

Plotting parameters:

```
txdb: new("TxDb", .xData = <environment>)  
      treat: c(query =  
"C:/Users/greenblatt/AppData/Local/R/win-library/4.3/GenomicPlot/extdata/treat_chr19.bam")  
      control: c(input =  
"C:/Users/greenblatt/AppData/Local/R/win-library/4.3/GenomicPlot/extdata/input_chr19.bam")  
      feature: alist[["feature"]]  
param: list(offset = -1, fix_width = 0, fix_point = "start", norm = TRUE, useScore =  
FALSE, outRle = TRUE, useSizeFactor = FALSE, genome = "hg19")
```