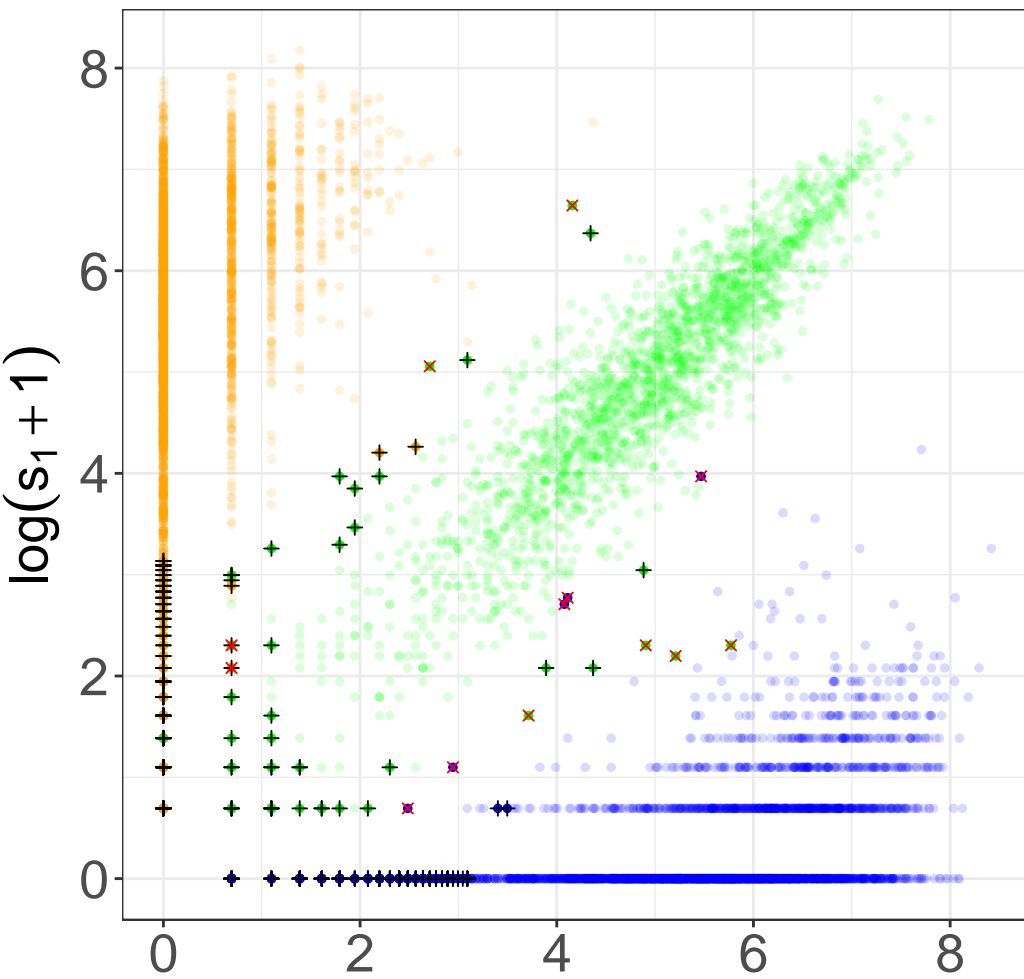
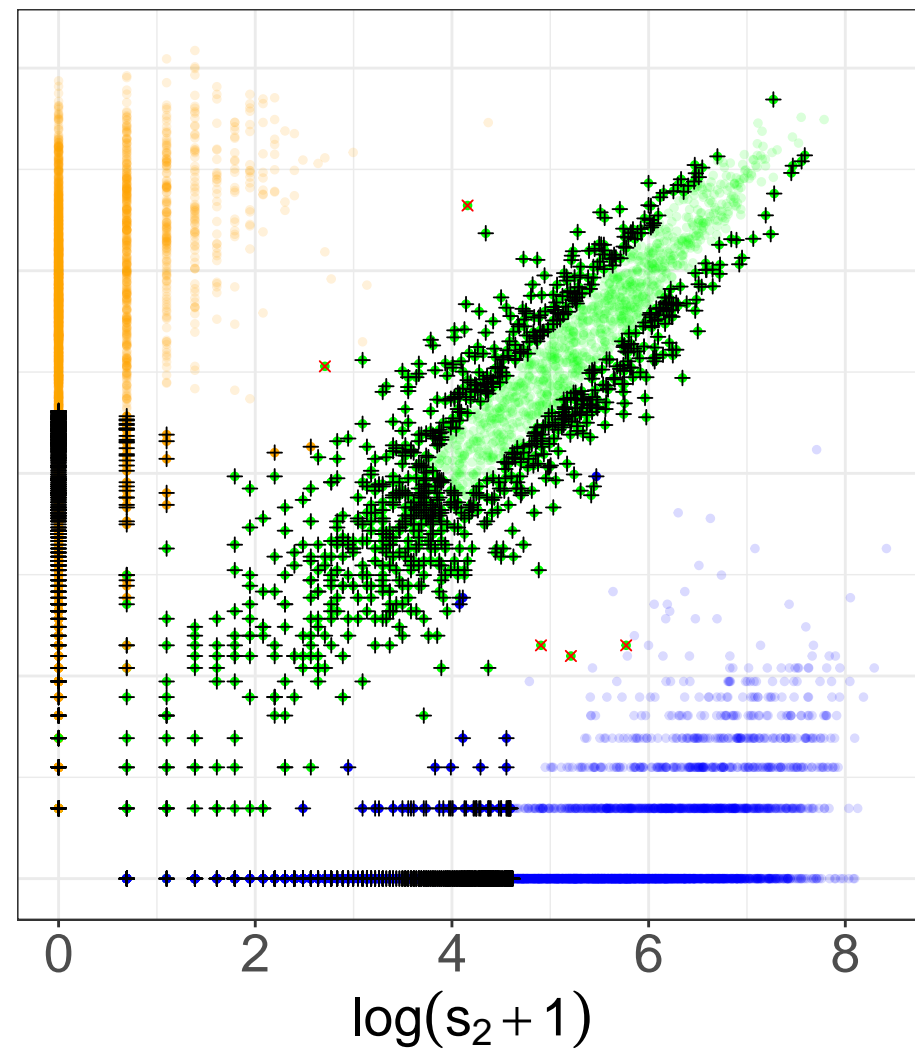


