

UNRAVELING THE REGULATION OF FLAVONOID PRODUCTION IN GLOBE ARTICHOKE: ISOLATION AND CHARACTERIZATION OF THE FLAVONOL REGULATOR *CCMYB12*

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Flavonoids, a well-studied group of secondary metabolites, are known to exhibit health promoting effects such as antioxidant capacities, anti-cancer and anti-inflammatory activity. The globe artichoke [*Cynara cardunculus* var. *scolymus* (L.) Fiori] is a natural functional food rich in bioactive phenolic compounds, including flavonoids, and in fibers (inulin) and minerals. To study the regulation of flavonoid biosynthesis, a R2R3-MYB transcription factor, *CcMYB12*, was isolated from artichoke leaves.

Phylogenetic analysis showed that this protein belongs to the MYB subgroup 7 (flavonol-specific MYB), which includes Arabidopsis AtMYB12, grapevine VvMYBF1, and tomato SIMYB12. *CcMYB12* transcripts were detected specifically in artichoke immature inflorescence and young leaves. Electrophoretic mobility shift assays (EMSAs) revealed that recombinant CcMYB12 protein is able to bind to canonical AC element, a DNA binding site ubiquitously present in the promoters of genes encoding flavonol biosynthetic enzymes. Arabidopsis and tobacco plants overexpressing *CcMYB12* protein were generated and characterized. Finally, by targeted metabolic and molecular analyses, we show that the ectopic expression of *CcMYB12* in Arabidopsis and tobacco plants promotes flavonol biosynthesis at expenses of anthocyanins.