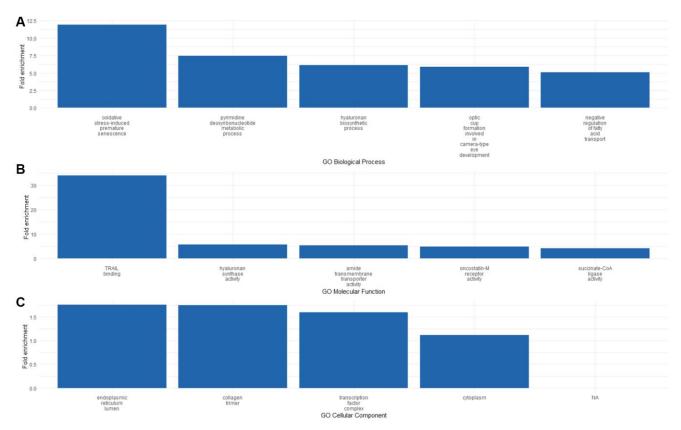
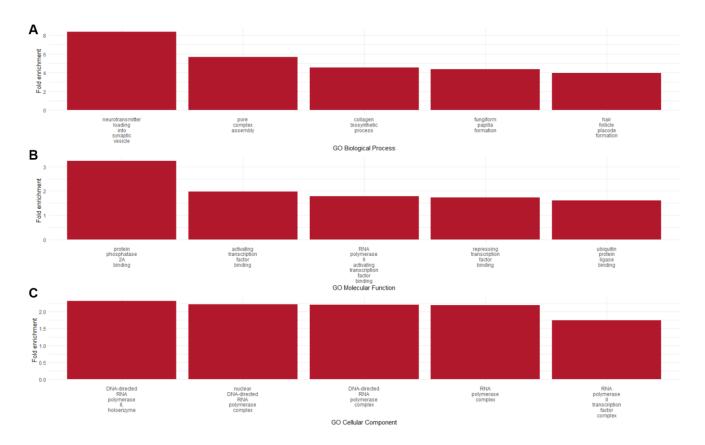
SUPPLEMENTARY FIGURES



Supplementary Figure 1. Gene Ontology (GO) terms enriched for the hypermethylated CpG sites after supplementation. The Top 5 most enriched GO terms for Biological Processes (A), Molecular Function (B), and Cellular Component (C) were included. A nominal *p*-value <0.001 was used to assign significance.



Supplementary Figure 2. Gene Ontology (GO) terms enriched for the hypomethylated CpG sites after supplementation. The Top 5 most enriched GO terms for Biological Processes (A), Molecular Function (B), and Cellular Components (C) were included. A nominal *p*-value <0.001 was used to assign significance.