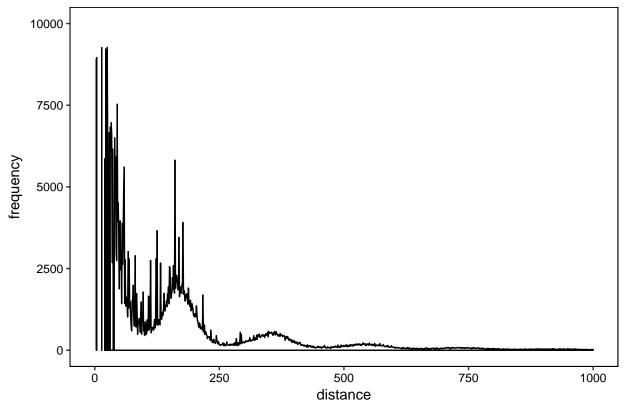
methylation distance on promoters

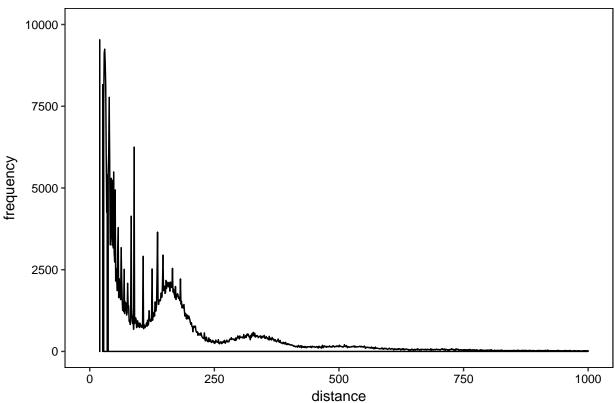
isaclee 5/6/2019

write out the names of high vs low expression genes subset gene windows by expression get distances between open marks

highexpression_promoters



lowexpression_promoters



```
## # A tibble: 2 x 3
## # Groups:
               regname [2]
     regname
                              region dist
##
     <chr>
                              <chr>
                                     <dbl>
##
## 1 highexpression_promoters all
                                     108.
## 2 lowexpression_promoters
                                      92.6
## # A tibble: 2 x 3
     regname
##
                              median mean
##
     <chr>
                               <dbl> <dbl>
## 1 highexpression_promoters
                                149. 157.
## 2 lowexpression_promoters
                                102. 117.
```

