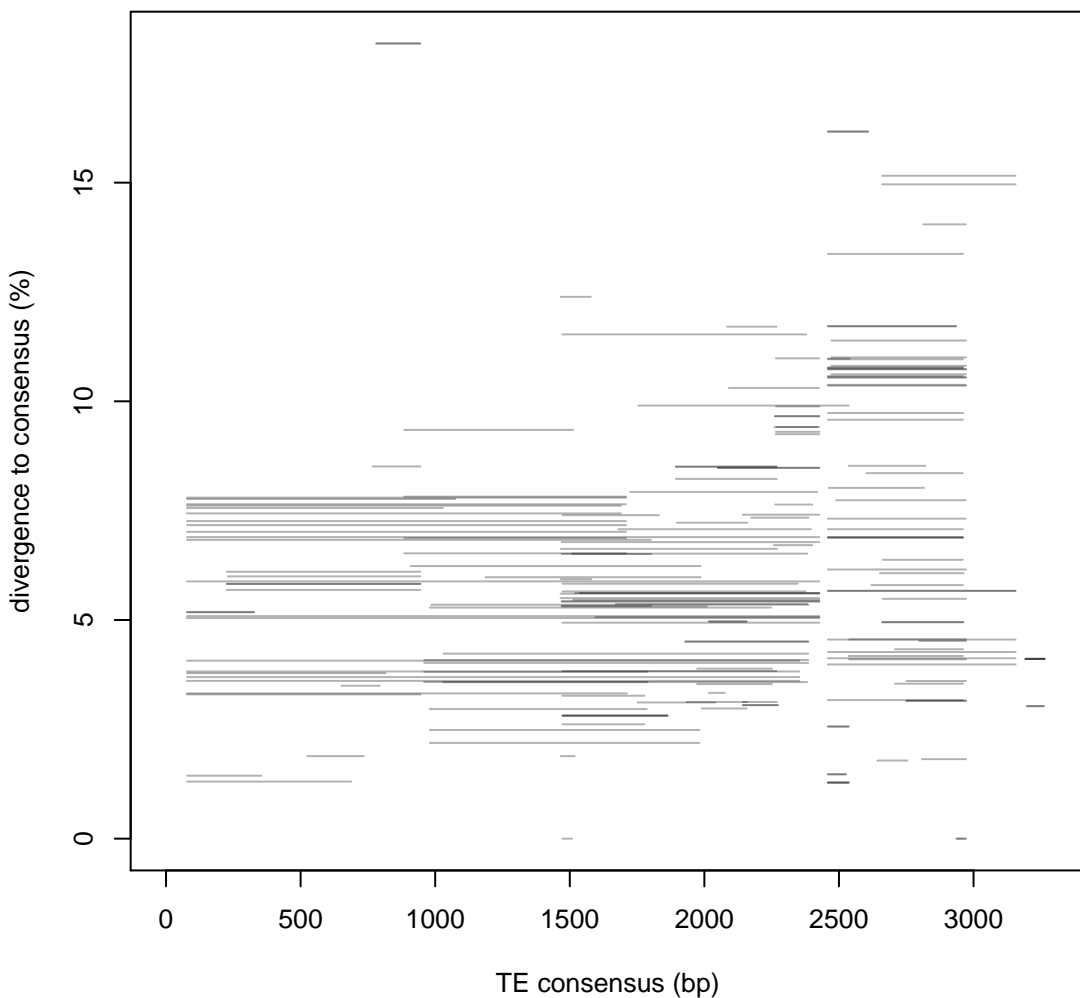
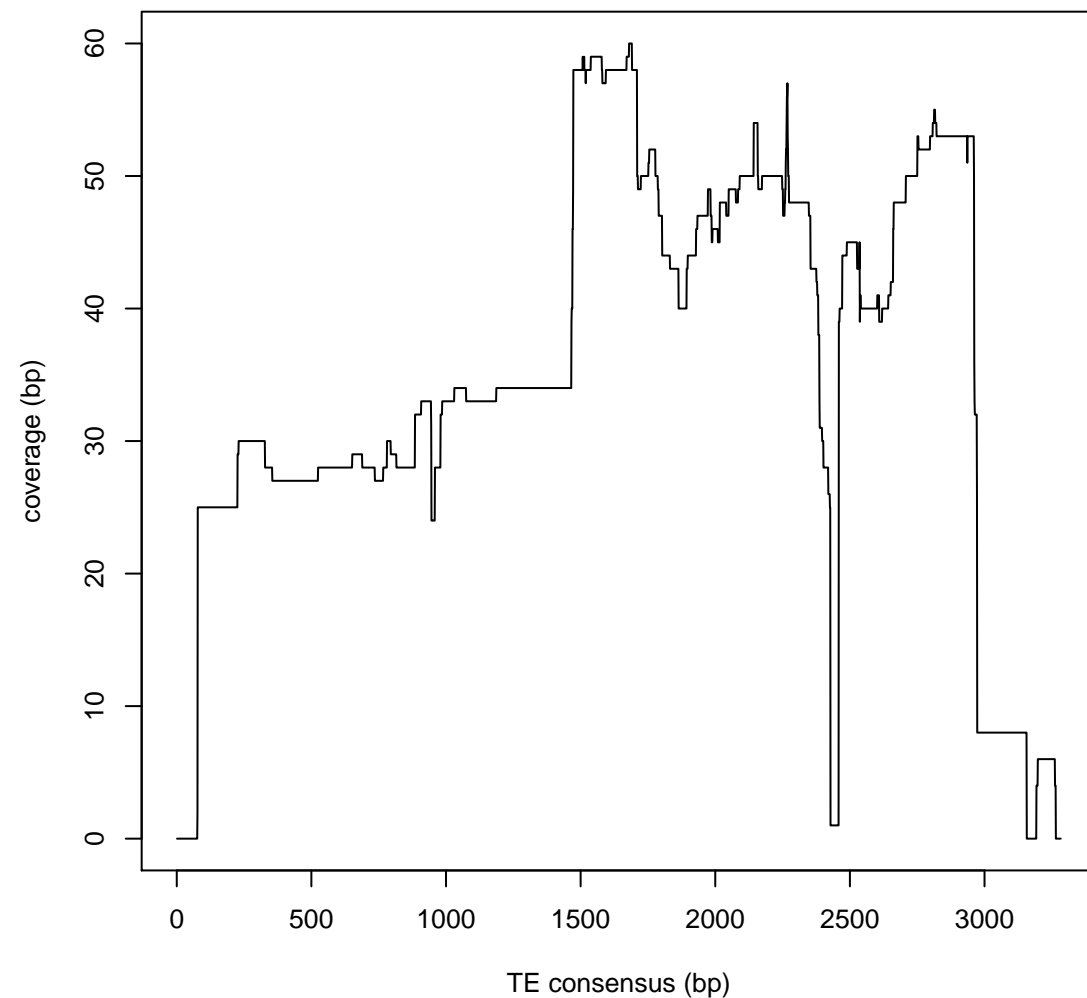


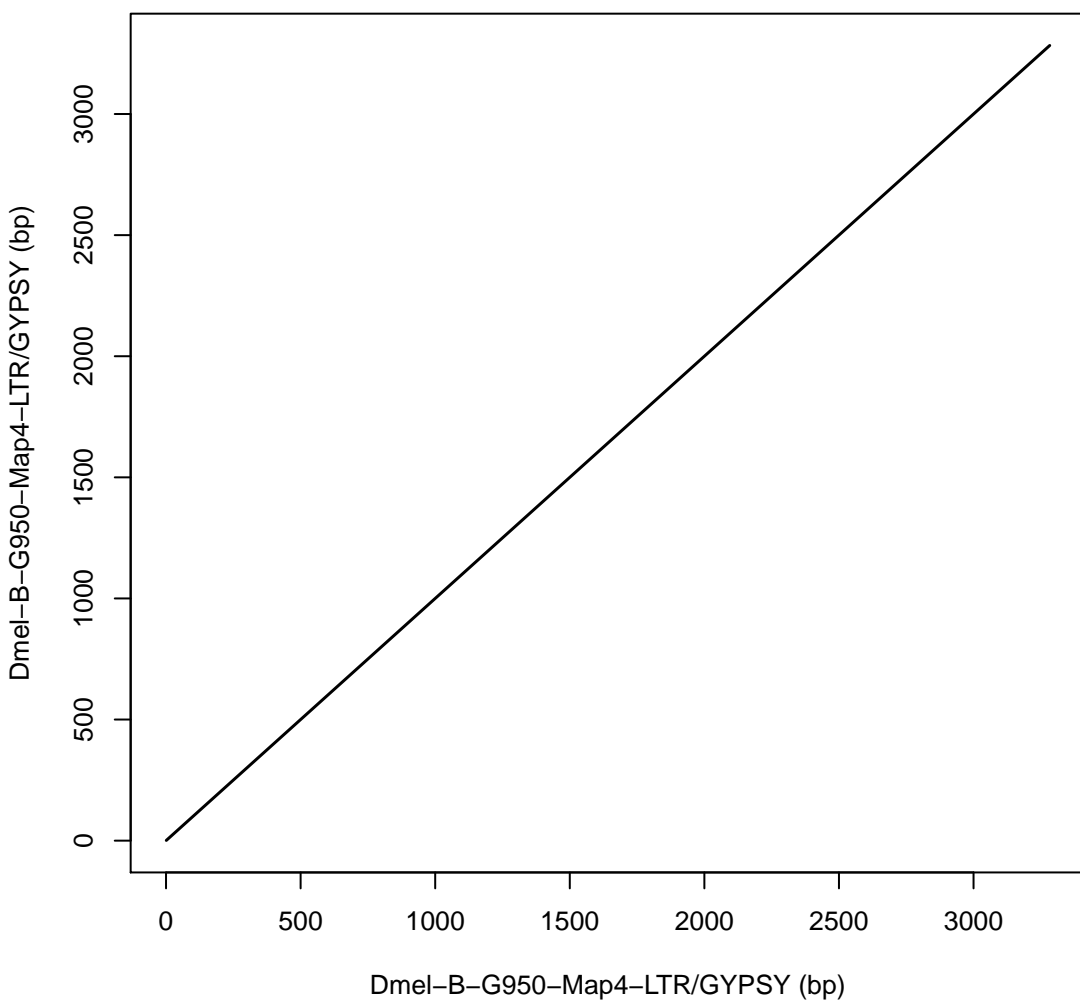
TE: Dmel-B-G950-Map4-LTR/GYPSY
consensus size: 3283bp; fragments: 192; full length: 0 (>=2954.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

