Confirming of Origin of Magnoliae Flos on the DNA analysis method and Identification

Jong-Hun Noh, Sun Hua, Baigalmaa Jigden, Deok-Chun Yang*

Korean Ginseng Center for Most Valuable product and Ginseng Genetic Resource Bank, Kyung Hee University, Yongin 446-701, Korea. *Corresponding author. Tel: 031-201-2100, E-mail: dcyang@khu.ac.kr

Most species in the genus Magnolia have been used for oriental medicinal purposes in the name of "Flos Magnoliae(Sin-ii)". Among the "Magnolia" species, five species, Manolia denudata, Manolia biondii, Manolia liliiflora, Manolia sprengeri, Manolia kobus were most used as publicly certified medicinal material in Korea, China. Phylogenetic analysis, using the neighbor joining method of 22 species of the "Magnolia" genus and its allies based on tRNA coding leucine/phenylalanie (trnL-F) sequences and NADH dehydrogenase subunit F (ndhF) sequence in chloroplast DNA were presented in this study. For the identification of "Magnolia" species, polymerase chain reaction (PCR) analysis of chloroplast DNA regions such as trnL-F and ndhF have proven an appropriate method. A single nucleotide polymorphism (SNP) has been identified between genuine "Sin-ii" or their fraudulent and misuse. Specific PCR primers were designed from this polymorphic site within the sequence data, and were used to detect true plants via multiplex PCR.