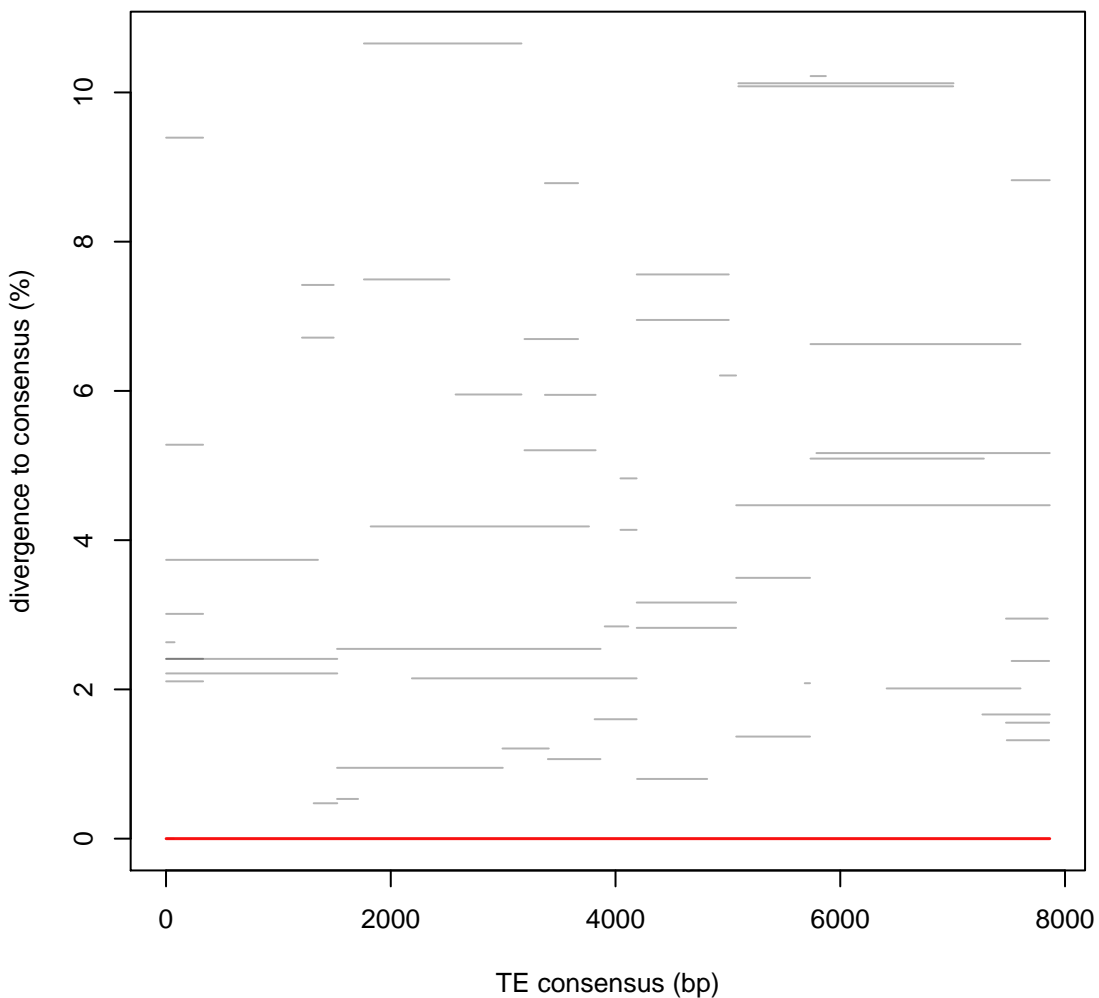
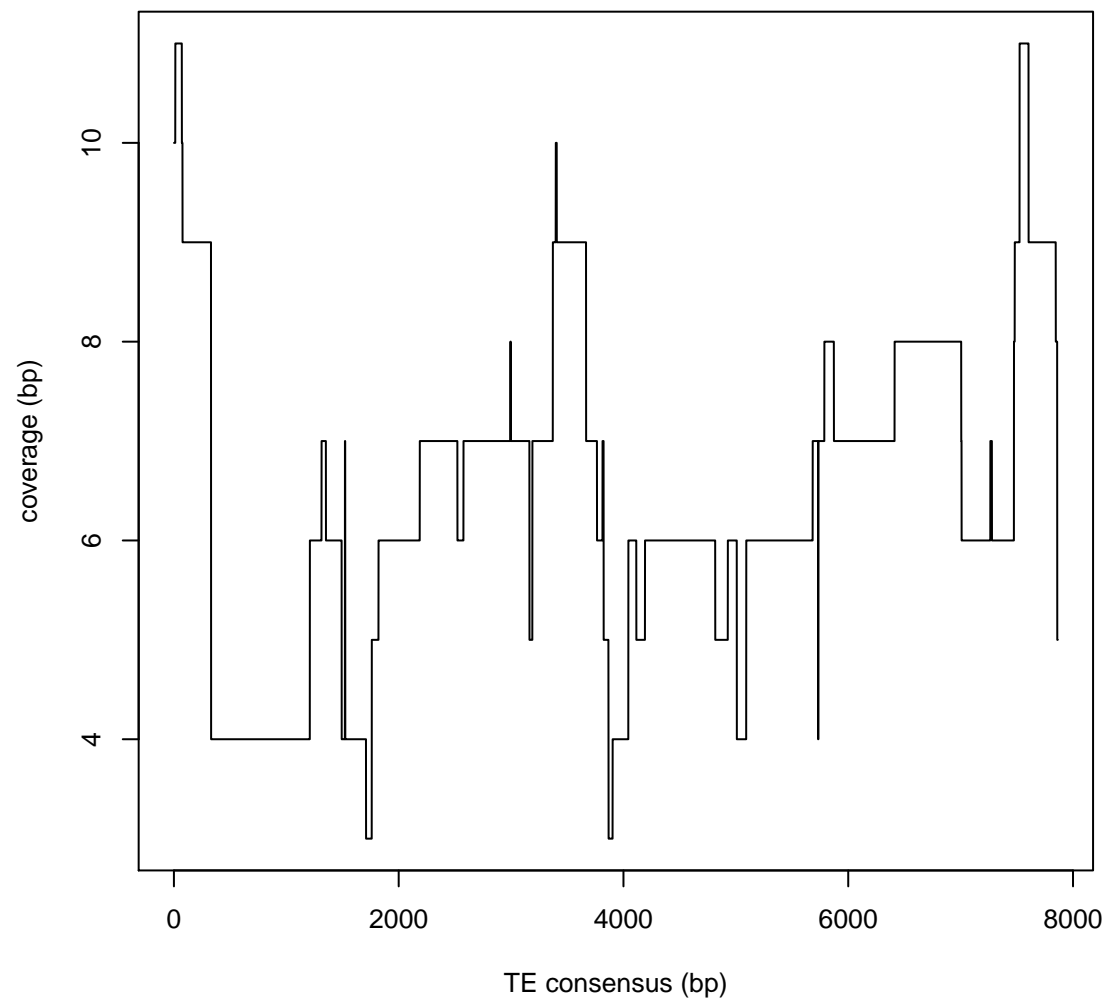


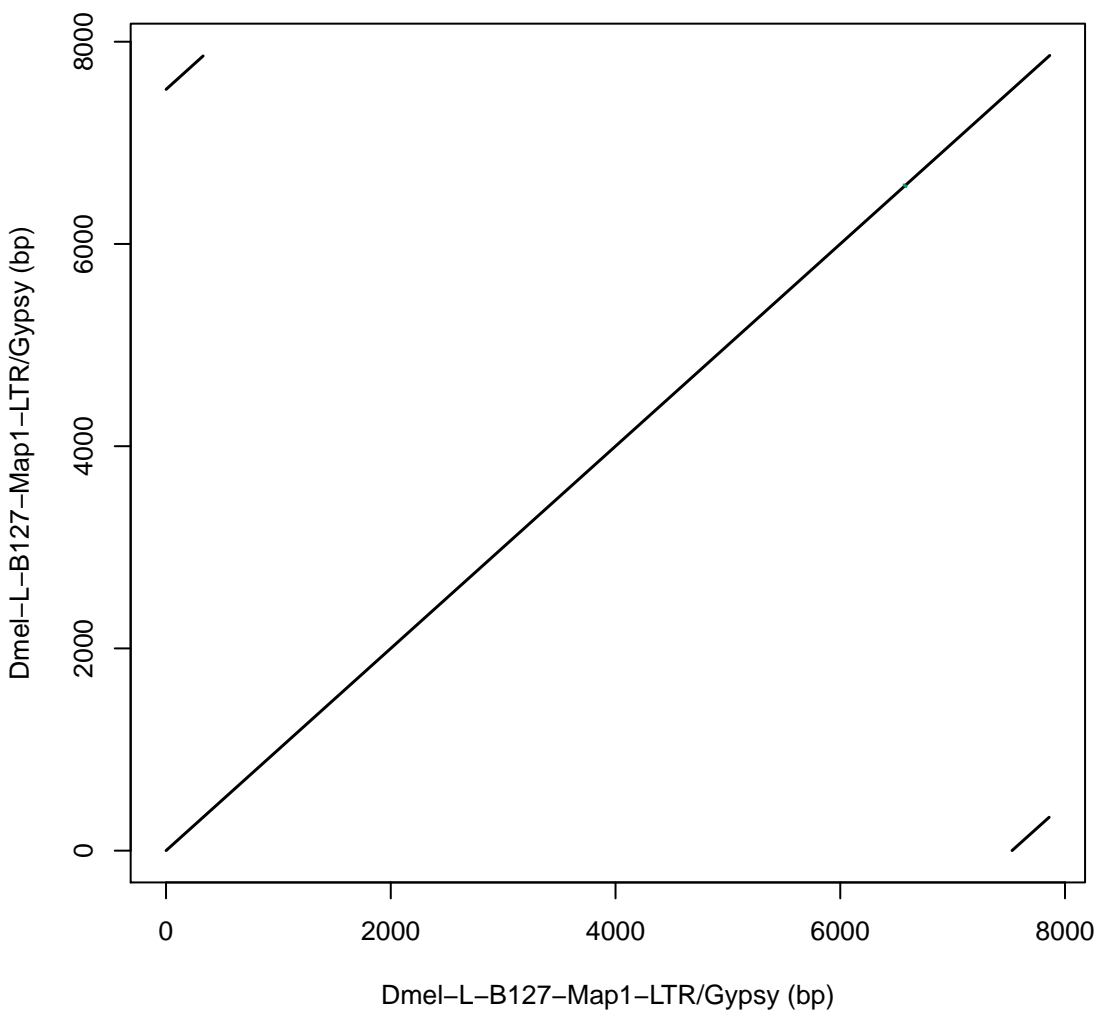
TE: Dmel-L-B127-Map1-LTR/Gypsy
consensus size: 7864bp; fragments: 55; full length: 1 (≥ 7077.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

