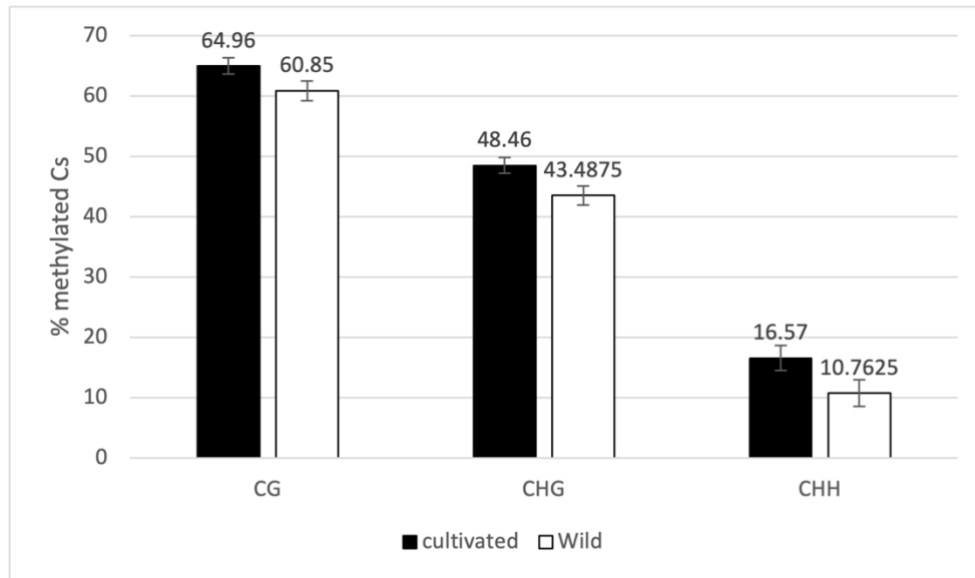




Supplementary Figure 1: Analysis of differences in global levels of DNA methylation in intergenic regions of cultivated and wild *V. vinifera* accessions. Bars show the average percentage of methylation per sequence context (CG, CHG, and CHH) in cultivated (*V. vinifera* ssp. *vinifera* (n = 10); black bars), and wild type (*V. vinifera* ssp. *sylvestris* (n = 8); white bars) accessions. Error bars indicate the calculated Standard Deviation. ** T-test, p-value < 0.01.