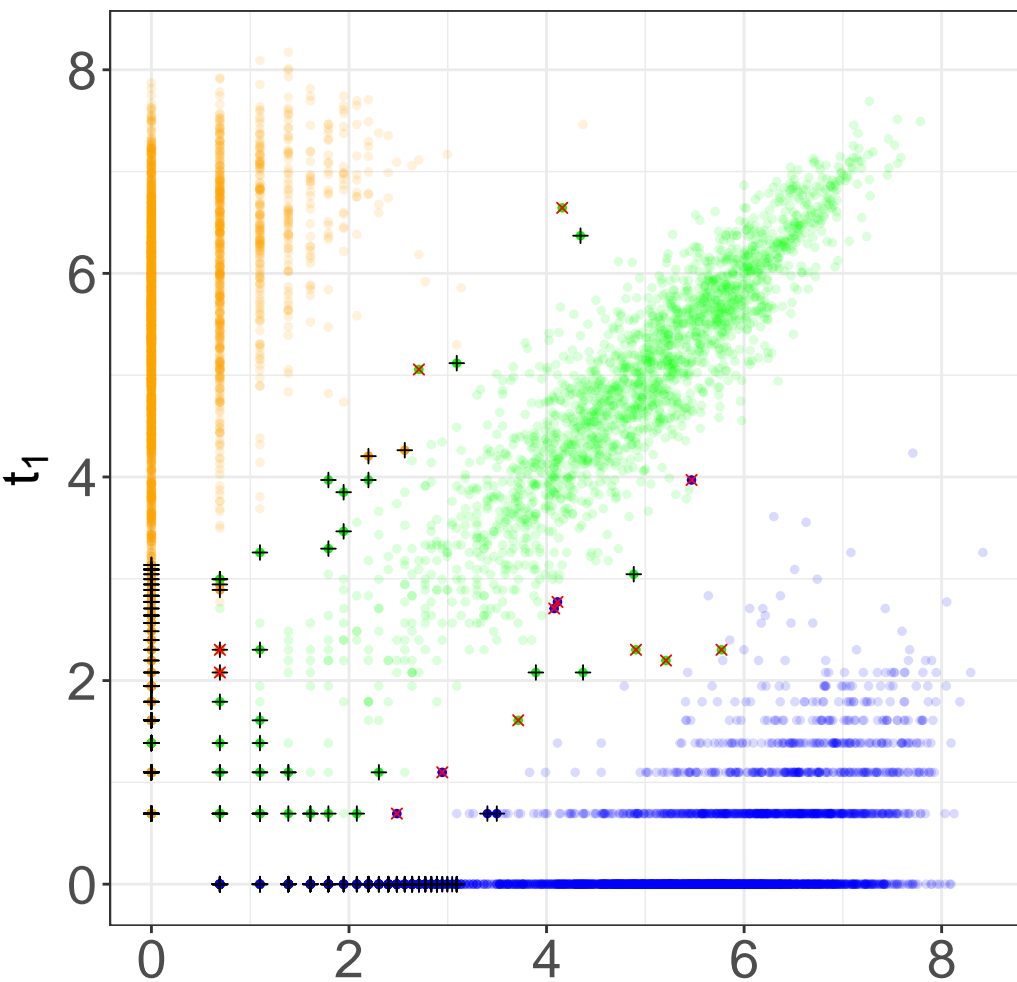
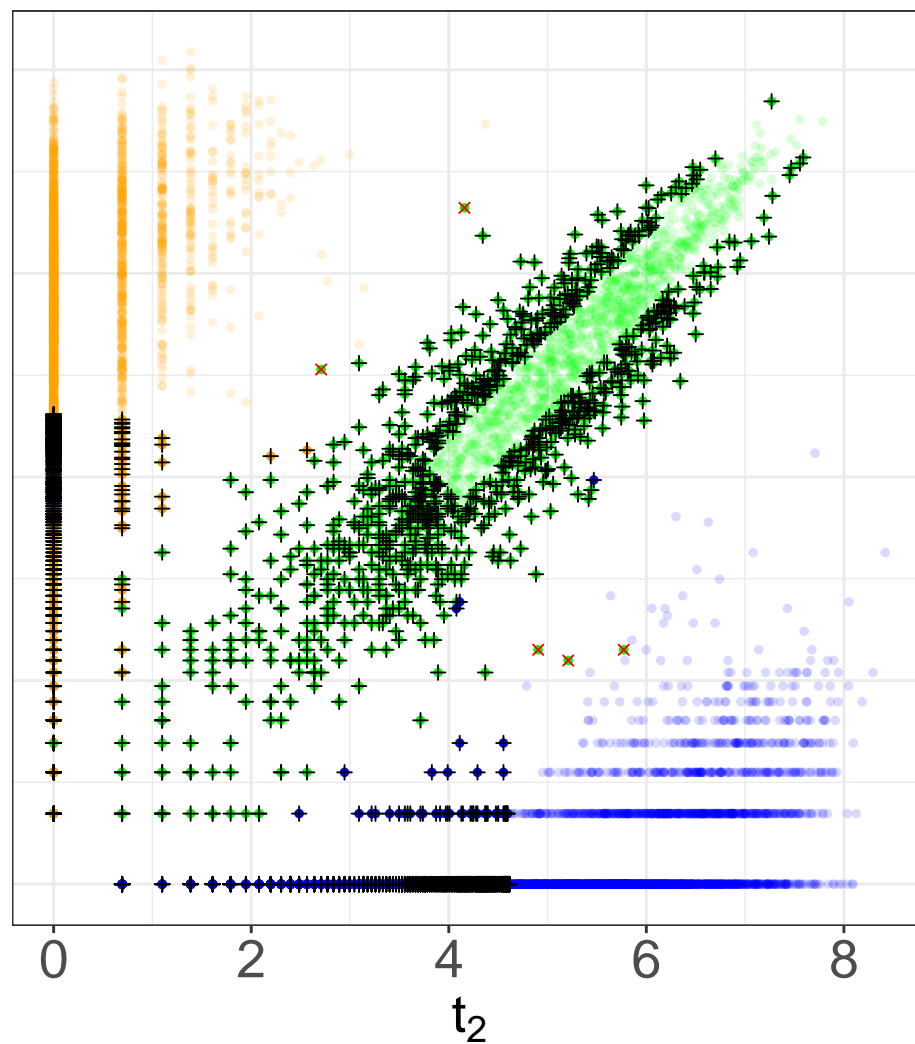


