram-stain-negative and oval-shaped. Cell size is 0.7-0.9 µm. Colonies are circular, smooth, entire, and yellow pigmented after 3 days of incubation at 28°C on Marine agar. Positive for reduction of nitrates to nitrite $(NO_3 > NO_2)$, D-glucose, D-mannitol, D-maltose, gluconate, malate, citrate, and phenyl-acetate; but negative for indole production, glucose acidification, arginine dihydrolase, urease, β -glucosidase (esculin hydrolysis), protease (gelatin hydrolysis), β -Galactosidase (PNPG), L-arabinose, D-mannose, N-acetyl-D-glucosamine, caprate, and adipate. D-mannitol, D-glucose, citrate, D-sucrose, D-maltose, L-alanine; but does not utilize salicin, D-melibiose, L-fucose, D-sorbitol, L-arabinose, propionate, caprate, valerate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, L-rhamnose, N-acetyl-D-glucosamine, D-ribose, inositol, itaconate, suberate, malonate, acetate, lactate, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, acid phosphatase, and naphtol-AS-BI-phosphohydrolase; but negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α -chymotrypsin, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, *N*-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase. Strain WS2 (=NIBRBAC000509142) was isolated from a marine soil sample, Jeju Island, Korea.

Description of Pedobacter caeni WS11

Cells are Gram-stain-negative and rod-shaped. Cell size is 1.0-1.2 µm. Colonies are irregular, smooth, entire, and white pigmented after 3 days of incubation at 28°C on R2A. Positive for reduction of β -glucosidase (esculin hydrolysis), β -Galactosidase (PNPG), D-glucose, L-arabinose, D-mannose, N-acetyl-D-glucosamine, D-maltose, and malate; but negative for nitrates to nitrite $(NO_3^- > NO_2^-)$, reduction of nitrates to nitrogen, indole production, glucose acidification, arginine dihydrolase, urease, protease (gelatin hydrolysis), D-mannitol, gluconate, caprate, adipate, citrate, and phenyl-acetate. Utilizes D-glucose, salicin, D-melibiose, L-arabinose, N-acetyl-D-glucosamine, D-sucrose, and D-maltose; but does not utilize D-mannitol, L-fucose, D-sorbitol, propionate, caprate, valerate, citrate, L-histidine, 2-ketogluconate, 3-hydroxy-butyrate, 4-hydroxy-benzoate, L-proline, L-rhamnose, D-ribose, inositol, itaconate, suberate, malonate, acetate, lactate, L-alanine, 5-ketogluconate, glycogen, 3-hydroxy-benzoate, and L-serine. Enzymatic activity reaction was positive for alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, trypsin, α -chymotrypsin, acid phosphatase, naphtol-AS-BI-phosphohydrolase, α -galactosidase, β -galactosidase, α -glucosidase, β -glucosidase, N-acetyl- β -glucosaminidase, and α -mannosidase; but negative for lipase (C14), cystine arylamidase, β -glucuronidase, and α -fucosidase. Strain WS11 (= NIBRBAC000509143) was isolated from an agricultural soil sample, Gyeongsangnam-do, Korea.

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