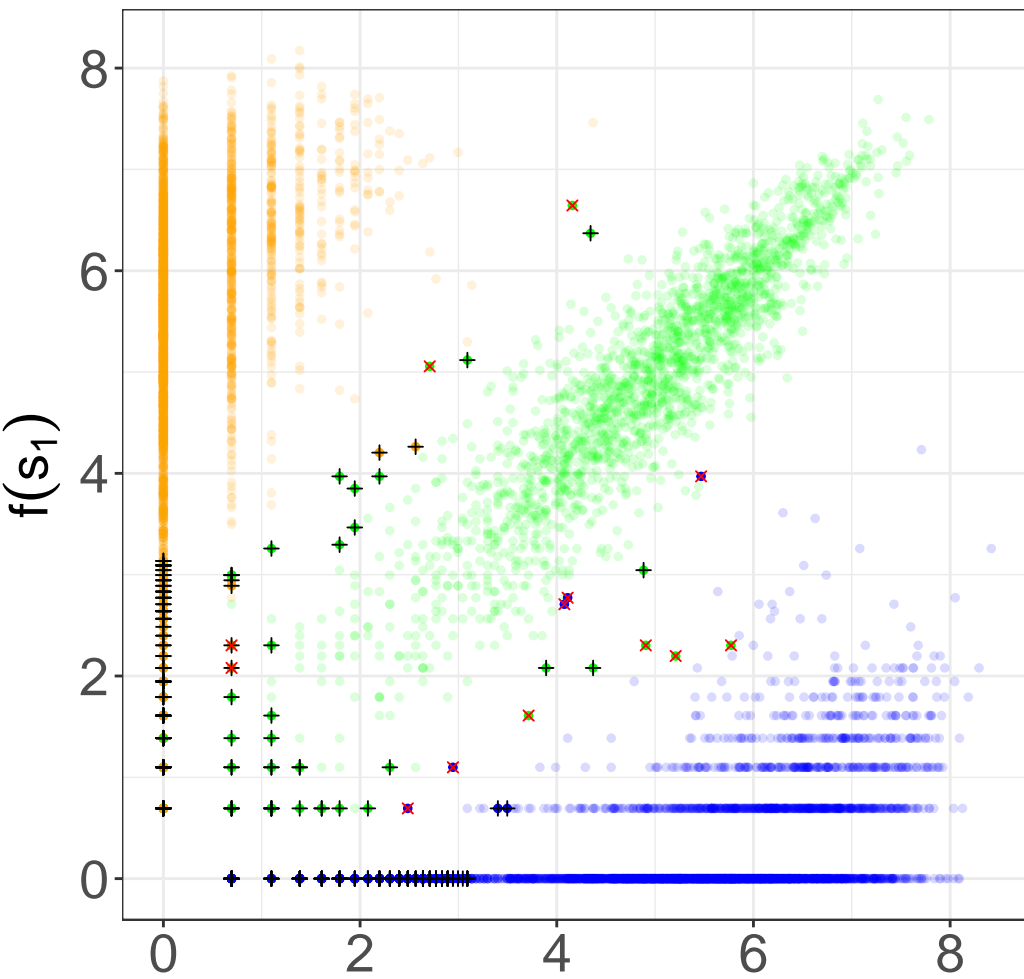
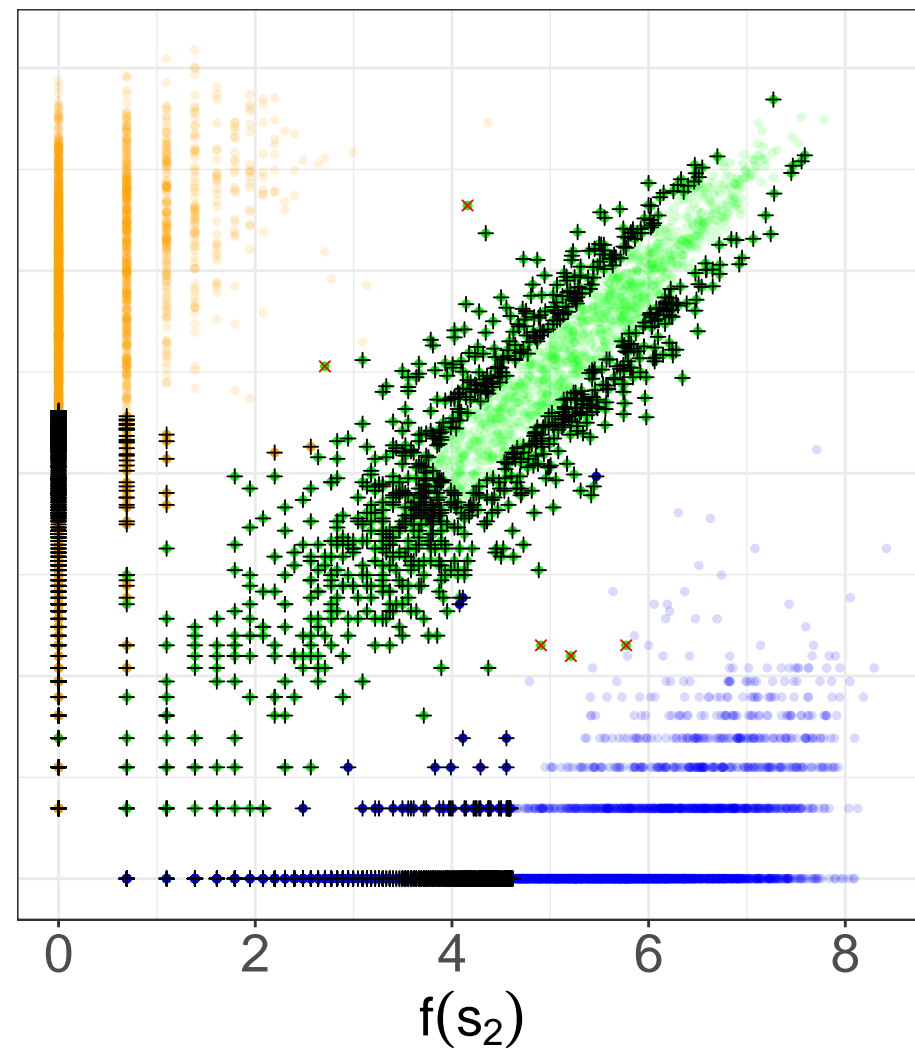


