

Luteolin *O*-methyltransferase from Soybean

Lee Hyo Jung, Kim Bong Gyu, Ahn Joong-Hoon*

Bio/Molecular Informatics Center, Department of Molecular Biotechnology,
KonKuk University, Seoul

TEL: +82-2-450-3764, FAX: +82-2-3437-6106

Abstract

O-methyltransferase (OMTs) catalyze the transfer of a methyl group from S-adenosine-L-methionine to a hydroxyl group of an acceptor molecule to form methyl ether derivatives and modify basic backbone of secondary metabolites. OMTs could be classified as two groups depending on molecular weight. The first group is 23,000 to 27,000 Da and is represented by the caffeoyl coenzyme A OMTs (CCoAOMTs) which have been known to be involved in the biosynthesis of monolignol. The other is about 38,000 to 43,000 and use caffeic acid, flavonoids, sugars, and other scent compounds. A new O-methyltransferase, SOMT-9 was cloned from the tissues of *Glycine max* and it turned out to encode a protein whose molecular weight is 27-kDa. SOMT-9 was expressed as a GST-fusion protein in *E. coli* and several substrates such as caffeic acid, narigenin, kampferarol, quercetin, and luteolin were tested. HPLC analysis of reaction product showed that SOMT-9 used flavonoids containing ortho hydroxyl groups such as luteolin and quercetin. The NMR result showed that SOMT-9 transfer a methyl group to 3'-OH group of lueolin, which was determined to be the best substrate.

References

1. Ibdah, M., X.-H. Zhang, J. Schnot, and Vogt, T., 2003. A novel Mg²⁺-dependent O-methyltransferase in the phenylpropanoid metabolism of *Mesembryanthemum crystallinum*. *J. Biol. Chem.* 278: 43961-43972.
2. Yoon, Y.; Yi, Y. S.; Lee, Y.; Kim, S.; Kim, B. G.; Ahn, J. _H.; Lim, Y. Characterization of O-methyltransferase ScOMT1 cloned from *Streptomyces coelicolor* A3(2), *Biochim. Biophys. Acta*, 2005, 1730, 85 - 95.