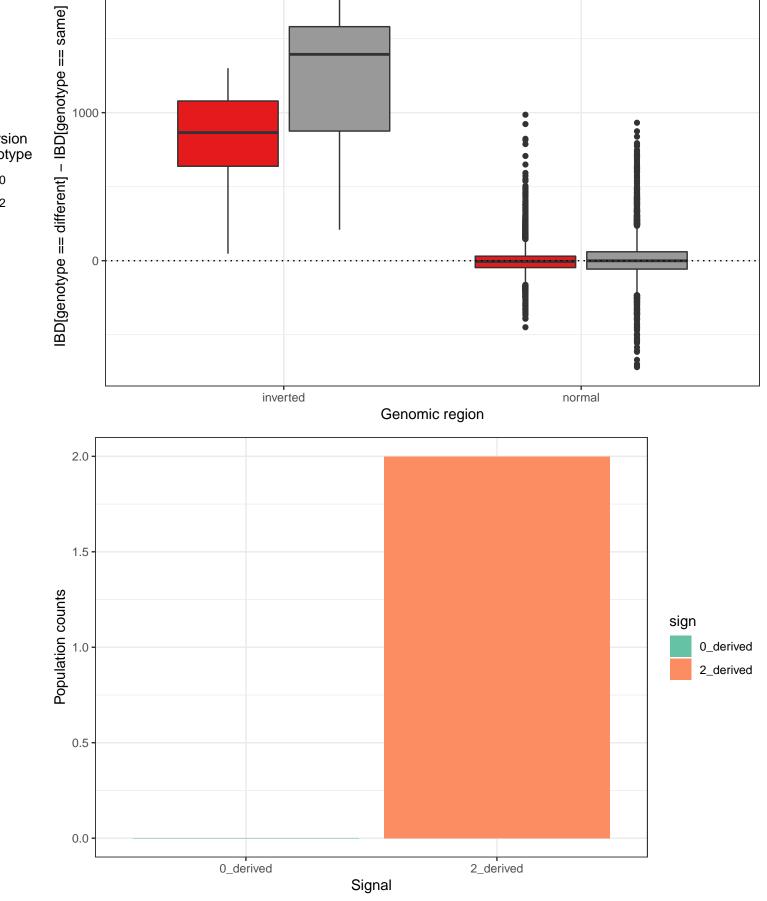
Petiolaris Ha412HOChr05 pos1 PET\_23 4000 3000 2000 IBD distance Inversion genotype PET\_24 Different - Same 4000 3000 2000 0 -MB

Petiolaris Ha412HOChr05 pos1



Genomic region

