

Coordinates (nucleotides)

Cluster Read Counts (Peaks in middle of clusters) input1.bam input2.bam 20 0e+00 2e+05 4e+05 6e+05 0e+00 2e+05 4e+05 6e+05 input1.bam input2.bam 1000 200 2e+05 0e+00 2e+05 0e+00 4e+05 6e+05 8e+05 4e+05 6e+05 8e+05 input1.bam input2.bam 1000 500 200000 400000 600000 800000 1000000 200000 400000 800000 1000000 600000 input1.bam input2.bam 1000 2000 4e+05 6e+05 0e+00 2e+05 4e+05 6e+05 8e+05 0e+00 2e+05 8e+05 input2.bam input1.bam 80 9 9 20 500000 0 500000 1000000 1000000 input1.bam input2.bam 4 30 20 10 0 0 500000 500000 1000000 1500000 1000000 1500000 input1.bam input2.bam 200 100 0 500000 1000000 1500000 0 500000 1500000 1000000 input1.bam input2.bam 50 500000 1000000 0 500000 1000000 1500000 1500000

Read Counts

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