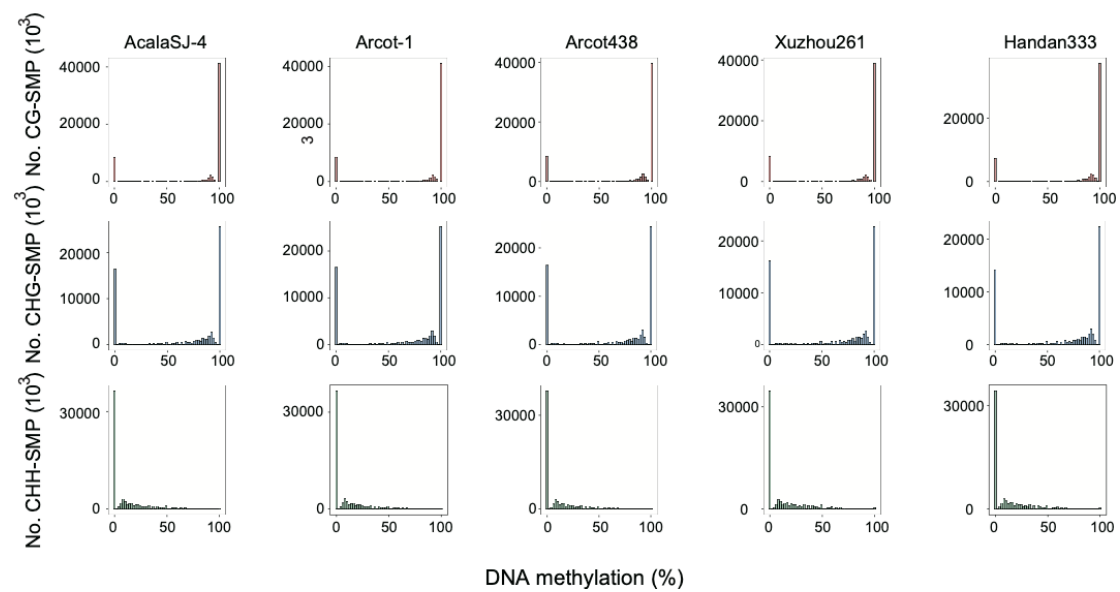




Supplementary information, Fig. S3. Bar plot showed the distribution of methylated levels for CG, CHG, and CHH loci in five randomly sampled accessions. The accessions, named acalaSJ-4, Arcot-1, Arcot438, Xuzhou261, and Handan333, are arranged from left to right. The top to bottom order indicates the distribution of DNA methylation levels of CG, CHG, and CHH, respectively.