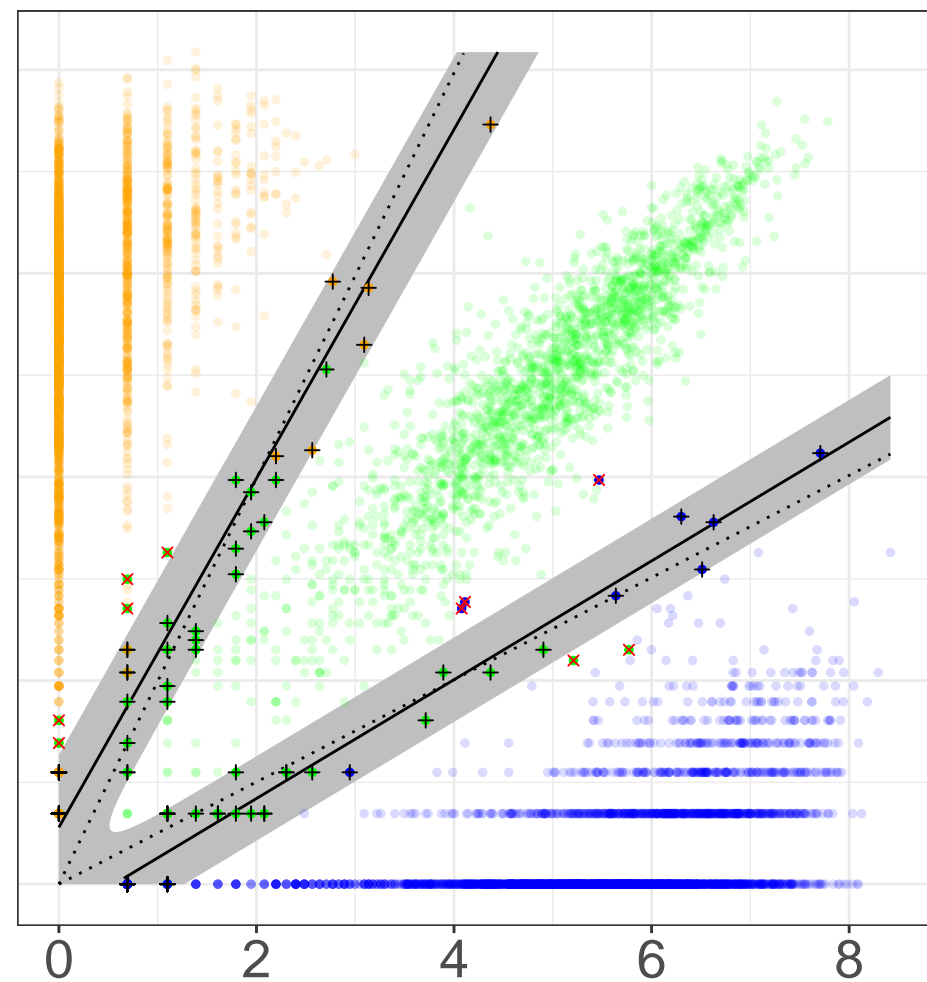
HID_SNP



UCPH



SMLR



Threshold

■ $q = 0.9$

Missed

+ No-call

x Wrong

Method

— $\beta_0 \neq 0$ $\beta_0 = 0$

True genotype

● a_1a_1 ● a_1a_2 ● a_2a_2