

Parameter summary

Chromosome name:
mito_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

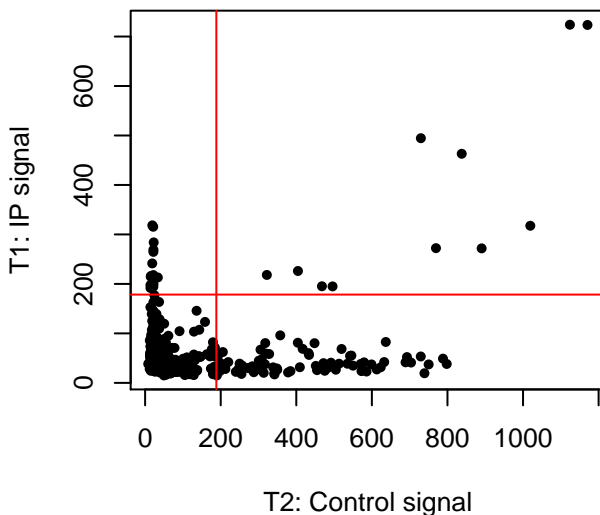
Result summary

Chromosome name:
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of significant windows
(before the merging procedure): 0

of detected bPeaks
(after the merging procedure): 0

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