Figure S2. Distribution of the ability to suppress CMS phenotype across sunflower samples with different allelic states for 8 statistically significant markers: A)S13_170494693, B)S13_171053833, C)S13_173268042, D)S13_173832391, E)S13_174474103, F)S13_174474122, G)S13_174809087, H)S13_178217103