

Metabolic engineering of *Arabidopsis thaliana* to increase carotenoid content by using the *Erwinia herbicola crtI* and *Escherichia coli dxs* genes

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In carotenoid biosynthesis, phytoene desaturase gene *crtI* and DXP synthase gene *dxs* are both rate-limiting step enzymes. These two genes were cloned from *Erwinia herbicola* and *Escherichia coli* chromosomal DNA, respectively. The genes were functionally expressed in lycopene producing *E. coli* and increased the carotenoid production. *Arabidopsis thaliana* was transformed with the genes which were expressed under the control of CaMV 35S promoter. Transit peptide sequence of *Brassica rapa* Rubisco small subunit was located in front of the genes to target the expressed enzymes to plastid in which carotenoids were synthesized. Recognition of these genes and expression level were demonstrated by PCR and southern blotting, and Northern blotting, respectively. Transgenic plants, which were transformed with *E. herbicola crtI*, showed resistance against bleaching herbicide norflurazon, which specifically inhibited phytoene desaturase encoded by native genes of *Arabidopsis thaliana*. Also, *crtI* and *dxs* transgenic plants showed higher amount of carotenoid contents than control transgenic plants.

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