

**Supplementary Table S1.** Summary of the closest affiliations of the representative isolates in the GenBank according to the 16S-rRNA gene sequences.

Isolate No.	Accession Number	Sequence Coverage	Closest NCBI match/Closest type strain	% Identity
PM03	KC683731	100	<i>Bacillus amyloliquefaciens</i> strain BVC18 (JQ660601)	100
PM05	HQ878378	100	<i>Microbacterium assamensis</i> (FR832516)	100
PM16	KC683732	100	<i>Pantoea stewartii</i> AIMSTTP2S ( JN835507)	93
PM219	KC683733	100	<i>Pantoea stewartii</i> strain S9-116 ( JQ660286)	99
PM221	KC683734	100	<i>Enterobacter cloacae</i> strain OVC28 (JQ660744)	100
PM223	KC683735	100	<i>Bacillus cereus</i> Kt7-14 (JF460754)	99
PM224	HQ878379	100	<i>Enterobacter cloacae</i> strain MSSRFS8 (HQ701802)	100
PM228	KC683736	100	<i>Bacillus</i> sp. ZW2531-1(EF567395)	99
PM232	HQ878380	100	<i>Enterobacter cloacae</i> strain MSSRFS8 (HQ701802)	100
PM251	KC683737	100	<i>Stenotrophomonas maltophilia</i> strain YPA 1-1 (JQ308611 )	100
PM253	KC683738	100	<i>Enterobacter cloacae</i> strain MSSRFS8 (HQ701802)	100
PM366	KC683739	100	<i>Bacillus nealsonii</i> strain KTNB19 ( FJ864726)	99
PM368	KC683740	100	<i>Stenotrophomonas maltophilia</i> strain EU84( JF681290)	100
PM376	KC683741	100	<i>Pseudomonas putida</i> strain DS8 ( JQ791183)	100
PM377	KC683742	100	<i>Pseudomonas</i> sp. AB_13 ( JQ033386)	97
PM379	KC683743	99	<i>Bacillus cereus</i> DF2C3 (DQ298085)	98
PM380	KC683744	100	<i>Bacillus subtilis</i> strain 1A337 (JQ746567)	99
PM381	KC683745	100	<i>Bacillus amyloliquefaciens</i> MH18B1 (JN558839)	99
PM386	KC683746	100	<i>Enterobacter cloacae</i> strain MSSRFS8 (HQ701802)	100
PM389	JF899310	100	<i>Pseudomonas aeruginosa</i> ASFP-38 (HQ018741)	97
PM4106	KC683747	99	<i>Enterobacter</i> sp. J54C13 (GQ901869)	96
PM4107	JF899311	100	<i>Pseudomonas aeruginosa</i> KZ3 (FJ695213)	99
PM5143	KC683748	100	<i>Pseudomonas fluorescens</i> A506 (CP003041)	100
PM5153	KC683749	100	<i>Pseudomonas fluorescens</i> A506 (CP003041)	99
PM7220	KC683750	100	<i>Microbacterium arborescens</i> strain S1-81 (JQ660028)	100
PM7265	JF899312	100	<i>Pseudomonas aeruginosa</i> strain BP C2 ( JQ866912)	100
PM8311	KC683751	100	<i>Ochrobactrum anthropi</i> strain BDUEBTVRMK33 (GU596495)	99
PM8335	KC683752	100	<i>Bacillus flexus</i> strain NIO24 ( JQ818411)	100
PM8337	KC683753	95	<i>Enterobacter</i> sp. Wy2-D9 ( JN986806)	98
PM8342	KC683754	100	<i>Enterobacter asburiae</i> strain M-T-MRS_23 (JQ795793)	100

