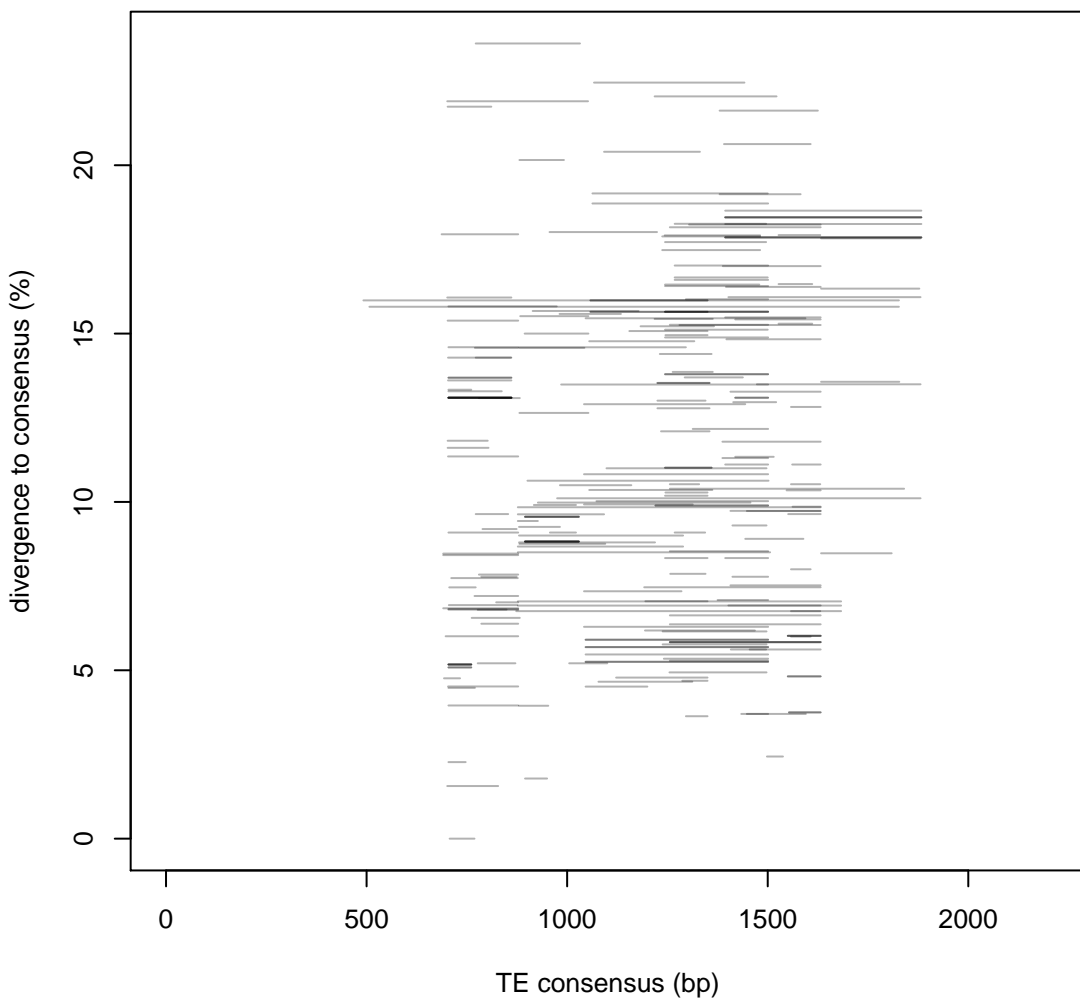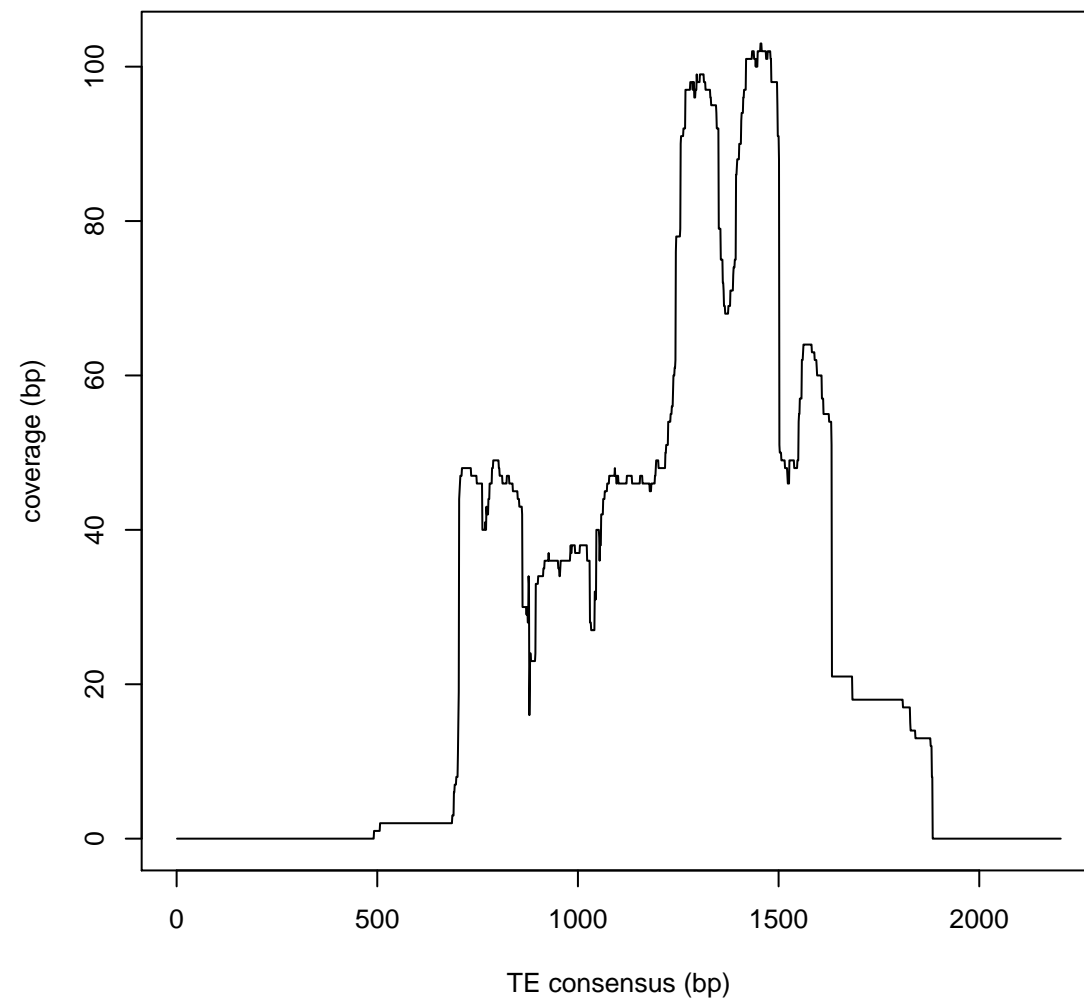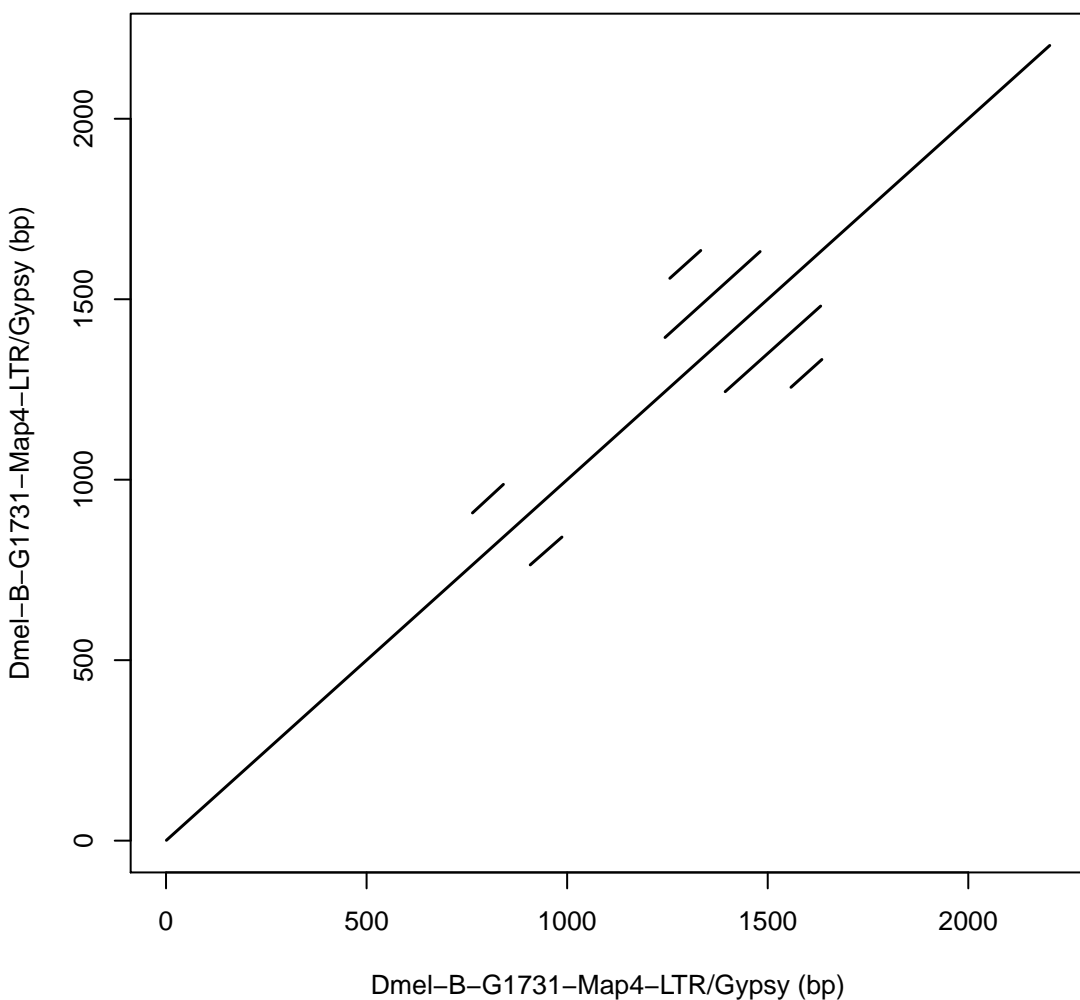## TE: Dmel−B−G1731−Map4−LTR/Gypsy
**consensus size: 2203bp; fragments: 254; full length: 0 (>=1982.7bp)**



## TE consensus genomic coverage



## TE consensus self dotplot (blastn)



## TE consensus structure and protein hits

no orf >400 bp detected