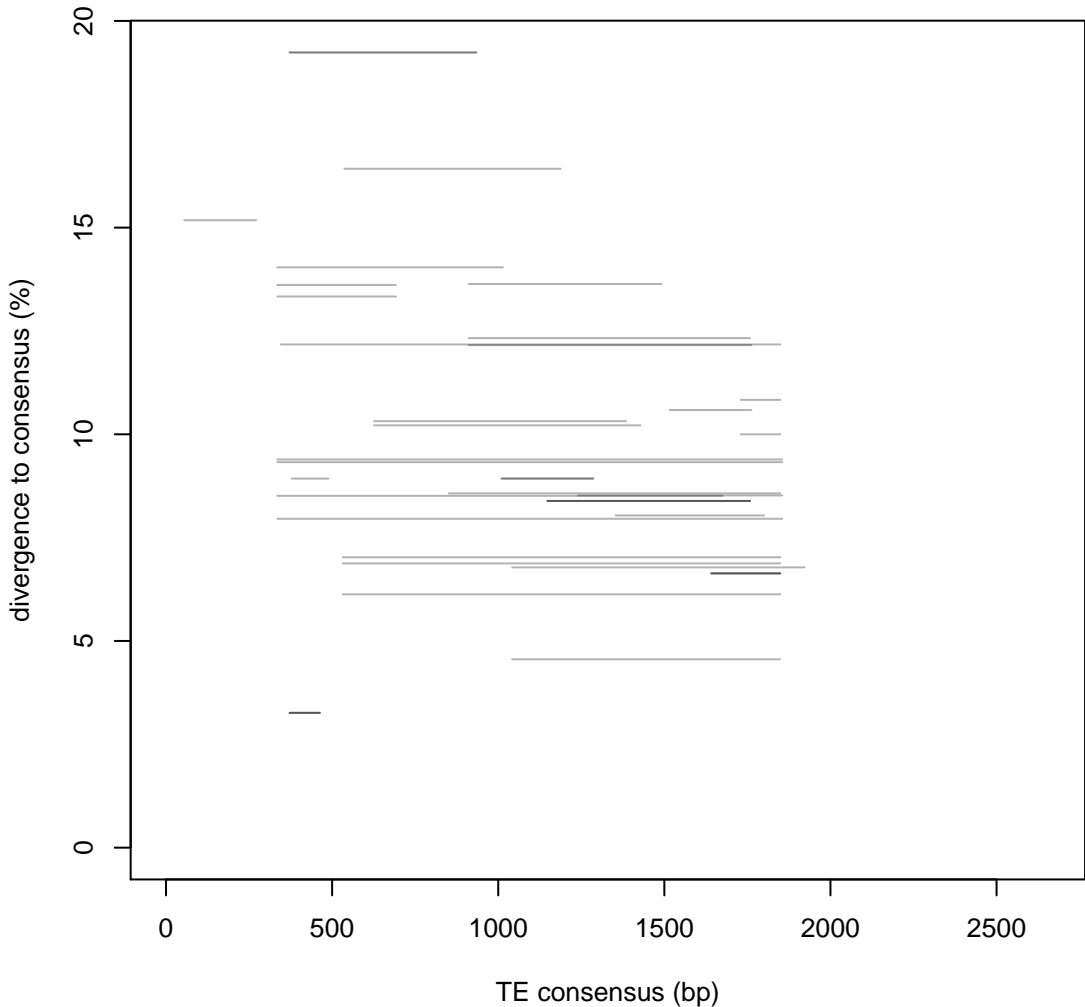
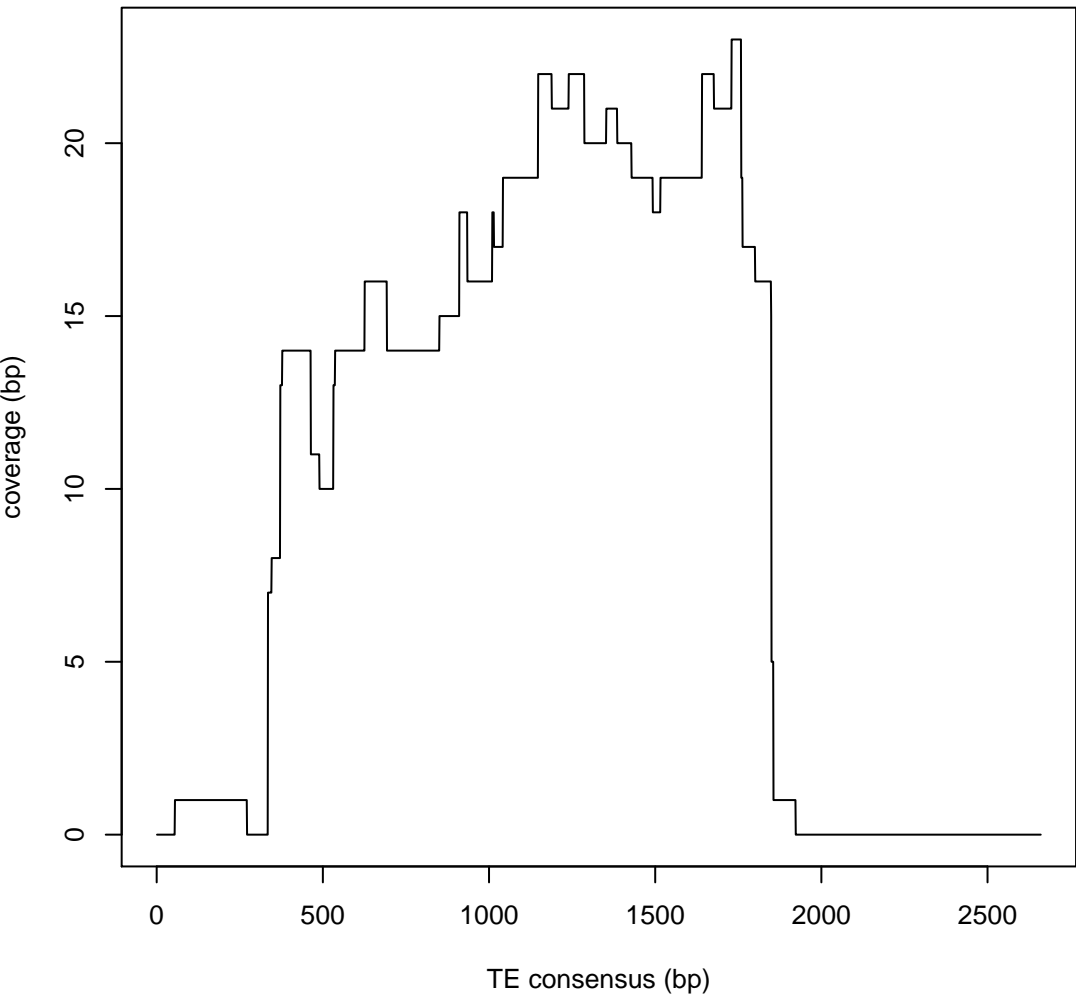


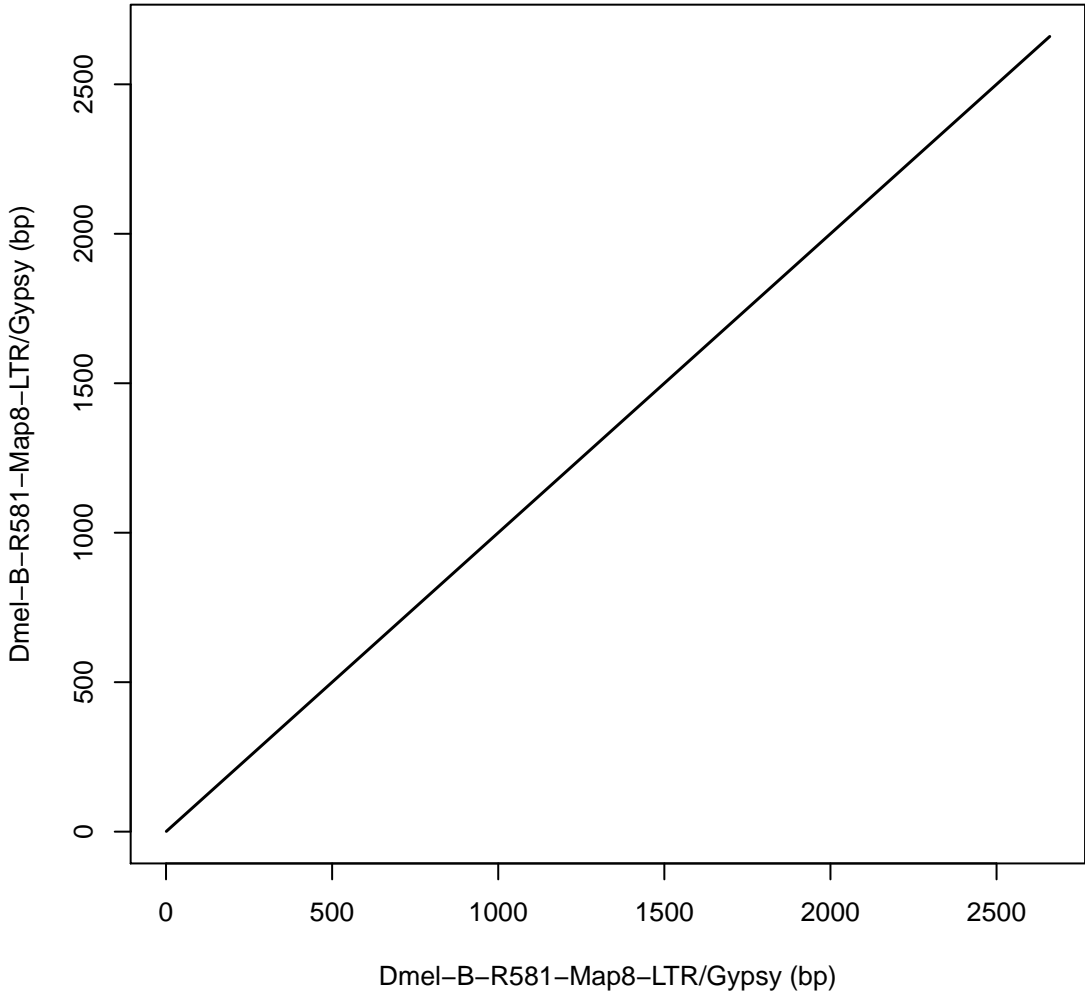
TE: Dmel-B-R581-Map8-LTR/Gypsy
consensus size: 2660bp; fragments: 40; full length: 0 (>=2394bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

