

## Parameter summary

Chromosome name:  
ChrL\_C\_glabrata\_CBS138

Window size: 150

Window overlap: 50

IP threshold (T1):  $> x \times 2$

Control threshold (T2):  $< x \times 2$

Log2FC threshold (T3):  $> 2$

Average log2 signals ([0,1]) (T4):  $> 0.9$

## Result summary

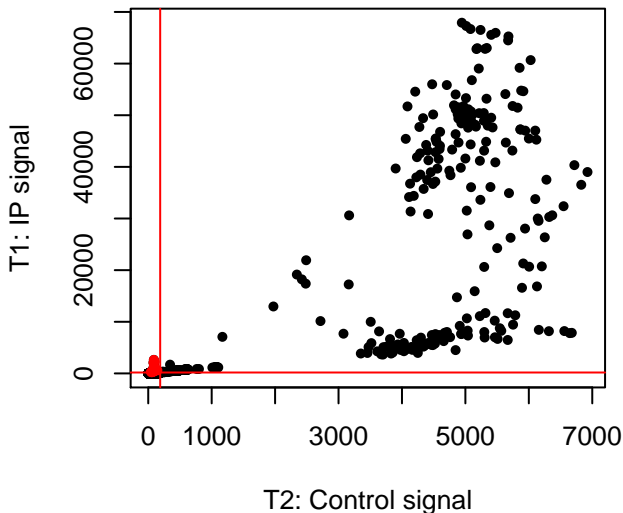
Chromosome name:  
ChrL\_C\_glabrata\_CBS138

# of significant windows  
(before the merging procedure): 24

# of detected bPeaks  
(after the merging procedure): 8

Bed file name:  
eaks\_YAP1-ChrL\_C\_glabrata\_CBS138.l

**Chromosome:**  
**ChrL\_C\_glabrata\_CBS138**



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