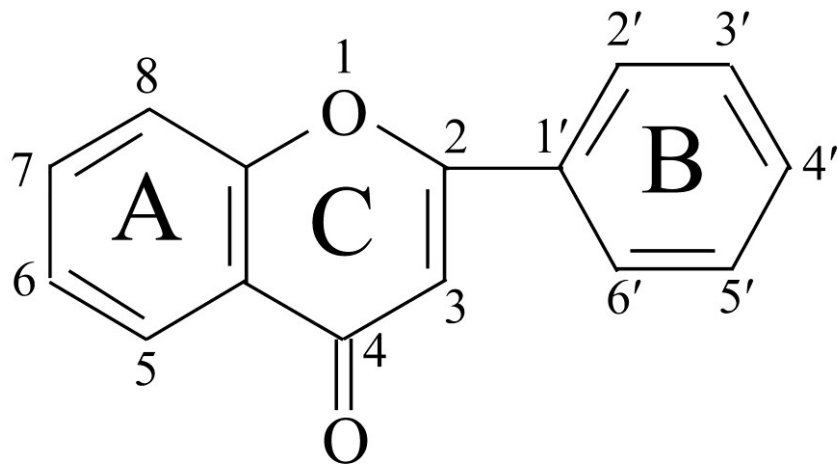


Quantitative Phosphoproteomic and Metabonomic Analyses Reveal GmMYB173

Optimizes Flavonoid Metabolism in Soybean to Survive Salt Stress

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Supplementary Figure S1. General structure of flavonoids.

113	114	115	116	117	118	119	121
C1	C2	C3	C4	T1	T2	T3	T4

Supplementary Figure S2. Sample set of quantitative phosphoproteomic analysis.

One eight-plex iTRAQ set was used for analyzing the four biological replications of control (C1~C4) and NaCl treated (T1~T4) groups of soybean roots.