

## 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

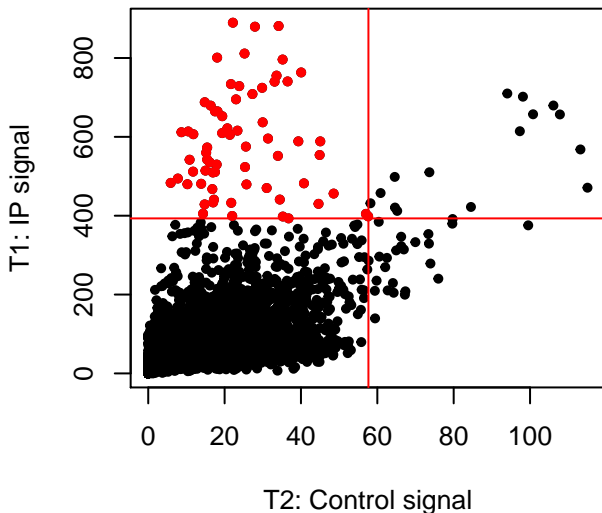
Chromosome name:  
chrV

# 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**

