

Parameter summary

Chromosome name:
chrV

Window size: 150

Window overlap: 50

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

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

Log2FC threshold (T3): $> \ 2$

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

Result summary

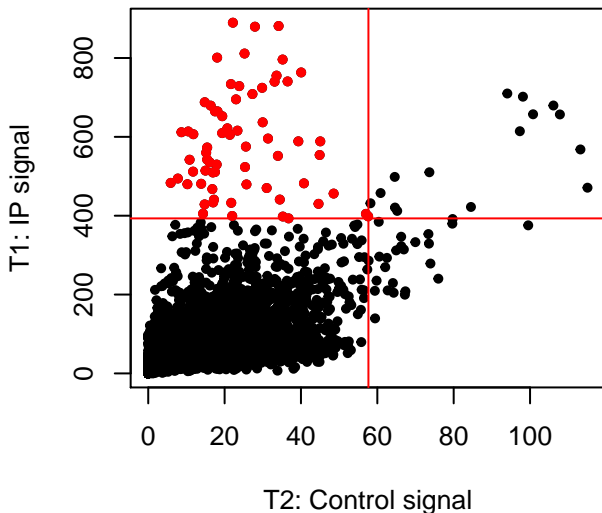
Chromosome name:
chrV

of significant windows
(before the merging procedure): 66

of detected bPeaks
(after the merging procedure): 17

Bed file name:
bPeaks_results_mock.bed

Chromosome:
chrV



Chromosome:
chrV

