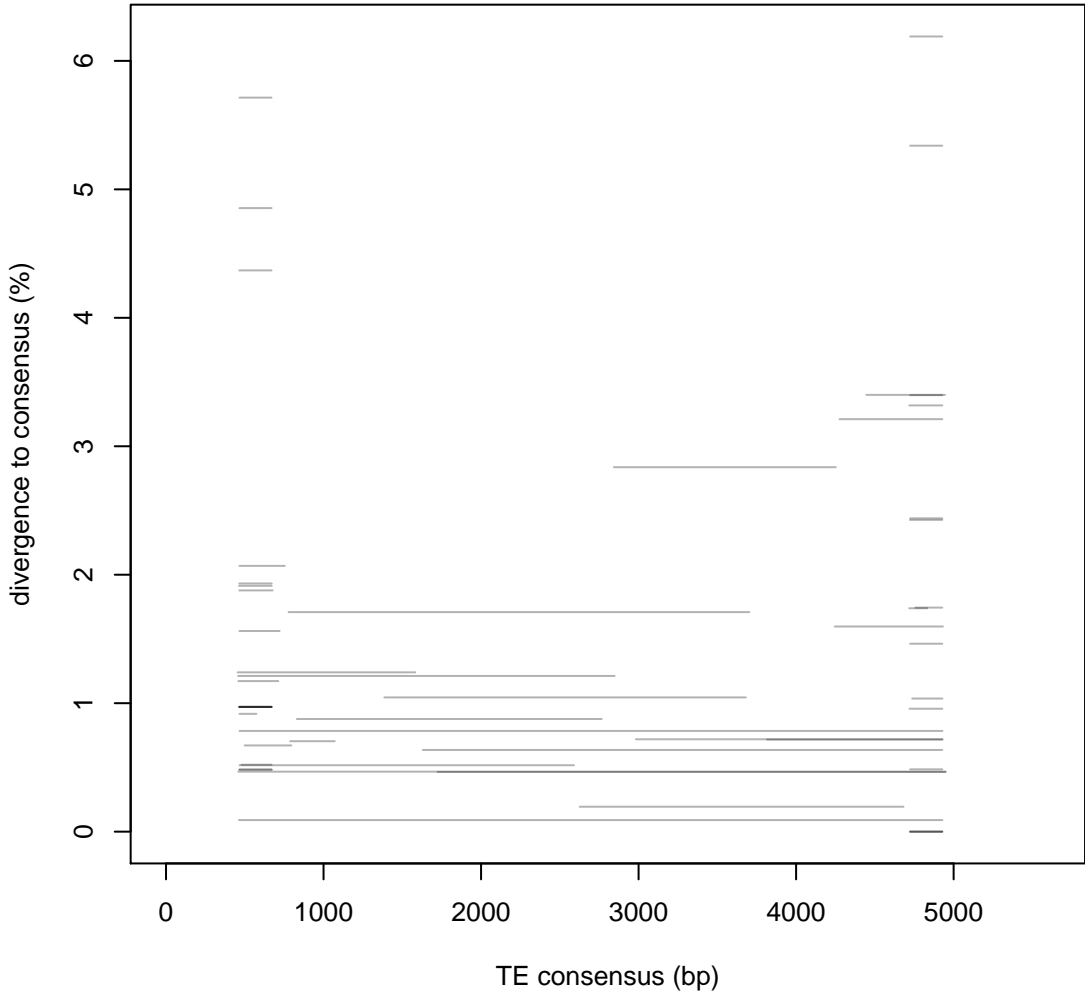
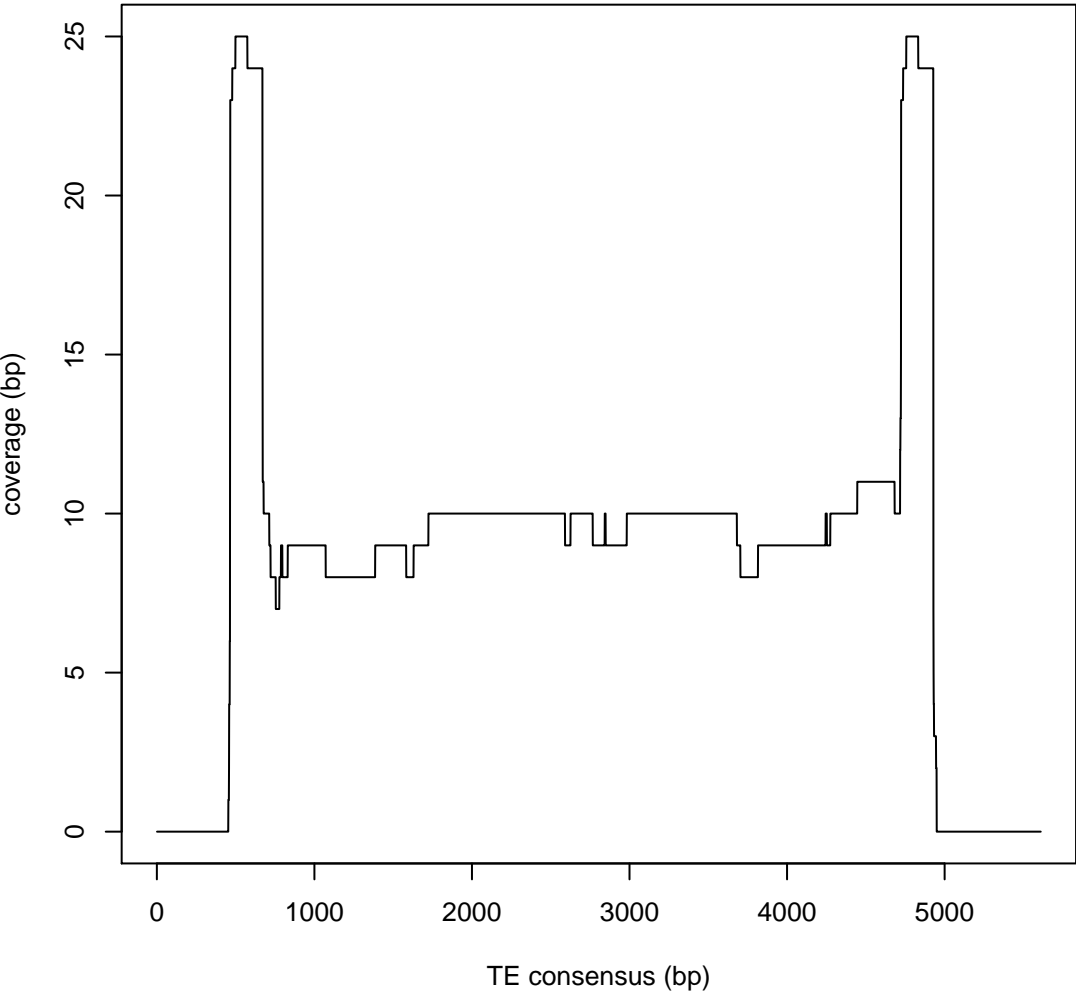


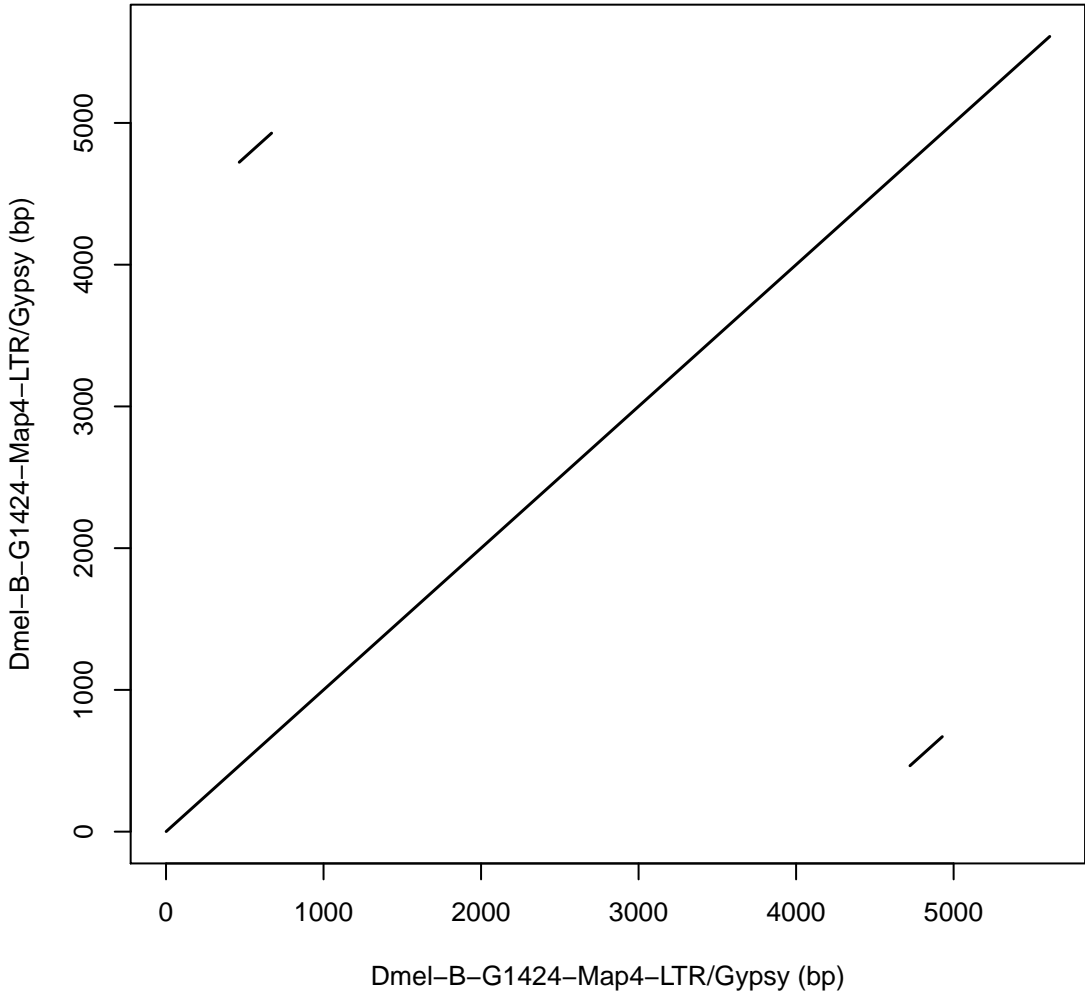
TE: Dmel-B-G1424-Map4-LTR/Gypsy
consensus size: 5610bp; fragments: 53; full length: 0 (>=5049bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

