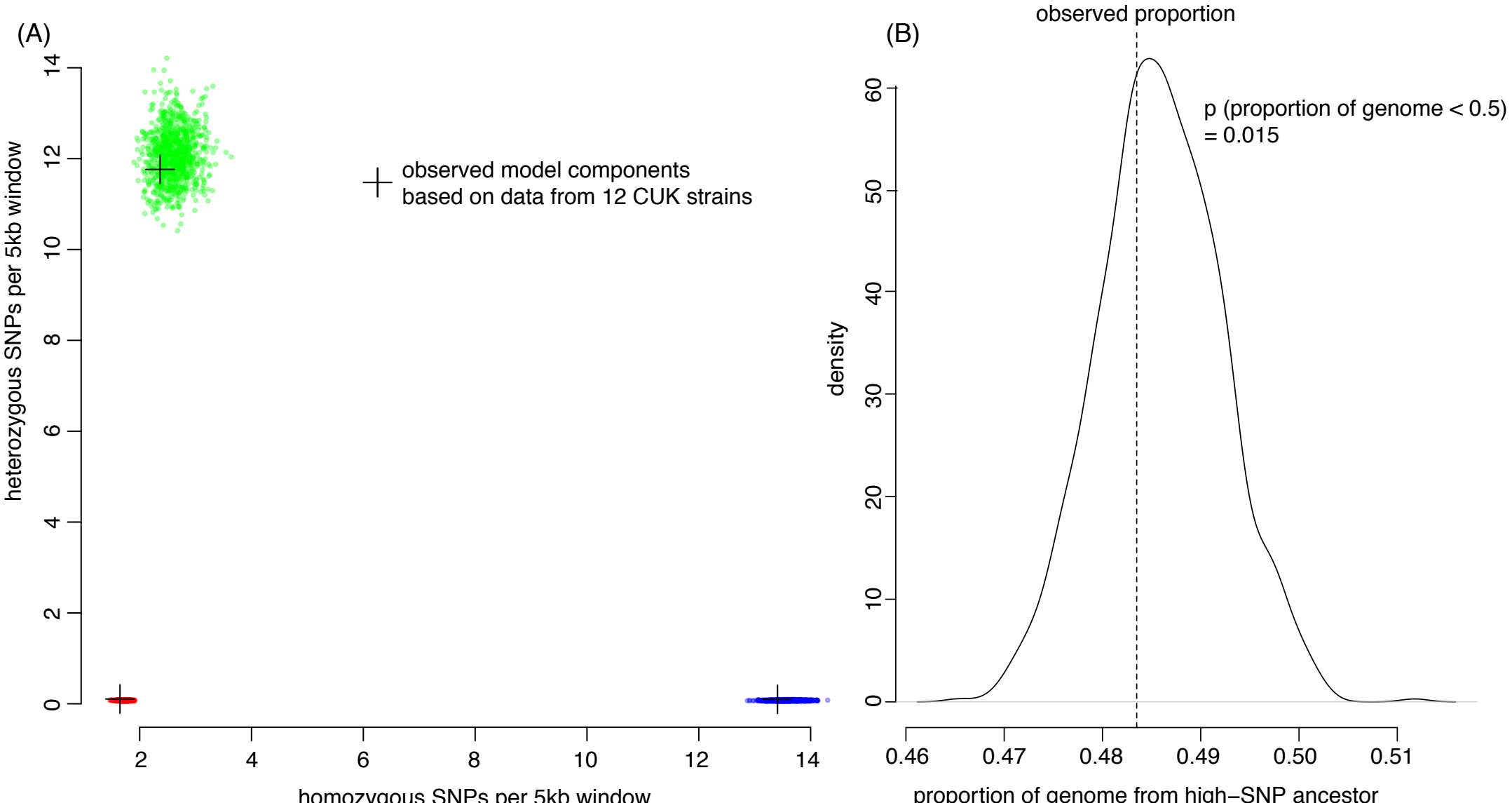
(A)
observed model components
based on data from 12 CUK strains
heterozygous SNPs per 5kb window
homozygous SNPs per 5kb window
0
2
4
6
8
10
12
14
(B)
observed proportion
p (proportion of genome < 0.5)
= 0.015
density
0
10
20
30
40
50
60
0.46
0.47
0.48
0.49
0.50
0.51
proportion of genome from high−SNP ancestor