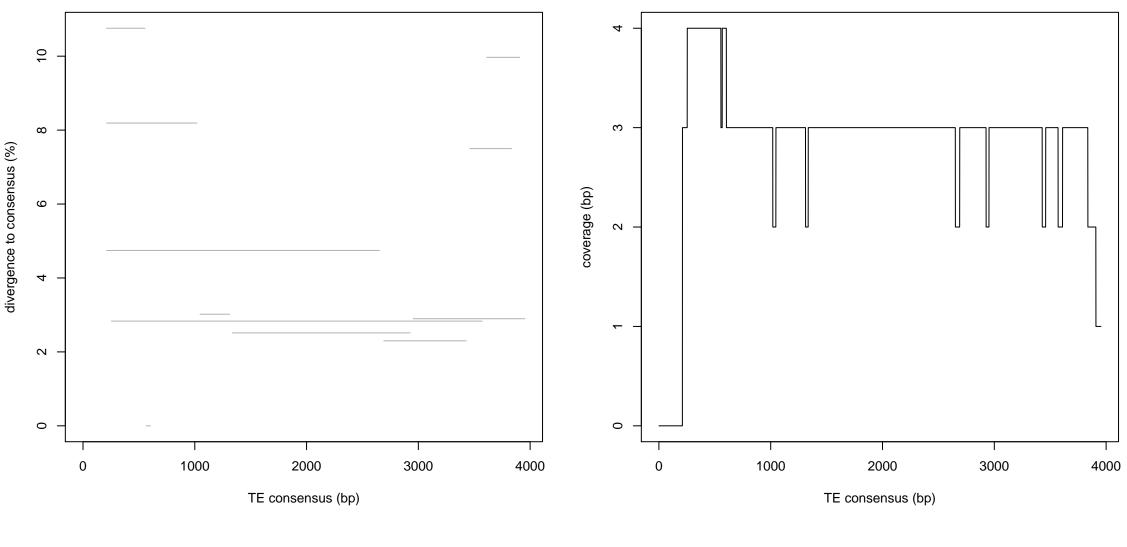
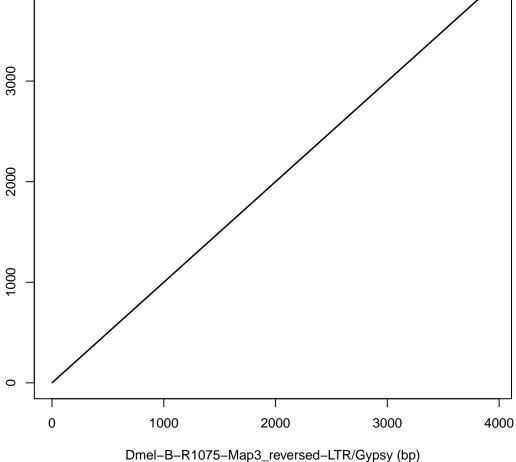
## TE consensus genomic coverage





4000

Dmel-B-R1075-Map3\_reversed-LTR/Gypsy (bp)



## TE consensus structure and protein hits

