## Isolation of *Burkholderia koreensis* sp. nov. at the Field of Ginseng

Ho-Bin Kim, Min-Ju Park, Hwa Son, Byung-Man Kong, Deok-Chun Yang\*
Dept. of Oriental Medicinal Material & Processing, College of Life Science,
Kyung Hee University, Yongin 449-701, Korea

\*E-mail. dcyang@khu.ac.kr

Gram-negative, slightly curved rod-shaped bacterium DCY02<sup>T</sup> was isolated at the ginseng field in Daejeon, South Korea. On the basis of 16S rRNA gene sequence similarity, the strain DCY02<sup>T</sup> was shown to belong to the family *Burkholderiaceae* of the β-Proteobacteria related to *Burkholderia terricola* LMG 20594<sup>T</sup> (97.8%), *Burkholderia caledonica* LMG 19076<sup>T</sup> (97.7%), *Burkholderia xenovorans* LMG 21463<sup>T</sup> (97.7%) and *Burkholderia fungorum* LMG 16225<sup>T</sup> (97.4%). Phylogenetic distance from any validly described species within the genus *Burkholderia* was less than 97%. Chemotaxonomic data (major ubiquinone - Q-8; major fatty acids - Summed feature 7 (C18:1 ω7c/ω9t/ω12t), Summed feature 4 (C16:1 ω7c/C15:0 iso 2OH), C16:0 and Summed feature 3 (C14:0 3OH/C16:1 iso) supported the affiliation of the strains to the genus *Burkholderia*. The results of DNA-DNA hybridization, physiological and biochemical tests allowed genotypic and phenotypic differentiation of the strain from the validly published *Burkholderia koreensis* sp. nov. is proposed, with the type strain DCY02<sup>T</sup> (=KCTC 12389<sup>T</sup> =NBRC 100965<sup>T</sup>). The NCBI GenBank accession number for 16S rRNA gene sequence of strain DCY02<sup>T</sup> is AB201286.