

Parameter summary

Chromosome name:
ChrF_C_glabrata_CBS138

Window size: 150

Window overlap: 50

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

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

Log2FC threshold (T3): > 2

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

Result summary

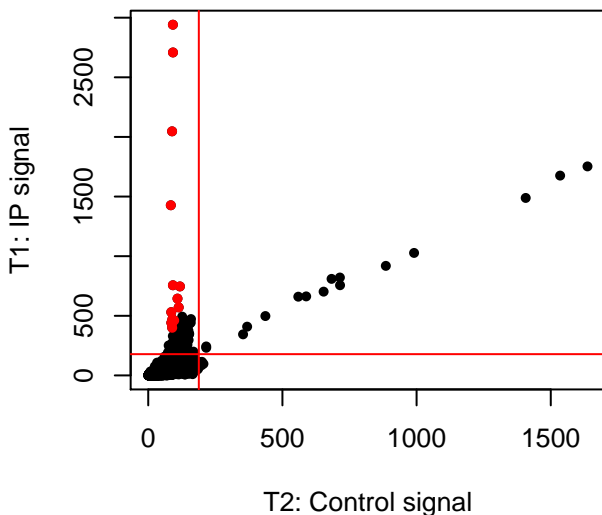
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of significant windows
(before the merging procedure): 16

of detected bPeaks
(after the merging procedure): 5

Bed file name:
saks_YAP1-ChrF_C_glabrata_CBS138.l

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