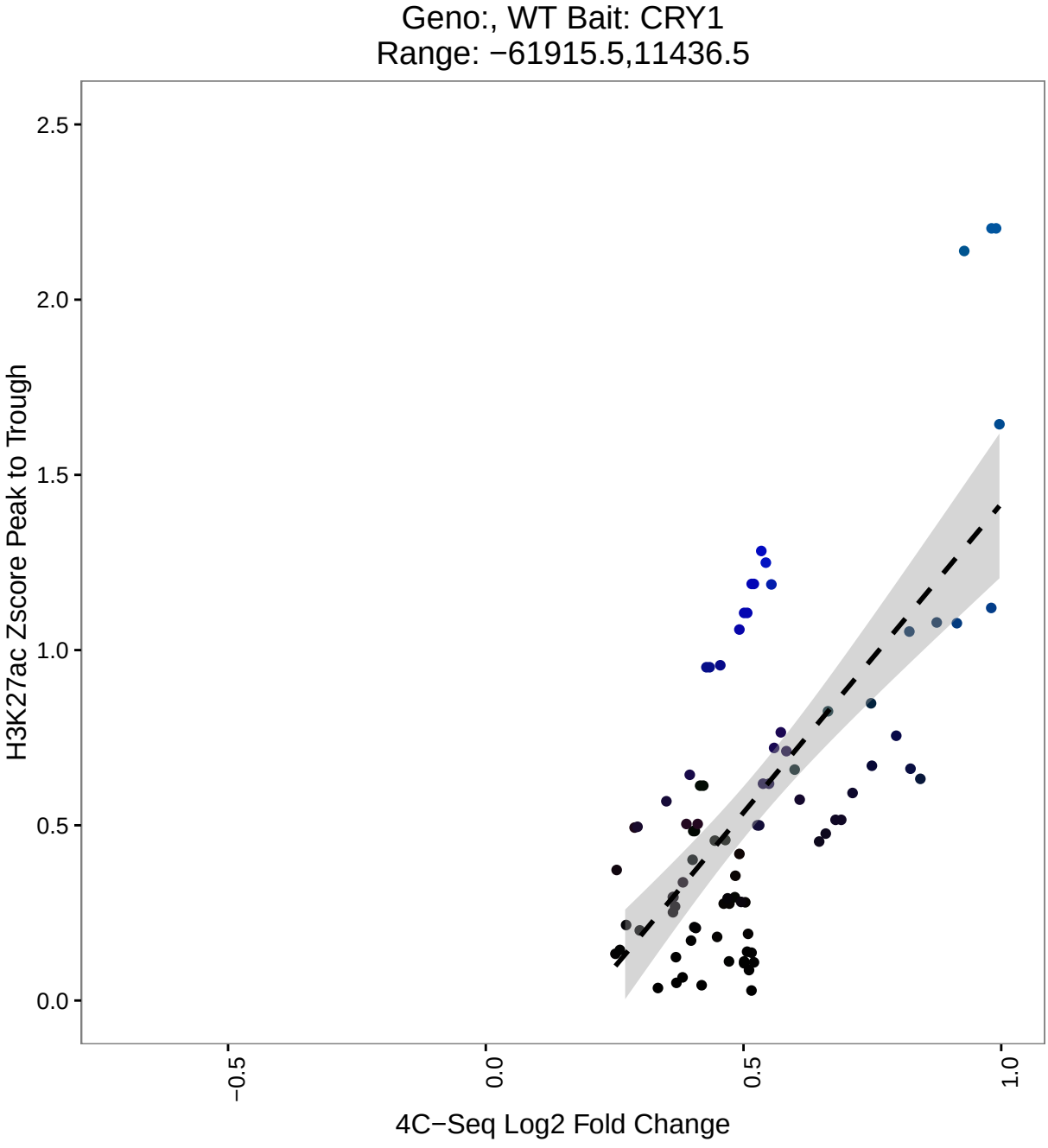
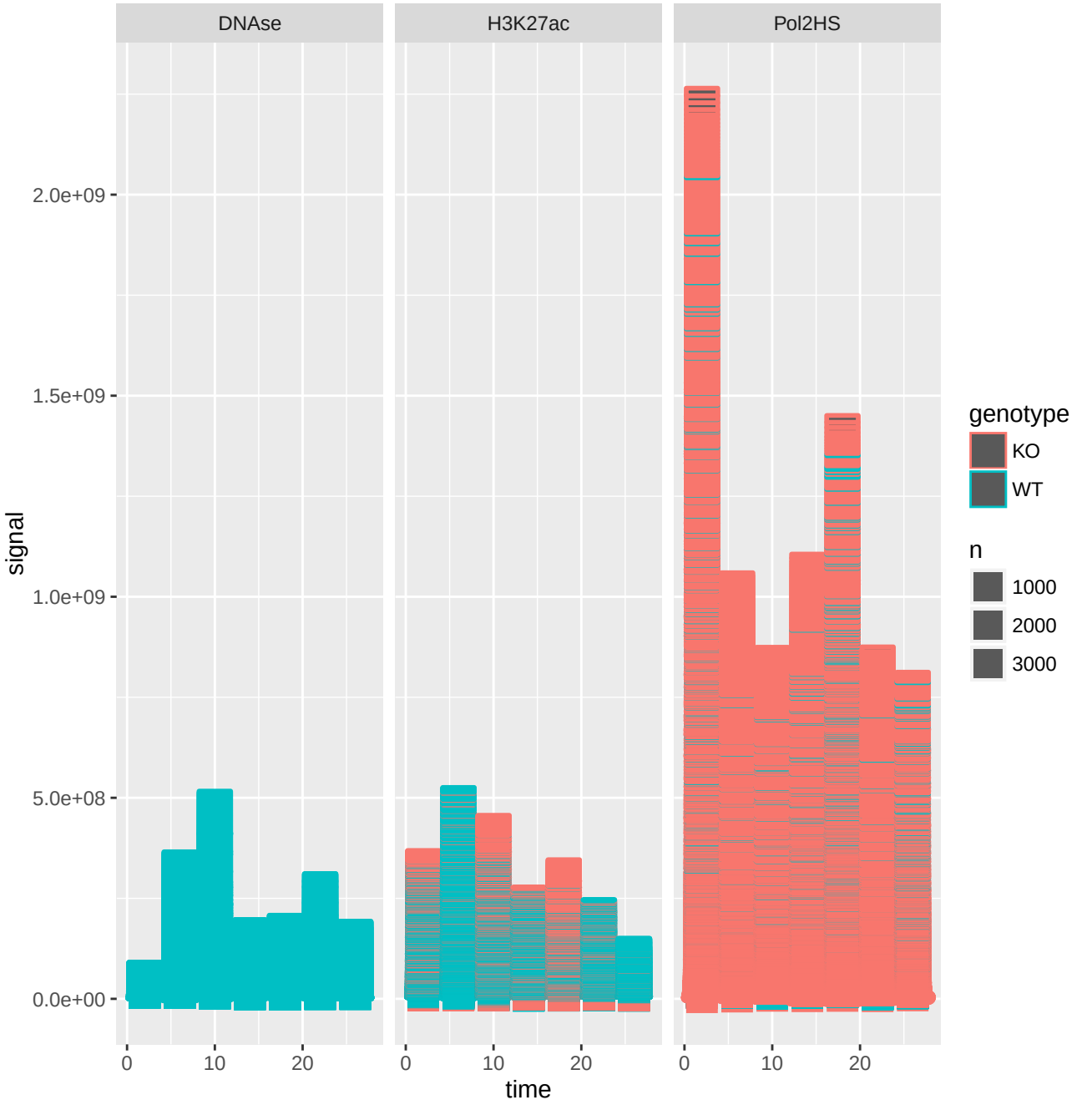

Geno:, WT Bait: CRY1
Range: −61915.5,11436.5
H3K27ac Zscore Peak to Trough
2.5
2.0
1.5
1.0
0.5
0.0
−0.5
0.0
0.5
1.0
4C−Seq Log2 Fold Change


DNAse
H3K27ac
Pol2HS
signal
2.0e+09
1.5e+09
1.0e+09
5.0e+08
0.0e+00
0
10
20
time
genotype
KO
WT
n
1000
2000
3000