

**Figure S1**

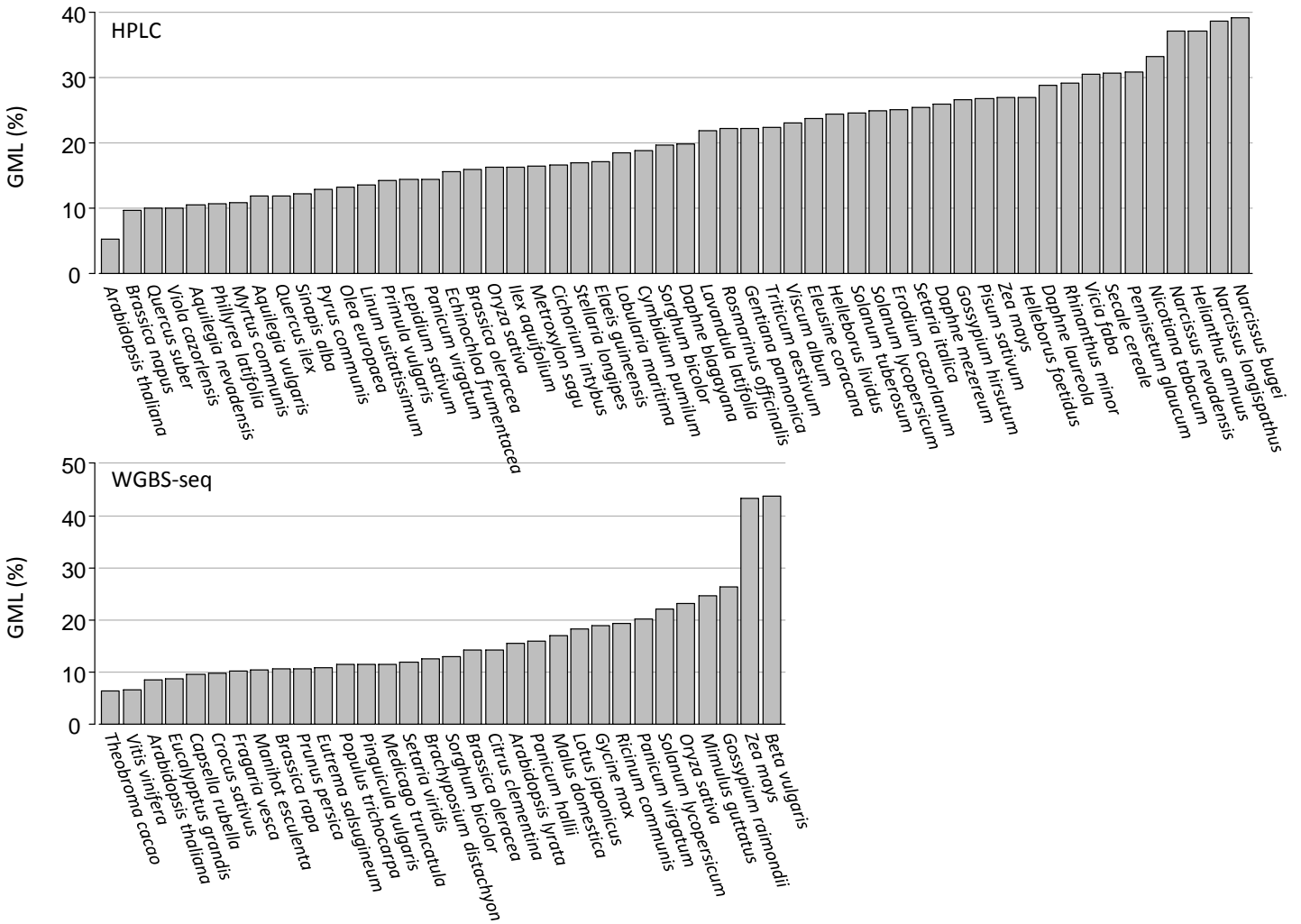


Figure S1: GMLs of plant species whose methylomes have been analyzed by whole genome bisulfite sequencing (WGBS-seq) or high-performance liquid chromatography (HPLC)

**Figure S2**

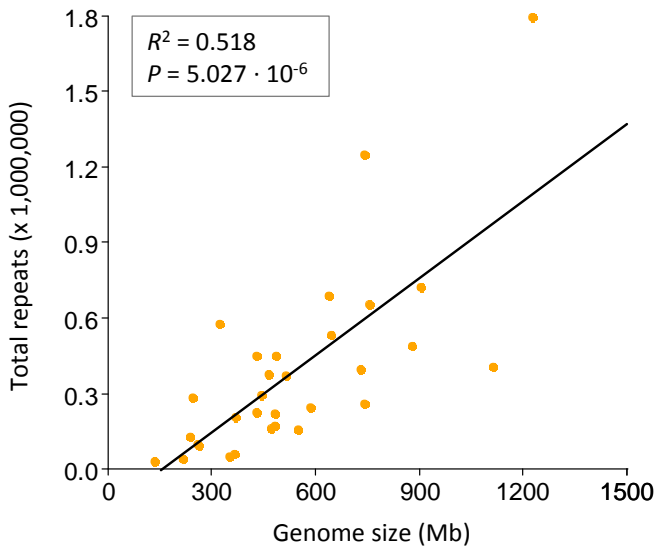


Figure S2: Correlation between genome size and total number of repeats in the genome, using the taxa presented in S1