

BIOCHEMICAL CHARACTERIZATION OF MORE AXILLARY GROWTH1 IN STRIGOLACTONE BIOSYNTHESIS

Takahito Nomura¹, Kaori Yoneyama¹, Kohki Akiyama², Xiaonan Xie¹, Toshiyuki Ohnishi³, Shinjiro Yamaguchi⁴, and Koichi Yoneyama¹

¹*Utsunomiya University*, ²*Osaka Prefecture University*, ³*Shizuoka University*, ⁴*Tohoku University*

Strigolactones (SLs) function as host recognition signals for root parasitic plants and symbiotic arbuscular mycorrhizal fungi in the rhizosphere and as plant hormones regulating shoot and root architecture in plants. Carotenoid isomerase D27, carotenoid cleavage dioxygenases CCD7 and CCD8, and cytochrome P450 monooxygenase MORE AXILLARY GROWTH1 (MAX1) were identified as SL biosynthesis enzymes by genetic screening of shoot branching mutants. D27, CCD7 and CCD8 convert β -carotene to carlactone (CL), an SL precursor having no canonical four-ring structure of SLs, by their sequential reactions. We have reported that *Arabidopsis* MAX1 catalyzes oxidation of the C-19 methyl group of CL to carboxylic acid, affording carlactonoic acid (CLA), while one of rice MAX1 homologs was reported to catalyze the conversion of CL to 4-deoxyorobanchol having the four-ring structure. In order to know which is the common reaction in MAX1 homologs is, we examined the enzymatic functions of MAX1 homologs in *Arabidopsis*, rice, maize and tomato using a yeast expression system. As a result, the conversion of CL to CLA was found to be a common function in MAX1 homologs but not that of CL to 4DO.