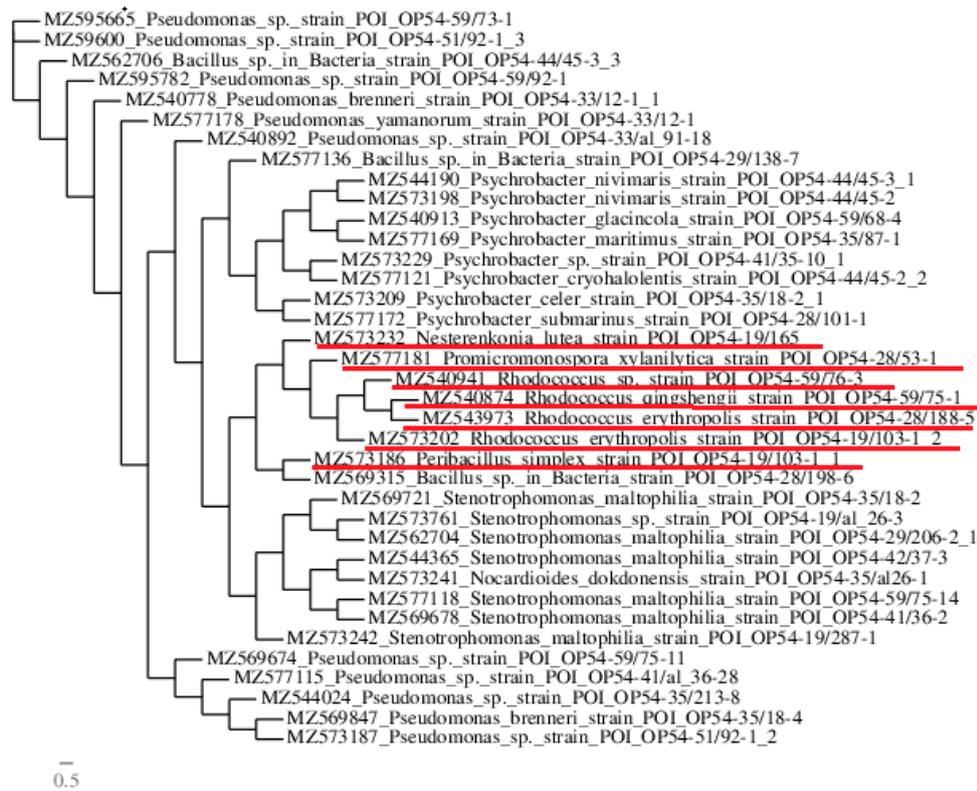
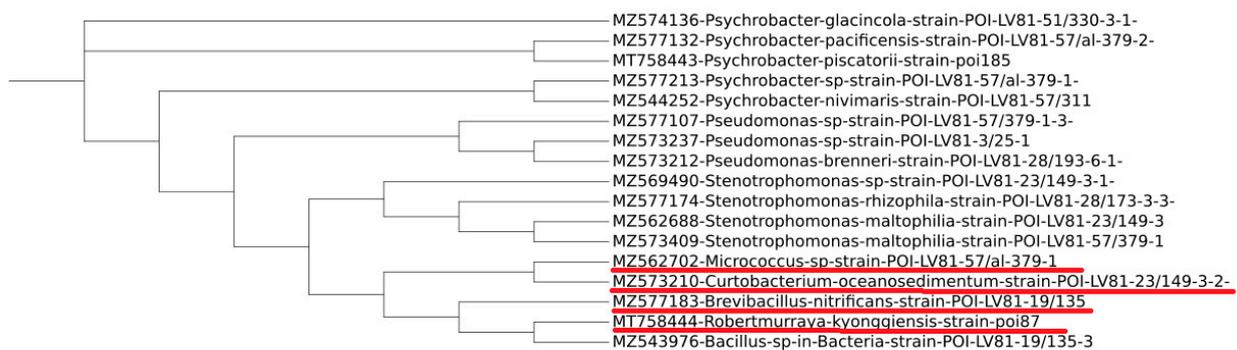


