

Supplementary information, Figure S1

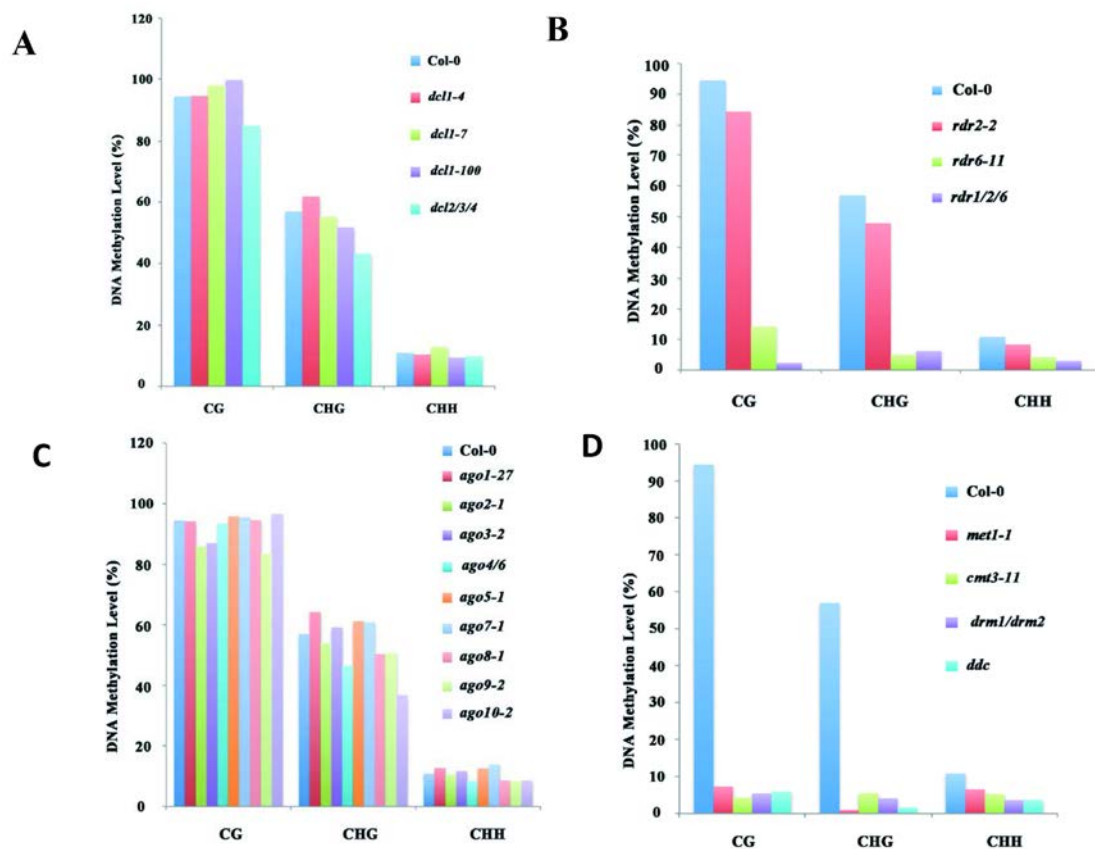


Figure S1 DNA methylation levels at the 5' UTR region of *HEI10*. Sodium bisulfite sequencing was performed using *HEI10*-specific primers in plants of the indicated genotypes: **(A)** *dcl* mutants; **(B)** *rdr* mutants; **(C)** *ago* mutants; **(D)** DNA methyltransferase mutants. At least 15 colonies were sequenced for each sample.