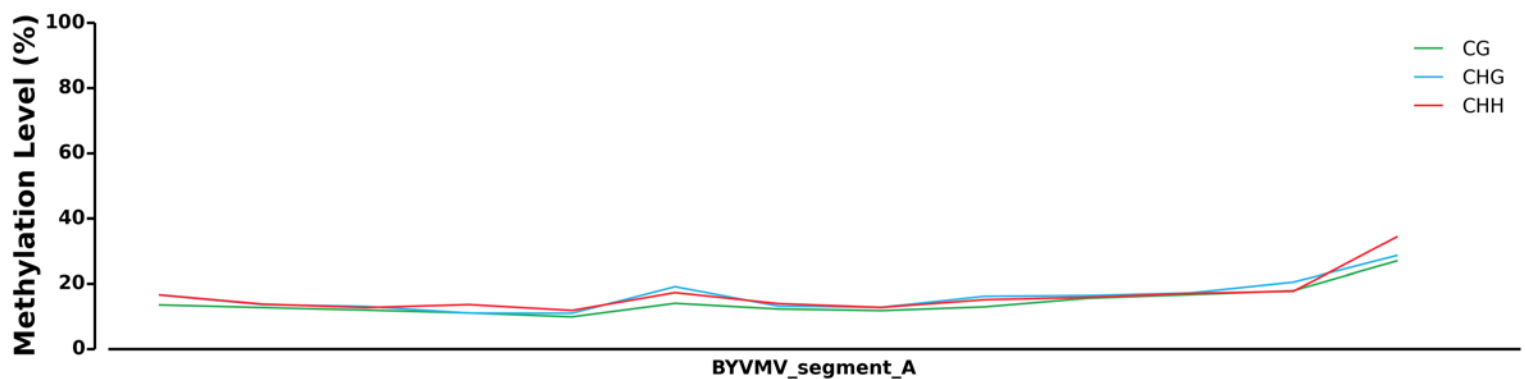
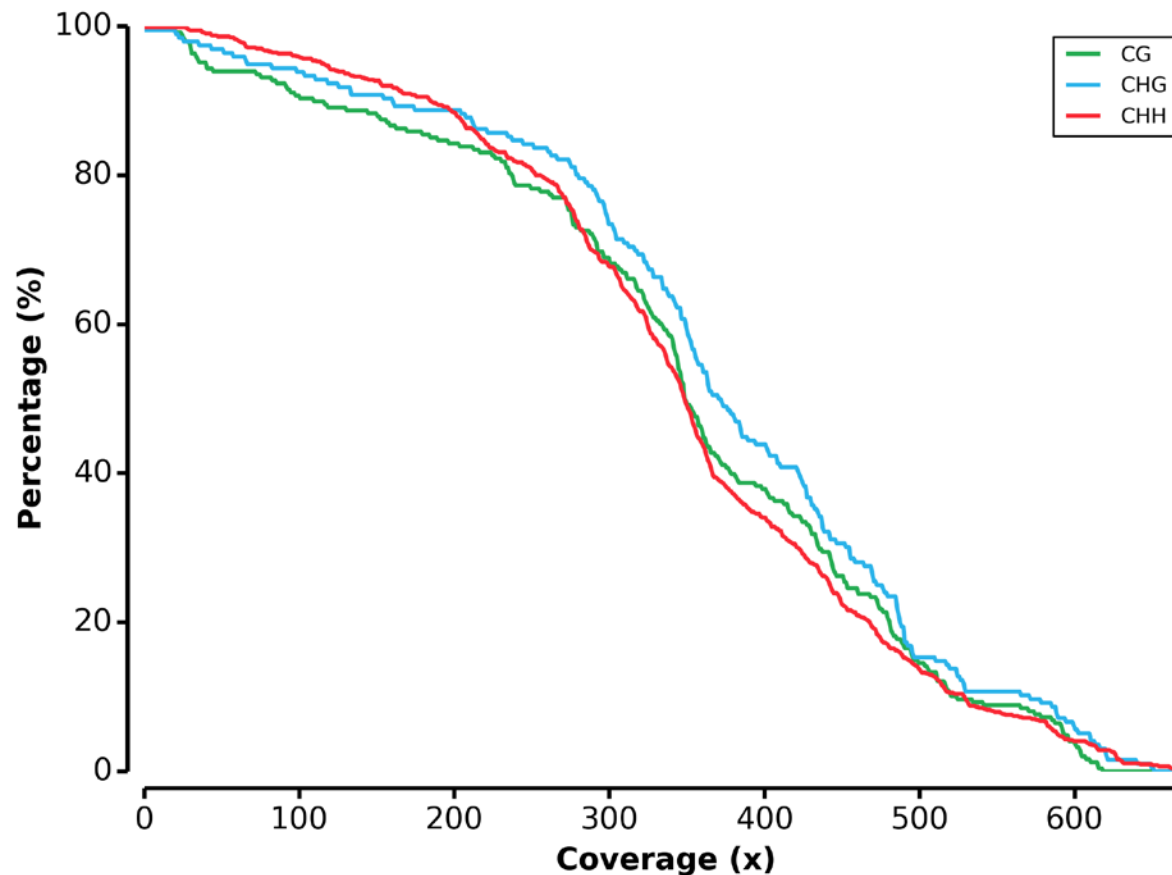


MethGo analysis of Bisulfite NGS Data for C2 mutant APTR (C2MAPTR) alone infiltrated plant samples



Supplementary Figure S5. MethGo analysis of Bisulfite NGS Data for C2 mutant APTR (C2MAPTR) alone infiltrated plant samples: A) Coverage map of ~2.8 million bisulfite NGS reads from C2MAPTR alone infiltrated plant sample. Y-axis represents the percentage of methylation sites covered in the **BYVMV** C2 mutant. X- axis represents the depth of coverage (X) in multiples of thousand. B) The genome-wide methylation map of BYVMV showing the level of methylation at different regions of C2 mutant **BYVMV**.