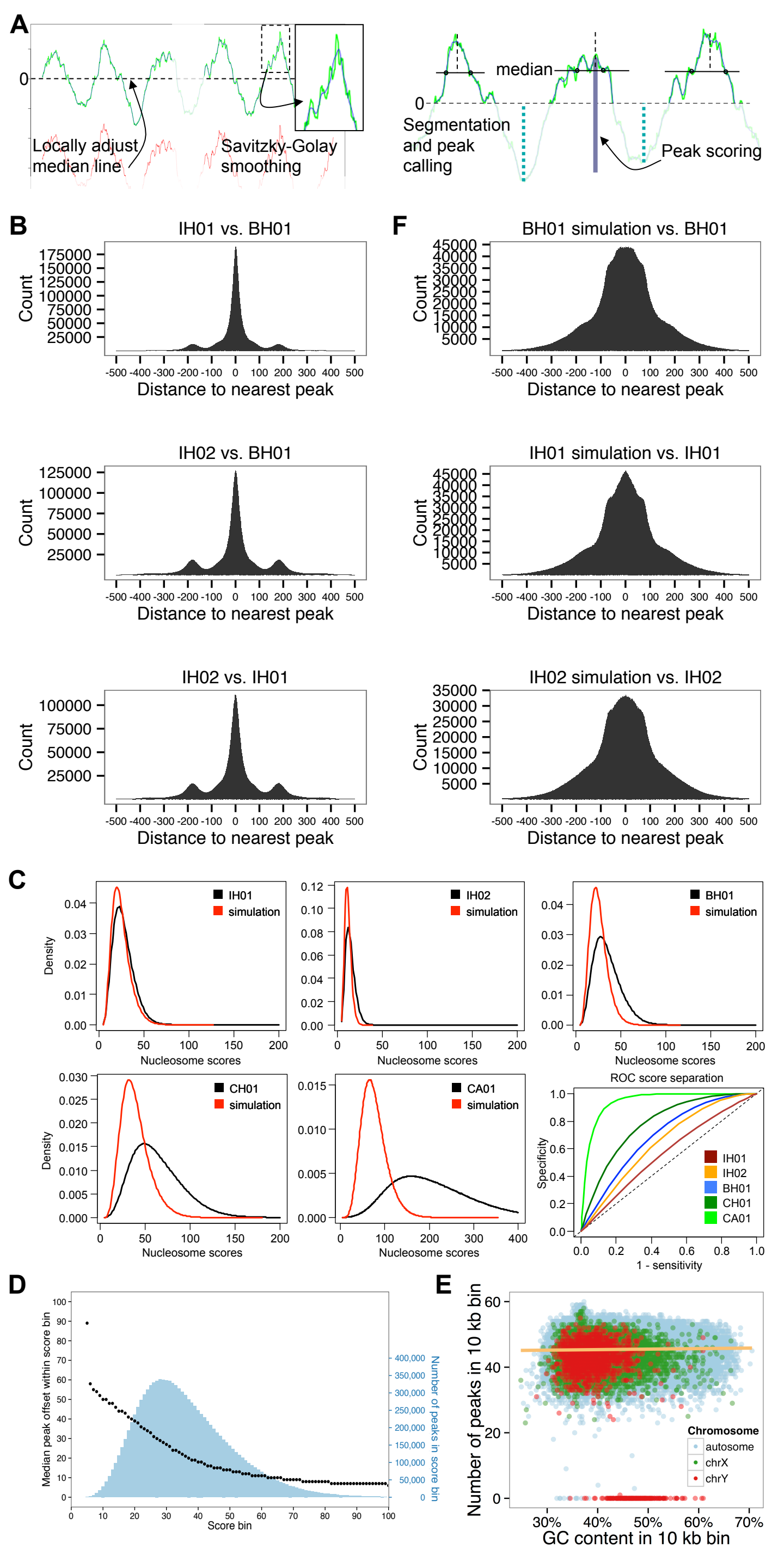
A
0
Locally adjust median line
Savitzky-Golay smoothing
median
0
Segmentation and peak calling
Peak scoring
B
IH01 vs. BH01
Count
Distance to nearest peak
IH02 vs. BH01
Count
Distance to nearest peak
IH02 vs. IH01
Count
Distance to nearest peak
F
BH01 simulation vs. BH01
Count
Distance to nearest peak
IH01 simulation vs. IH01
Count
Distance to nearest peak
IH02 simulation vs. IH02
Count
Distance to nearest peak
C
IH01
simulation
IH02
simulation
BH01
simulation
CH01
simulation
CA01
simulation
Density
Nucleosome scores
ROC score separation
Specificity
1 - sensitivity
IH01
IH02
BH01
CH01
CA01
D
Median peak offset within score bin
Number of peaks in score bin
Score bin
E
Number of peaks in 10 kb bin
GC content in 10 kb bin
Chromosome
autosome
chrX
chrY