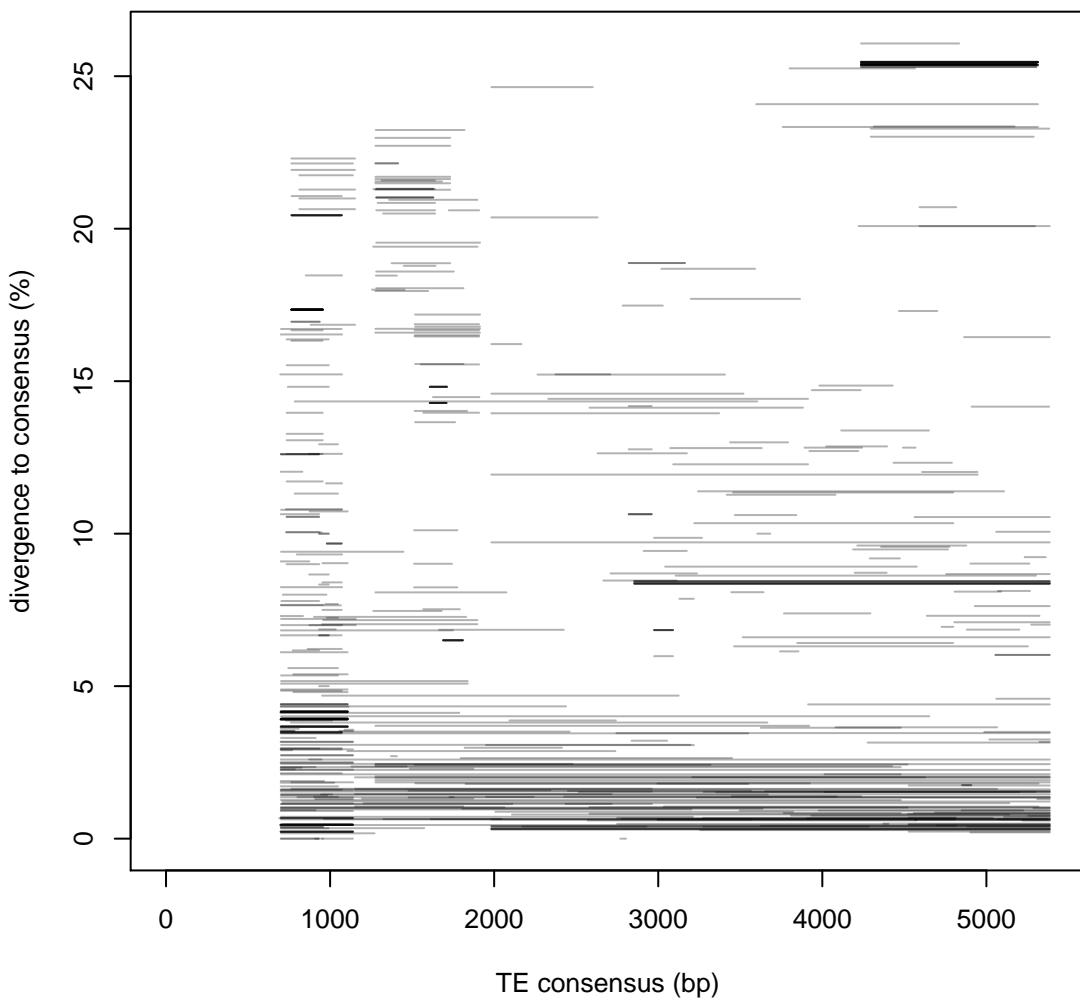
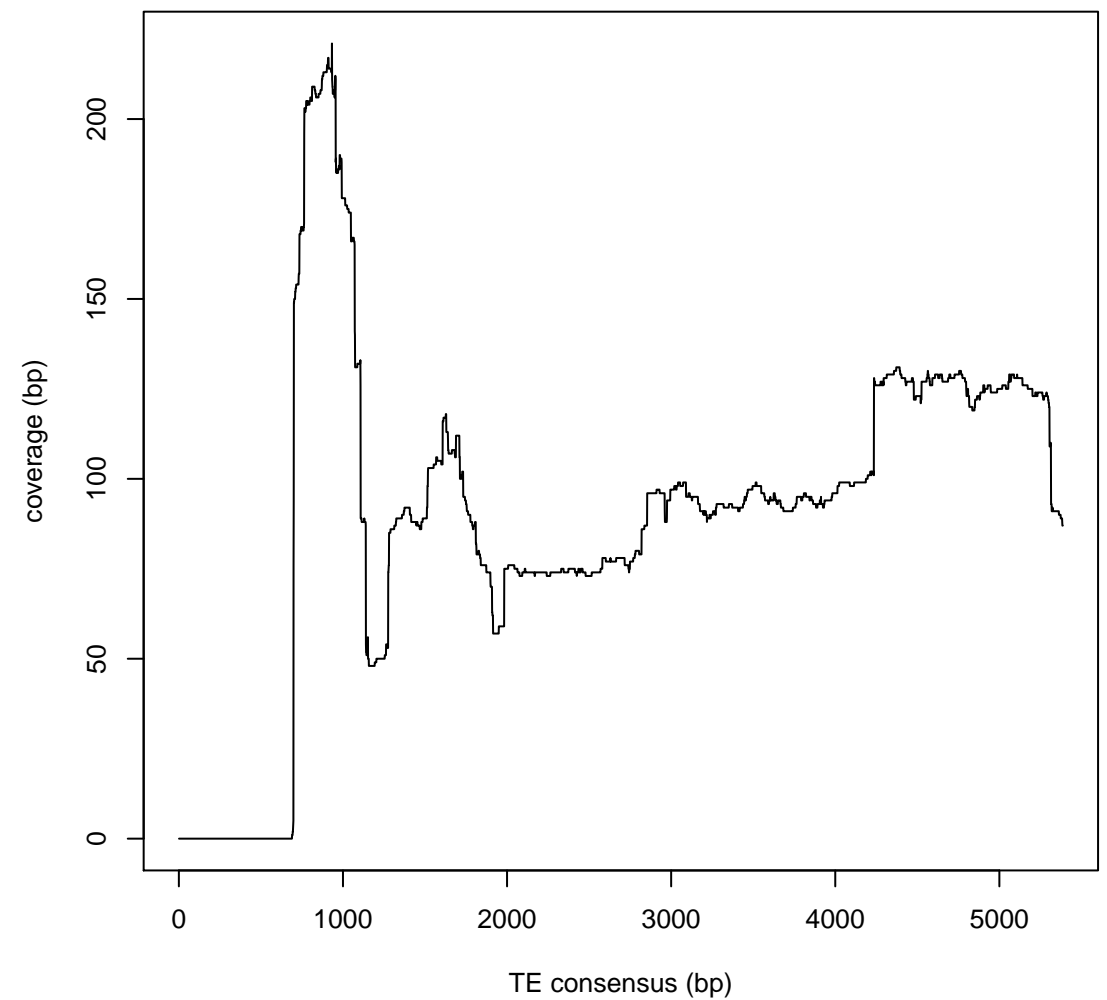


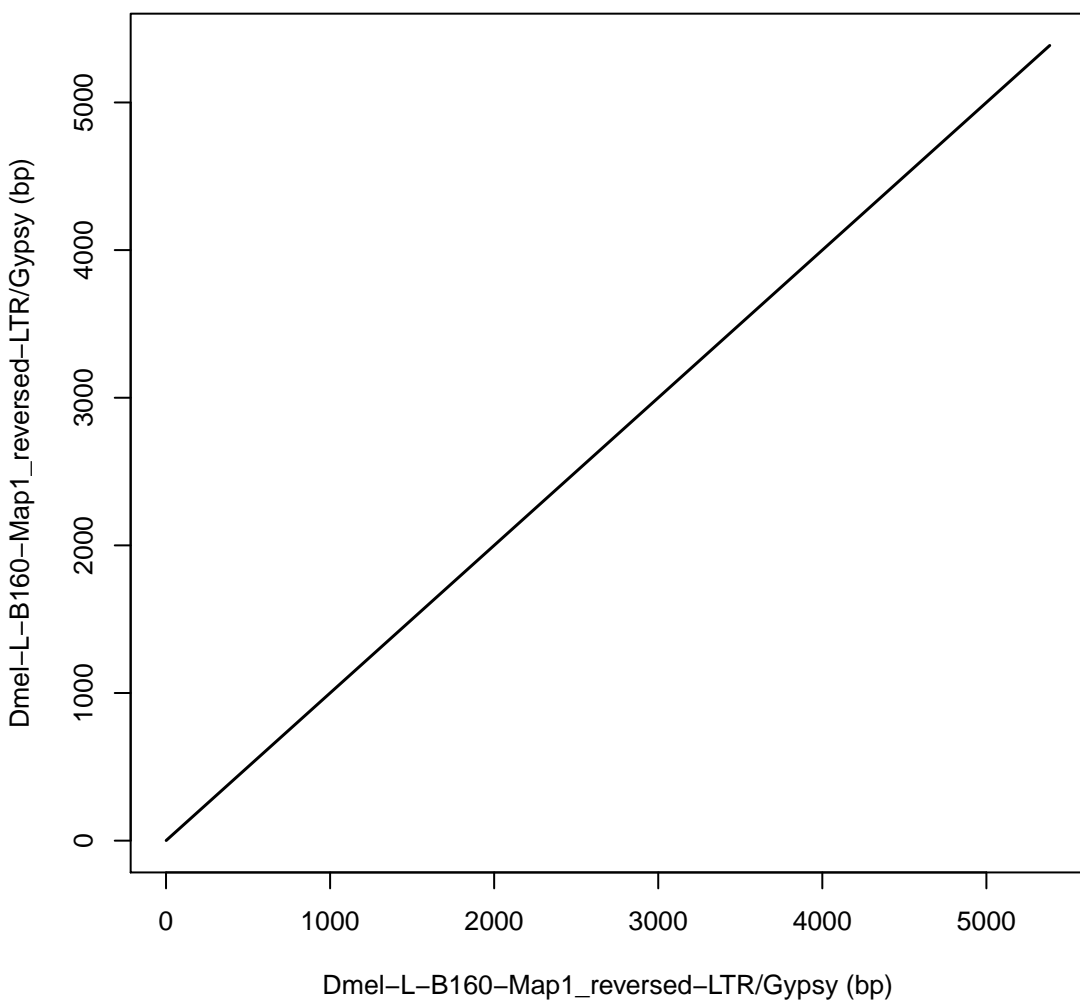
TE: Dmel-L-B160-Map1_reversed-LTR/Gypsy
consensus size: 5386bp; fragments: 558; full length: 0 (≥ 4847.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