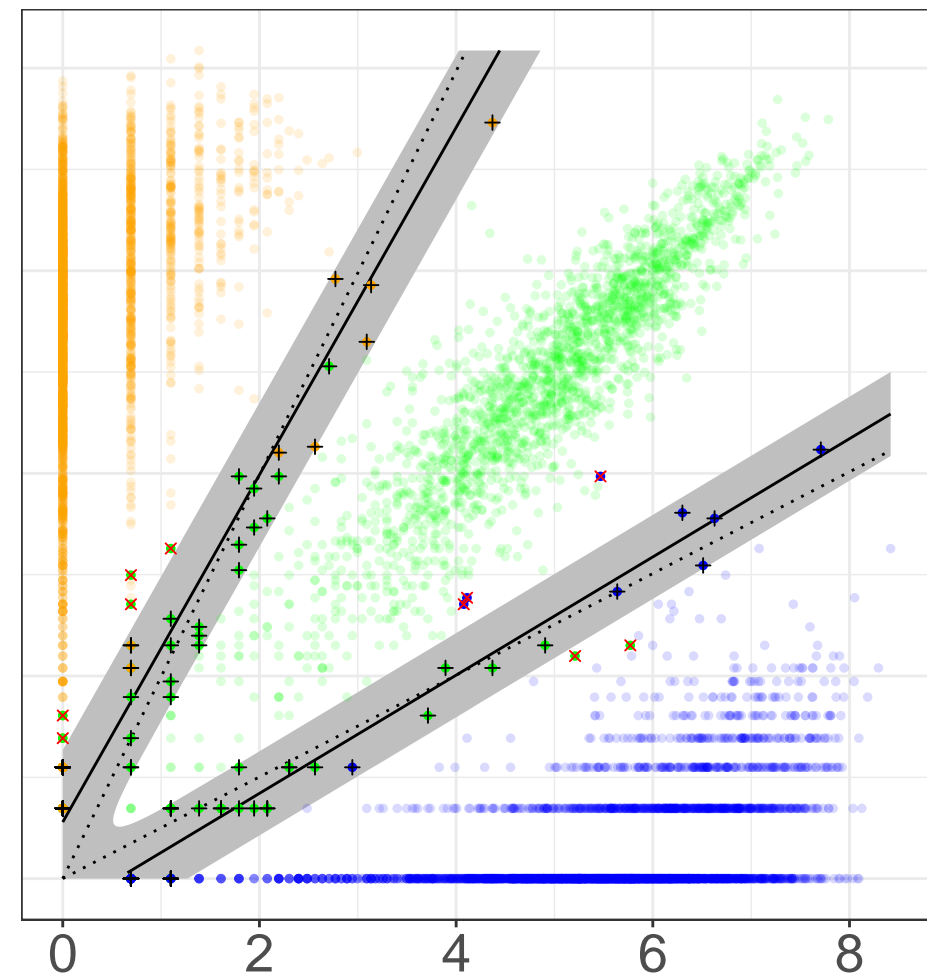
HID\_SNP



UCPH



SMLR



Threshold

 $q = 0.9$ 

Missed

+ No-call

x Wrong

Method

—  $\beta_0 \neq 0$ ....  $\beta_0 = 0$ 

True genotype

•  $a_1a_1$ •  $a_1a_2$ •  $a_2a_2$