Figure S1 Bayes Factors (BFs) describing the deviation of the genotype-environment associations of 266 putative candidate SNPs from a null demographic model (i.e. a co-variance matrix built with 1,745 control SNPs) obtained from BAYENV 2.0. Red dots are those SNPs above the BF=10 threshold (horizontal discontinuous line), and which were assumed to be significantly associated with climate.