

Parameter summary

Chromosome name:
chrV

Window size: 150

Window overlap: 50

IP threshold (T1): $> \times 6$

Control threshold (T2): $< \times 4$

Log2FC threshold (T3): > 2

Average log2 signals ([0,1]) (T4): > 0.7

Result summary

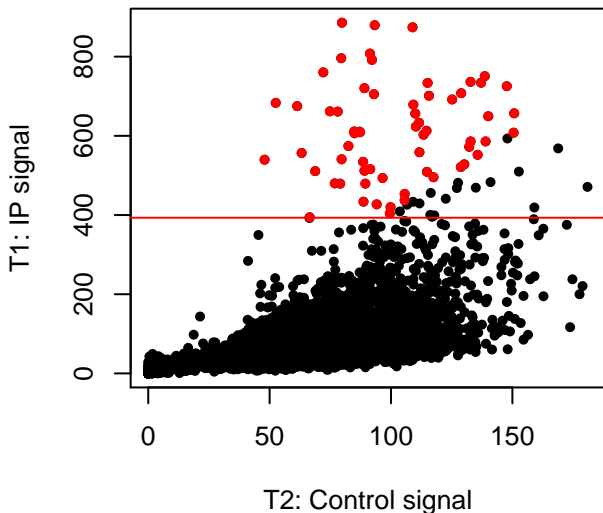
Chromosome name:
chrV

of significant windows
(before the merging procedure): 62

of detected bPeaks
(after the merging procedure): 19

Bed file name:
bPeaks_input_chrV.bed

Chromosome:
chrV



Chromosome:
chrV

