



Figure S6 | Distribution of allele frequencies and effects for HopAM1-induced responses. Local cell death response (A) and HopAM1-induced meristem chlorosis response (B). Effect size represents standardized regression coefficients or “beta”. They are interpreted as the phenotypic difference between genotypes carrying the minor allele-frequency allele from those with the reference allele and are expressed in standard deviations of the phenotype. Each dot represents a tested SNP. The top 500 SNPs are in blue.