# 1 Table S1. Identified soil bacteria obtained as culturable on MWA' containing 0.25%

### 2 D-glucose and 0.25% DL-malic acid

Soil Isolate	V5	V10	V20	V30
G'1	Stenotrophomonas sp.	NI	Arthrobacter sp.*	Paenibacillus sp.
G'2	Pseudomonas sp.	Luteibactor sp.	Leifsonia sp.*	Leifsonia sp.
G'3	Stenotrophomonas sp.	NI	Leifsonia sp.*	Burkholderia sp.
G'4	Pseudomonas sp.	Burkholderia sp.	Paenibacillus borealis*	Burkholderia sp.
G'5	Leifsonia sp.	NI	Leifsonia sp.	Leifsonia sp.
G'6	Pseudomonas sp.	Burkholderia sp.	Microbacteriaceae	Micrococcineae
G'7	Janthinobacterium sp.	NI	Paenibacillus sp.	Burkholderia sp.
G'8	Pseudomonas sp.	Leifsonia sp.	Arthrobacter sp.	Paenibacillus sp.
G'9		Burkholderia sp.	Microbacterium sp.	NI
G'10		Pseudomonas sp.	Arthrobacter sp.	NI
G'11		Pseudomonas sp.	Arthrobacter sp.	
G'12		Burkholderia sp.		

# 5 Table footnote

6	Conditions and methods for isolation of bacteria on MWA' were the same as for Table 4,
7	except 1.5% agar for gel matrix. Distinguishable colony-forming bacteria in the plate culture
8	were isolated as many as possible. Isolates were preliminary identified with their 16S rRNA
9	gene regions partially sequenced (approximately 150 pb containing the V6 region of their 16S
10	rRNA gene). Note that MWG'-culturable bacteria of genera Janthinobacterium,
11	Paenibacillus, Arthrobacter and Luteibacter obtained were all minor isolates.
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# 15 Table S2. Effect of gellan gum as matrix to increase acetylene reduction of microfloral

#### 16 communities in Shizunai farmland Andisol.

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	Medium ( $0.05\%$ D-mannitol-containing Winogradsky's mineral mix.)				
Source	Liquid	0.2% agar soft gel	0.3% gellan gum soft gel		
	Activity as nmol $d^{-1}$ of $C_2H_4$				
Shizunai 5	$0 \pm 0$	$0.7 \pm 0.6$	$42.7 \pm 20.1$		
Shizunai 10	$0.6 \pm 1.3$	$0.5 \pm 1.1$	$26.0 \pm 3.4$		
Shizunai 15	$2.4 \pm 1.4$	$3.2 \pm 0.2$	$308.0 \pm 412.9$		

19	After preculturing for 14 days, acetylene gas was injected to the headspace of each culture
20	vial to a 10% concentration (v/v), and the culture was further incubated for 7 days at $15^{\circ}$ C.
21	Three different conditions (liquid, 0.2% agar, and 0.3% gellan gum) for Winogradsky's
22	medium (0.05% D-mannitol as carbon source) were tested and incubation was as previously
23	described for acetylene reduction. Values are means and standard deviations for five
24	replicates.
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#### 32 Fig. S1. Appropriate temperature for N<sub>2</sub>-fixation by cryophilic bacterial microbiota

33 Responses of Beijerinckia indica subsp. indica IFO 3744 and V30 microfloral community to incubation temperature. The most active bacterial microbiota sample from East Siberian 34Taiga forest bed soil, V30, showed high acetylene reduction at 15°C when cultured in a gellan 3536 gum soft gel. This representative response of V30 as a cryophilic microbiota was also shown at higher temperature. Even in gellan gum medium, V30 failed to exhibit acetylene reduction 3738at either 20°C or 25°C. Thus, the profitable temperature range for acetylene reduction of 39 cryophilic bacterial microbiota was narrow, near 15°C. In contrast, B. indica subsp. indica 40IFO 3744, used as a reference bacterium for free-living N-fixers, showed higher acetylene reduction at 20°C and 25°C, but low at 15°C. In addition, this bacterium showed almost the 41 42same level of acetylene reduction in agar and gellan gum.

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#### A: MWA-grown bacteria





#### B: MWG'-culturable bacteria



# Fig. S2 Phylogenetic alignment for 16S rRNA gene sequences of the soil bacteria from East Siberian forest bed soil

Phylogenetic trees were constructed for the bacterial isolates obtained from 1.0% 54sucrose-containing MWA plates (A) and 0.25% D-glucose plus 0.25% DL-malic 5556acid-containing MWG (MWG') plates (B). For the phylogenetic analysis, Clustal W was used. In this program, 1.36 pkb (positioned from 86 to 1443 of Escherichia coli) of the 16S rRNA 57gene sequences among the collections of MWA-grown bacteria was compared. Genera 5859Pseudomonas and Burkholderia formed two large groups among the isolates on MWA plate culture. These two genera are diverse in bacterial species based on their clustering in tree of A. 60 61 A 1.24 kbp region (positioned from 105 to 1345 of E. coli) of the 16S rRNA gene sequences 62 was used in the clustering for the MWG'-culturable bacteria as shown in tree **B**. A cluster of 63 Pseudomonas spp. shows that the soil bacterial microbiota of samples is diverse at the species 64 level of genus Pseudomonas. Genera Luteibacter and Paenibacillus show a similar tendency. 65 66 67 68 69 70

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87 In this preliminary experiment, we used pure isolates to show acetylene reduction-positive 88 responses. Acetylene reduction of the tested bacteria in gellan gum medium is shown with 89 filled symbols, while open symbols indicate those cultured in agar medium. The assays for 90 each bacterial isolate were replicated five times, but the OD was not very consistent because 91of the presence of the gel in the medium. Nevertheless, this figure shows that some bacteria, 92particularly those cultured in gellan gum, showed slight positive correlation between OD and 93 acetylene reduction. This is reasonable because bacteria that are able to grow in 94nitrogen-deficient medium beyond an OD of 0.05 at 665 nm are most likely diazotrophic 95 bacteria. However, a few exceptions are also observed. For example, V20-A9 (Pseudomonas sp.) showed a relatively low OD, but it showed relatively high acetylene reduction in gellan 96 97 gum medium. Another remarkable bacterium was V30-A6 (Burkholderia xenovorans), which uniquely both grew in agar and gellan gum media and had the highest acetylene reduction 98 99 among tested bacteria. OD at 665 nm was measured in 96-well plates, in which 200 µl of 100 agitated gelling medium had been poured (depth of the medium, 5.2 mm), using a CENios 101 microplate reader (Tecan, Männedorf, Switzarland). The value was used without any 102correction for the light path length. Because there was less-thick medium in the well, OD 103 values were relatively low, but the standard curve gave reliable values.

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