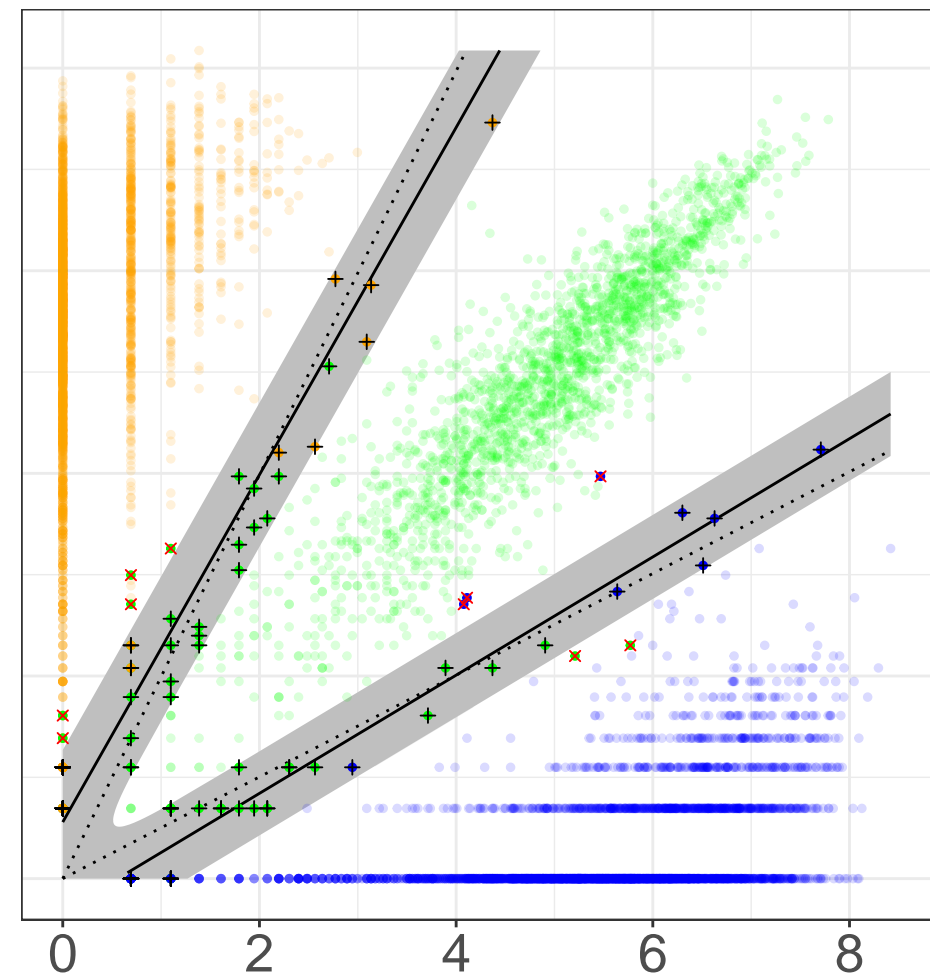
HID_SNP



UCPH



SMLR



Threshold

 $q = 0.9$

Missed

+ No-call

× Wrong

Method

— $\beta_0 \neq 0$ $\beta_0 = 0$

True genotype

• a_1a_1 • a_1a_2 • a_2a_2