**Table S1.** Additional materials 1. Composition of media used.

<b>Microbiological Media</b>	<b>Media composition (g/l)</b>
marine ammonium medium 1313 (Marine ammonium mineral salts)	NaCl - 20, (NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub> -1, CaCl <sub>2</sub> • 2H <sub>2</sub> O - 0.2, MgSO <sub>4</sub> • 7H <sub>2</sub> O - 1, FeSO <sub>4</sub> • 7H <sub>2</sub> O - 0.002, Na <sub>2</sub> WO <sub>4</sub> • 2H <sub>2</sub> O - 0.003, Na <sub>2</sub> MoO <sub>4</sub> • 2H <sub>2</sub> O - 0.020, microelement solution SL-10 - 1, KH <sub>2</sub> PO <sub>4</sub> - 0.36, K <sub>2</sub> HPO <sub>4</sub> - 2.34, pH 7.2
marine mineral medium (Marine salt medium)	KH <sub>2</sub> PO <sub>4</sub> - 0.5, NaH <sub>2</sub> PO <sub>4</sub> • H <sub>2</sub> O - 0.5, MgSO <sub>4</sub> • 7H <sub>2</sub> O - 0.5, NaCl - 4.0, NH <sub>4</sub> Cl - 0.5, microelement solution SL-10 - 1, pH 7.2
a modified Voroshilova-Dianova medium	K <sub>2</sub> HPO <sub>4</sub> - 1.0; KH <sub>2</sub> PO <sub>4</sub> - 1.0; NH <sub>4</sub> NO <sub>3</sub> - 1.0; MgSO <sub>4</sub> • 7H <sub>2</sub> O - 0.2; CaCl <sub>2</sub> • 2H <sub>2</sub> O - 0.02, a solution of trace elements SL-10, a solution of vitamins according to Volin, FeCl <sub>2</sub> -2 drops of a concentrated solution. Artificial seawater (g/l) was used as the basis for the medium: NaCl - 27.5, MgCl <sub>2</sub> - 5.0, MgSO <sub>4</sub> • 7H <sub>2</sub> O - 2.0, CaCl <sub>2</sub> - 0.5, KCl - 1.0, FeSO <sub>4</sub> - 0.001[30], pH 7.0-7.2.



**Figure S1.** Phylogenetic tree built on the basis of sequence analysis of 16S rRNA gene fragments of bacteria isolated from bottom sediments of the northern part of the Sea of Japan (gas-hydrate area). Scale bar length: 1 substitution per 100 nucleotides. Evolutionary distances were computed according to Maximum-likelihood phylogenetic tree based on 16S rRNA gene. The reliability of each branch was valuated by bootstrap analysis based on 1000 replications.



**Figure S2.** Phylogenetic tree built on the basis of sequence analysis of 16S rRNA gene fragments of bacteria isolated from bottom sediments of the northern part of the Sea of Japan (Non-gas-hydrate area). Scale bar length: 1 substitution per 100 nucleotides. Evolutionary distances were computed according to Maximum-likelihood phylogenetic tree based on 16S rRNA gene. The reliability of each branch was valuated by bootstrap analysis based on 1000 replications.

**Table S2.** Degree of biodegradation of hydrocarbons by strains isolated from bottom sediments of gas-hydrate area under aerobic and anaerobic conditions.

Strain code and NCBI number	The degree of biodegradation of hydrocarbons	
	under aerobic conditions	under anaerobic conditions
MZ573202 <i>Rhodococcus erythropolis</i> strain POI OP54-19/103-1(2)	78,71±1,94	
MZ540874 <i>Rhodococcus qingshengii</i> strain POI OP54-59/75-1	84,50±3,29	97,13±2,90
MZ540941 <i>Rhodococcus</i> sp. strain POI OP54-59/76-3	63,27±2,22	94,46±2,81
MZ543973 <i>Rhodococcus erythropolis</i> strain POI OP54-28/188-5	89,56±3,25	97,44±3,83
MZ577118 <i>Stenotrophomonas maltophilia</i> strain POI OP54-59/75-14	72,31±2,24	
MZ544365 <i>Stenotrophomonas maltophilia</i> strain POI OP54-42/37-3	63,51±1,95	
MZ573242 <i>Stenotrophomonas maltophilia</i> strain POI OP54-19/287-1	81,79±2,65	80,13±2,84
MZ573761 <i>Stenotrophomonas</i> sp. strain POI OP54-19/al 26-3	69,54±2,51	79,54±2,80
MZ562704 <i>Stenotrophomonas maltophilia</i> strain POI OP54-29/206-2(1)	51,98±1,77	84,74±2,43
MZ569721 <i>Stenotrophomonas maltophilia</i> strain POI OP54-35/18-2	97,38±3,83	100,00±3,77
MZ569678 <i>Stenotrophomonas maltophilia</i> strain POI OP54-41/36-1	84,02±2,61	94,64±3,33
MZ577121 <i>Psychrobacter cryohalolentis</i> strain POI OP54-44/45-2 (2)	75,71±2,78	
MZ573209 <i>Psychrobacter celer</i> strain POI OP54-35/18-2(1)	73,32±2,70	
MZ577169 <i>Psychrobacter maritimus</i> strain POI OP54-35/87-1	96,24±3,69	
MZ573198 <i>Psychrobacter nivimaris</i> strain POI OP54-44/45-2	78,18±2,69	92,30±3,60
MZ573229 <i>Psychrobacter</i> sp. strain POI OP54-41/35-10(1)	72,53±2,40	91,09±3,28
MZ540913 <i>Psychrobacter glacincola</i> strain POI OP54-59/68-4	79,98±2,85	98,73±3,10
MZ544190 <i>Psychrobacter nivimaris</i> strain POI OP54-44/45-3(1)	87,54±2,57	90,40±3,09
MZ577172 <i>Psychrobacter submarinus</i> strain POI OP54-28/101-1	86,89±3,40	97,28±3,33
MZ569847 <i>Pseudomonas brenneri</i> strain POI OP54-35/18-4	77,81±2,76	
MZ540778 <i>Pseudomonas brenneri</i> strain POI OP54-33/12-1(1)	76,84±2,54	93,17±2,92
MZ540892 <i>Pseudomonas</i> sp. strain POI OP54-33/al 91-18	83,83±3,20	96,77±3,25
MZ544024 <i>Pseudomonas</i> sp. strain POI OP54-35/213-8	56,07±1,71	89,33±2,58
MZ595665 <i>Pseudomonas</i> sp. strain POI OP54-59/73-1	77,12±2,89	84,89±2,51
MZ595782 <i>Pseudomonas</i> sp. strain POI OP54-59/92-1	70,53±2,16	97,22±3,62
MZ59600 <i>Pseudomonas</i> sp. strain POI OP54-51/92-1(3)	81,15±3,19	86,61±3,25
MZ577115 <i>Pseudomonas</i> sp. strain POI OP54-41/al 36-28	76,39±2,58	95,29±3,46
MZ569674 <i>Pseudomonas</i> sp. strain POI OP54-59/75-10	73,25±3,13	94,77±2,97
MZ573187 <i>Pseudomonas</i> sp. strain POI OP54-51/92-1(2)	67,37±2,40	85,01±2,96
MZ577178 <i>Pseudomonas yamanorum</i> strain POI OP54-33/12-1	68,01±1,09	84,91±2,13
MZ595666 <i>Bacillus</i> sp. (in: Bacteria) strain POI LV81-57/379-1(1)	71,18±2,30	92,00±2,90
MZ562706 <i>Bacillus</i> sp. (in: Bacteria) strain POI OP54-44/45-3(3)	87,82±3,45	89,23±2,72
MZ577136 <i>Bacillus</i> sp. (in: Bacteria) strain POI OP54-29/138-6	68,21±2,46	86,01±2,60
MZ569315 <i>Bacillus</i> sp. (in: Bacteria) strain POI OP54-28/198-6	88,22±2,72	80,95±2,41
MZ573241 <i>Nocardioides dokdonensis</i> strain POI OP54-35/al26-1	79,98±2,52	
MZ573232 <i>Nesterenkonia lutea</i> strain POI OP54-19/165	73,69±2,29	
MZ573186 <i>Peribacillus simplex</i> strain POI OP54-19/103-1(1)	87,64±3,45	92,40±2,66

