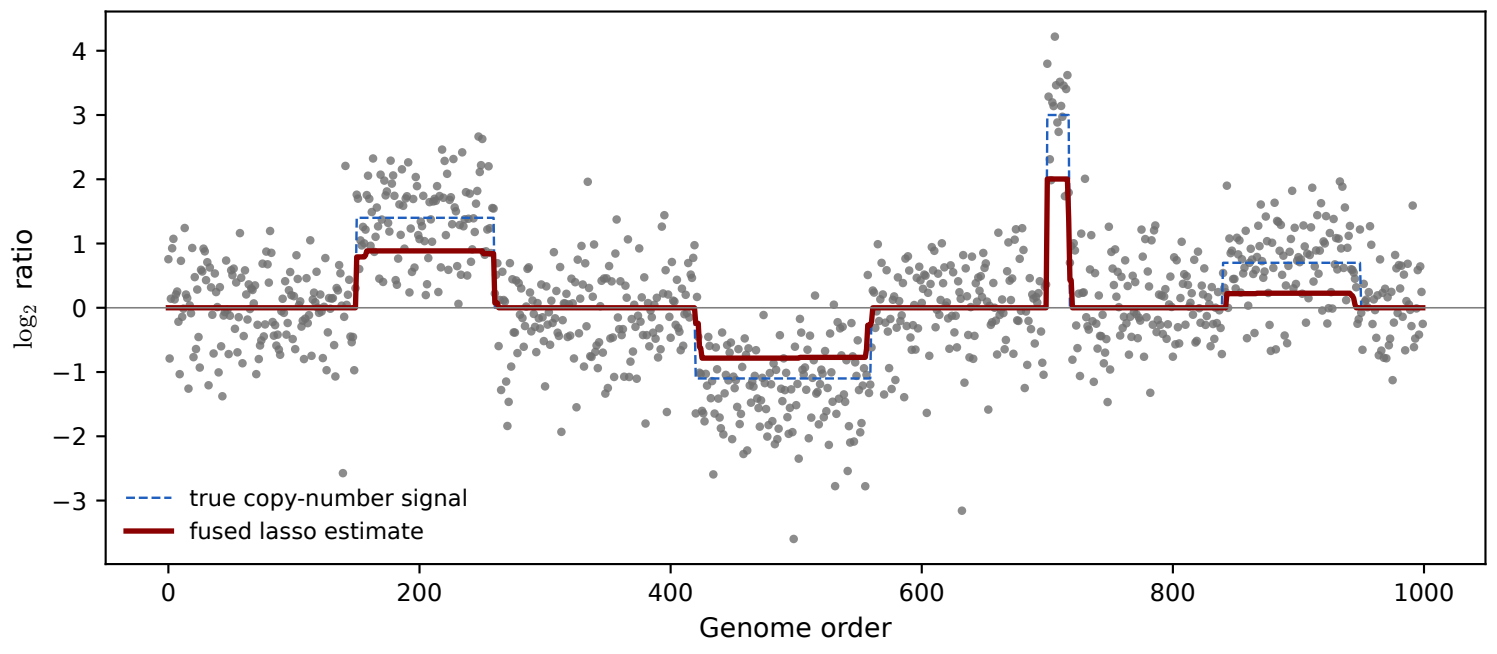

4
3
2
1
0
−1
−2
−3
log2 ratio
true copy-number signal
fused lasso estimate
0
200
400
600
800
1000
Genome order