

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Teosinte, landrace, and modern maize samples used in the study.

File Name: Supplementary Data 2

Description: Population-wide DMRs in the CG and CHG contexts.

File Name: Supplementary Data 3

Description: Selective sweeps detected between populations.

File Name: Supplementary Data 4

Description: Linkage disequilibrium (LD) analysis between DMR and local SNPs.

File Name: Supplementary Data 5

Description: The list of genes with CG teosinte-maize DMRs located at the exonic regions.

File Name: Supplementary Data 6

Description: The list of genes exhibiting interactive loops between genes and hypomethylated DMRs in maize located at the intergenic regions.

File Name: Supplementary Data 7

Description: Flowering time candidate genes located at sweep DMRs and interacting DMRs.