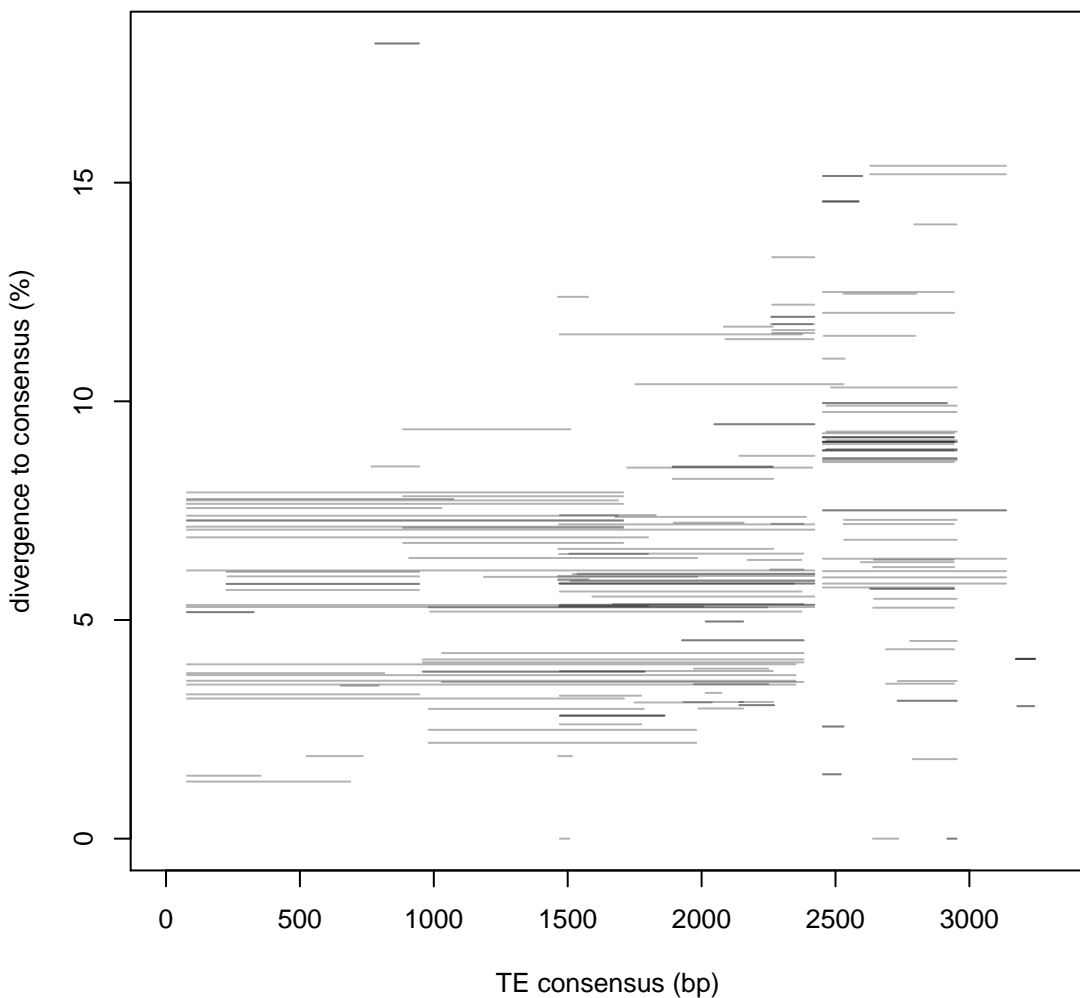
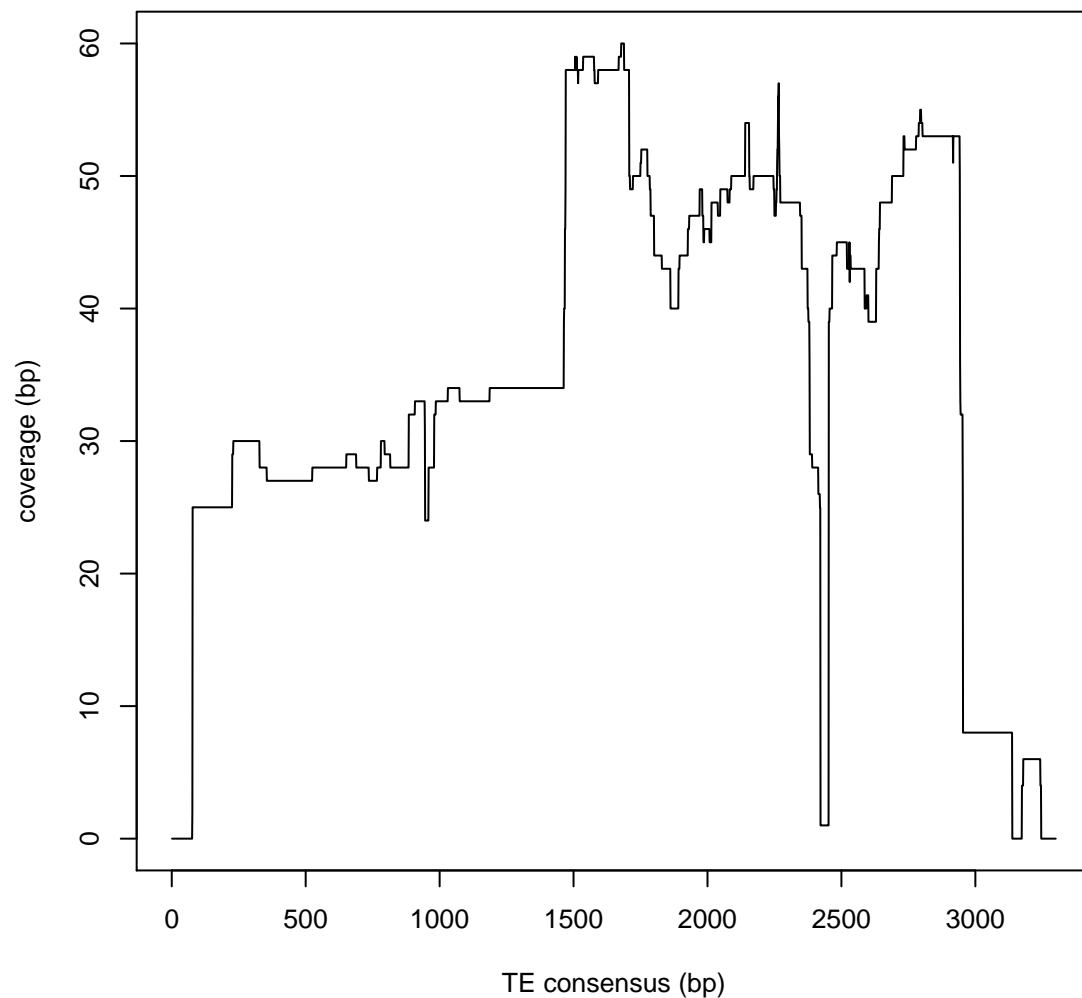


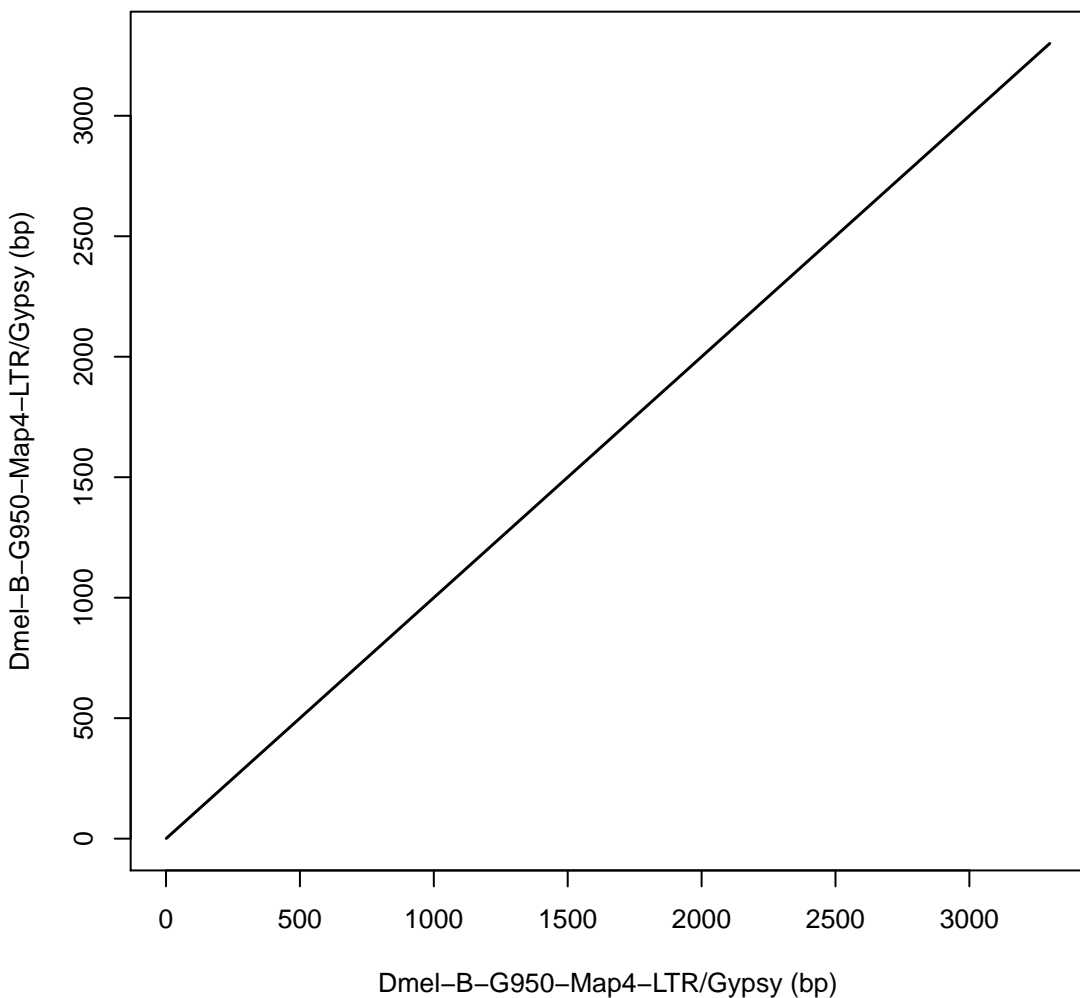
TE: Dmel-B-G950-Map4-LTR/Gypsy
consensus size: 3300bp; fragments: 192; full length: 0 (>=2970bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