**Table S3.** Degree of biodegradation of hydrocarbons by strains isolated from bottom sediments of non- gas-hydrate area under aerobic and anaerobic conditions.

Strain code and NCBI number	The degree of biodegradation of hydrocarbons	
	under aerobic conditions	under anaerobic conditions
MZ577132 <i>Psychrobacter pacificensis</i> strain POI LV81-57/al 379 (2)	80,21±2,75	
MZ577174 <i>Stenotrophomonas rhizophila</i> strain POI LV81-28/173-3(3)	86,33±2,67	
MZ573210 <i>Curtobacterium oceanosedimentum</i> strain POI LV81-23/149-3(2)	75,56±2,31	
MZ573212 <i>Pseudomonas brenneri</i> strain POI LV81-28/193-6(1)	70,33±2,72	92,35±3,27
MZ573237 <i>Pseudomonas sp.</i> strain POI LV81-3/25-1	53,07±2,08	76,62±2,22
MZ543976 <i>Bacillus sp. (in: Bacteria)</i> strain POI LV81-19/135-3	85,06±2,59	83,58±3,09
MZ544252 <i>Psychrobacter nivimaris</i> strain POI LV81-57/311	69,49±2,13	77,22±2,34
MZ573409 <i>Stenotrophomonas maltophilia</i> strain POI LV81-57/379-1	81,81±2,58	100,00±3,83
MZ562702 <i>Micrococcus sp.</i> strain POI LV81-57/al 379-1	78,77±2,76	72,34±2,56
MZ574136 <i>Psychrobacter glacincola</i> strain POI LV81-51/330-3(1)	77,34±2,89	88,26±3,35
MZ577107 <i>Pseudomonas sp.</i> strain POI LV81-57/379-1(3)	53,25±1,68	69,53±2,07
MZ569490 <i>Stenotrophomonas sp.</i> strain POI LV81-23/149-3(1)	72,56±2,59	88,03±2,57
MT758443 <i>Psychrobacter piscatorii</i> strain poi185	75,3±2,11	91,1±1,09
MZ577213 <i>Psychrobacter sp.</i> strain POI LV81-57/al 379 (1)	72,3±1,98	84,1±3,27
MZ562688 <i>Stenotrophomonas maltophilia</i> strain POI LV81-23/149-3	63,2±1,37	86,9±2,07
MZ577183 <i>Brevibacillus nitrificans</i> strain POI LV81-19/135	54,6±3,16	87,3±1,91
MT758444 <i>Robertmurraya kyonggiensis</i> strain poi87	75,1±2,09	94,2±2,01