The IP appears weak
The percentage enrichment of IP over Input is not statistically significant

Divergence test q–value (pFDR): 0.369995
TN: 0.413925 HN: 0.431307 TC: 0.252079 HC: 0.311304
Scaling factor: 0.413925 (scale input by this amount)
1.9966% of the genome is enriched for signal.
Cumulative % enrichment of IP over Input: 58.625%

The IP appears weak
Significant enrichment for signal in IP over Input
Divergence test q–value (pFDR): 6.84e-26
Scaling factor: 0.61192 (scale input by this amount)
2.2007% of the genome is enriched for signal.
Cumulative % enrichment of IP over Input: 19.1968%

Warnings:
Possible PCR amplification bias in Input, more than 25% of the reads map to less than 0.01% of the genome. Consider de–duplicating your reads and running CHANCE again.