PM9363	KC683755	99	<i>Stenotrophomonas maltophilia</i> strain YPA 1-1 (JQ308611)	99
PM9364	KC683756	100	<i>Stenotrophomonas maltophilia</i> ( HE664167)	99
PM9365	KC683757	100	<i>Arthrobacter woluwensis</i> ED (HM536961)	99
PM9366	KC683758	99	<i>Arthrobacter woluwensis</i> ED (HM536961)	99
PM9367	KC683759	100	<i>Bacillus cereus</i> Kt7-14 (JF460754)	99
PM9368	KC683760	99	<i>Rhodococcus kroppenstedtii</i> strain AHJ_3 (JN873342)	99
PM9369	KC683761	100	<i>Mycobacterium neoaurum</i> strain OS29 (JQ348111)	99
PM9370	KC683762	100	<i>Stenotrophomonas maltophilia</i> strain YPA 1-1(JQ308611 )	99
PM9376	KC683763	96	<i>Bacillus cereus</i> AIMST Musa3 (HQ694153)	99
PM9383	KC683764	100	<i>Enterobacter cloacae</i> strain RCT8 (HM805113)	100
PM9399	KC683765	97	<i>Bacillus stratosphericus</i> strain 290 (JQ860101)	99
PM9400	KC683766	100	<i>Microbacterium testaceum</i> StLB037 (AP012052)	99
PM9401	KC683767	100	<i>Arthrobacter woluwensis</i> ED (HM536961)	99
PM9402	KC683768	100	<i>Curtobacterium oceanosedimentum</i> strain S9-644 (JQ660303)	99
PM9403	KC683769	100	<i>Rhodococcus</i> sp. Eu-32 (DQ386111)	98
PM9404	KC683770	100	<i>Nocardioides</i> sp. DWM128 (HM854514)	99
PM9405	KC683771	99	<i>Acinetobacter calcoaceticus</i> (AY800383)	99
PM9406	KC683772	100	<i>Stenotrophomonas maltophilia</i> (HE664167)	100
PM9408	KC683773	100	<i>Pseudomonas oryzae</i> (JQ388741)	100
PM9409	KC683774	100	<i>Rhodococcus qingshengii</i> strain HWG-A33 (JQ684256)	99
PM9410	KC683775	100	<i>Nocardioides</i> sp. S2-186 ( JQ660079)	99
PM9411	KC683776	100	<i>Stenotrophomonas maltophilia</i> strain C-J-R2A1	99
PM9424	KC683777	100	<i>Arthrobacter woluwensis</i> ED (HM536961)	100
PM9426	KC683778	100	<i>Arthrobacter woluwensis</i> MS4_481 (HM032831)	99
PM9427	KC683779	100	<i>Arthrobacter woluwensis</i> ED (HM536961)	99
PM9429	KC683780	97	<i>Pantoea stewartii</i> strain S9-116 (JQ660286)	99
PM9430	KC683781	100	<i>Bacillus cereus</i> Kt7-14 (JF460754)	100
PM9432	KC683782	92	<i>Bacillus cereus</i> strain H9B-3 (HQ238861)	99
PM9433	KC683783	98	<i>Pantoea stewartii</i> AIMST Nmie4 (JF819695)	99
PM9435	KC683784	99	<i>Pantoea stewartii</i> isolate 40G RT2A (EF189919)	100
PM9436	KC683785	100	<i>Microbacterium testaceum</i> StLB037 (AP012052)	100
PM9437	KC683786	100	<i>Agrobacterium larrymoorei</i> strain S3-89 (JQ660107)	99
PM9438	KC683787	100	<i>Microbacterium testaceum</i> StLB037 (AP012052)	98
PM10455	KC683788	100	<i>Rhizobium</i> sp. HPCkC-Ca(i) (HE802763)	100
PM10458	KC683789	100	<i>Bacillus pumilus</i> FS55 (AF260751)	99
PM10459	KC683790	99	<i>Ochrobactrum</i> sp. MVS1(JN089705)	97
PM10461	KC683791	100	<i>Rhizobium</i> sp. HPCkC-Ca(i) ( HE802763)	100
PM10462	KC683792	100	<i>Microbacterium testaceum</i> StLB037 (AP012052)	99
PM10463	KC683793	100	<i>Acinetobacter</i> sp. XJ104-YF-7(JQ726509)	98

PM10476	KC683794	100	<i>Ochrobactrum intermedium</i> strain TND41(JQ660549)	100
PM10478	KC683795	100	<i>Ochrobactrum intermedium</i> strain TND41(JQ660549)	99
PM10490	JF899314	99	<i>Pseudomonas aeruginosa</i> NQ8 (EU939706)	97
PM10491	JF899315	100	<i>Pseudomonas aeruginosa</i> PB28 (JN408334)	100
PM10503	KC683796	92	<i>Bacillus amyloliquefaciens</i> Kk3-3 (JF460724)	99
PM10509	KC683797	100	<i>Pseudomonas fluorescens</i> strain hswx163 (JQ236807)	100
PM10510	KC683798	100	<i>Bacillus pumilus</i> strain Y24 (JQ798393)	100