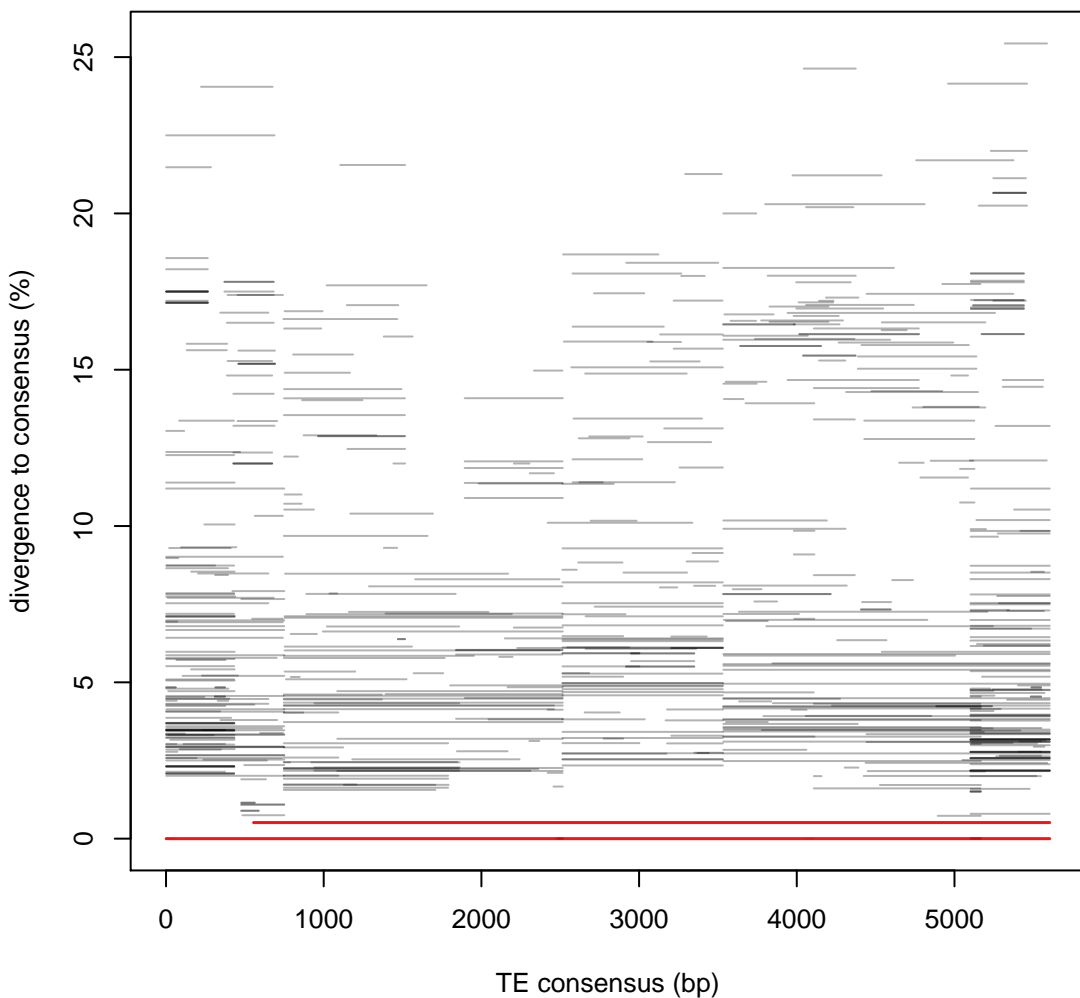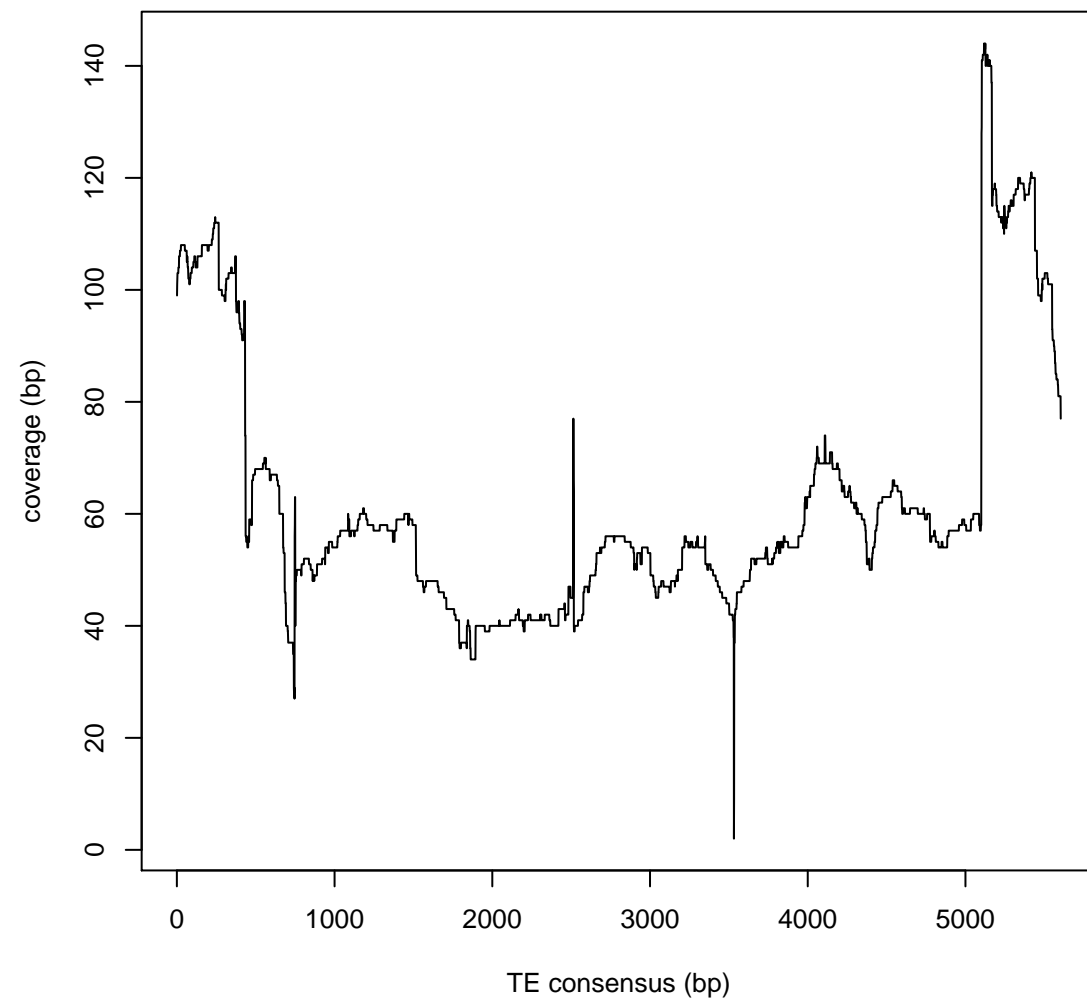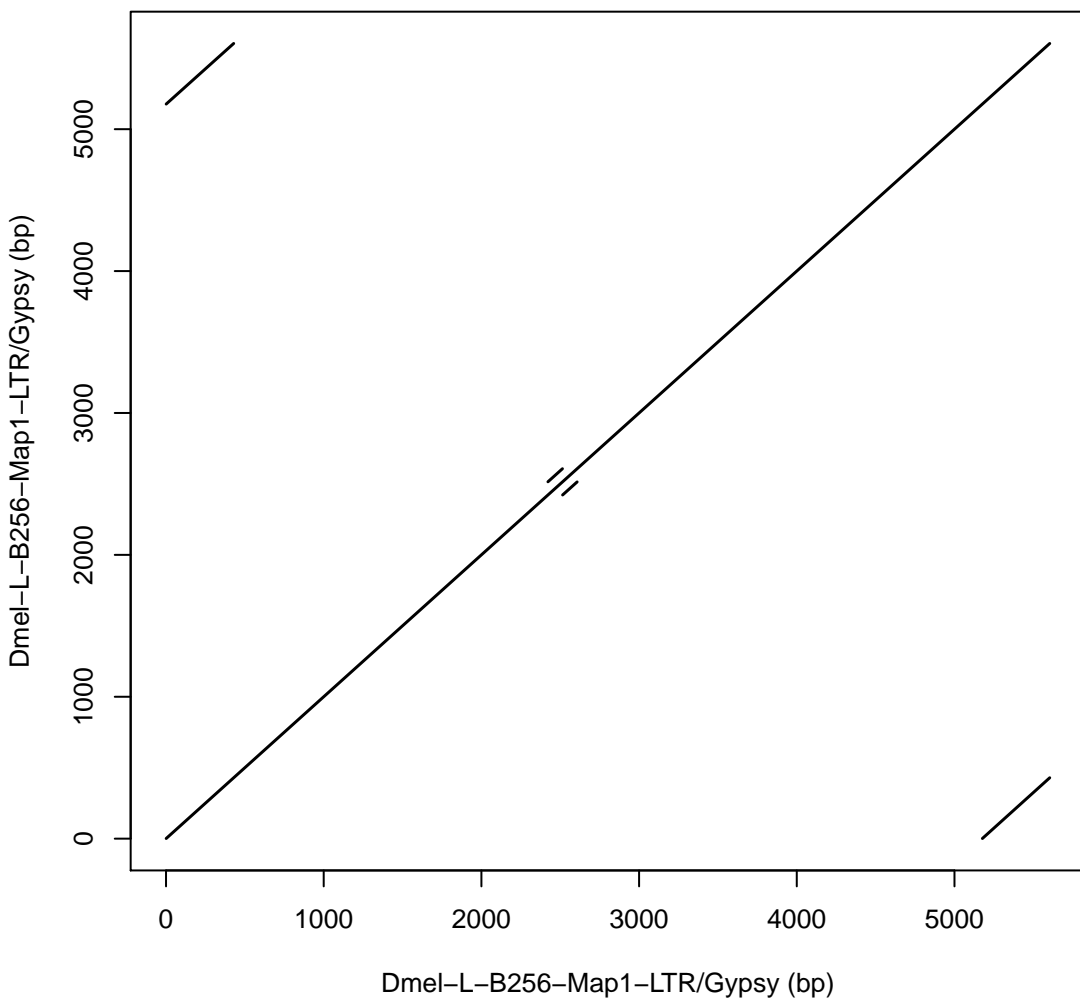## TE: Dmel−L−B256−Map1−LTR/Gypsy
### consensus size: 5604bp; fragments: 626; full length: 2 (>=5043.6bp)



## TE consensus genomic coverage



## TE consensus self dotplot (blastn)



## TE consensus structure and protein hits

no orf >400 bp detected