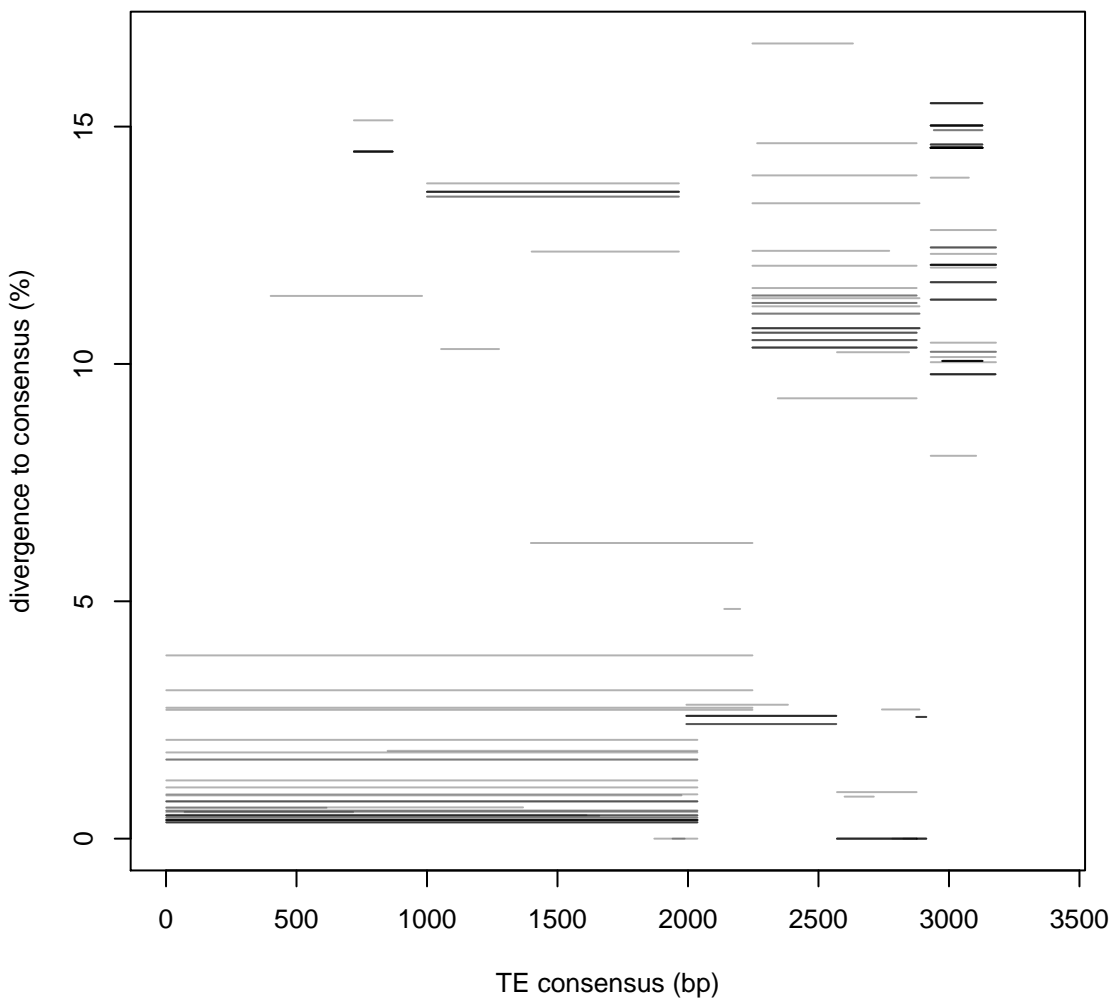
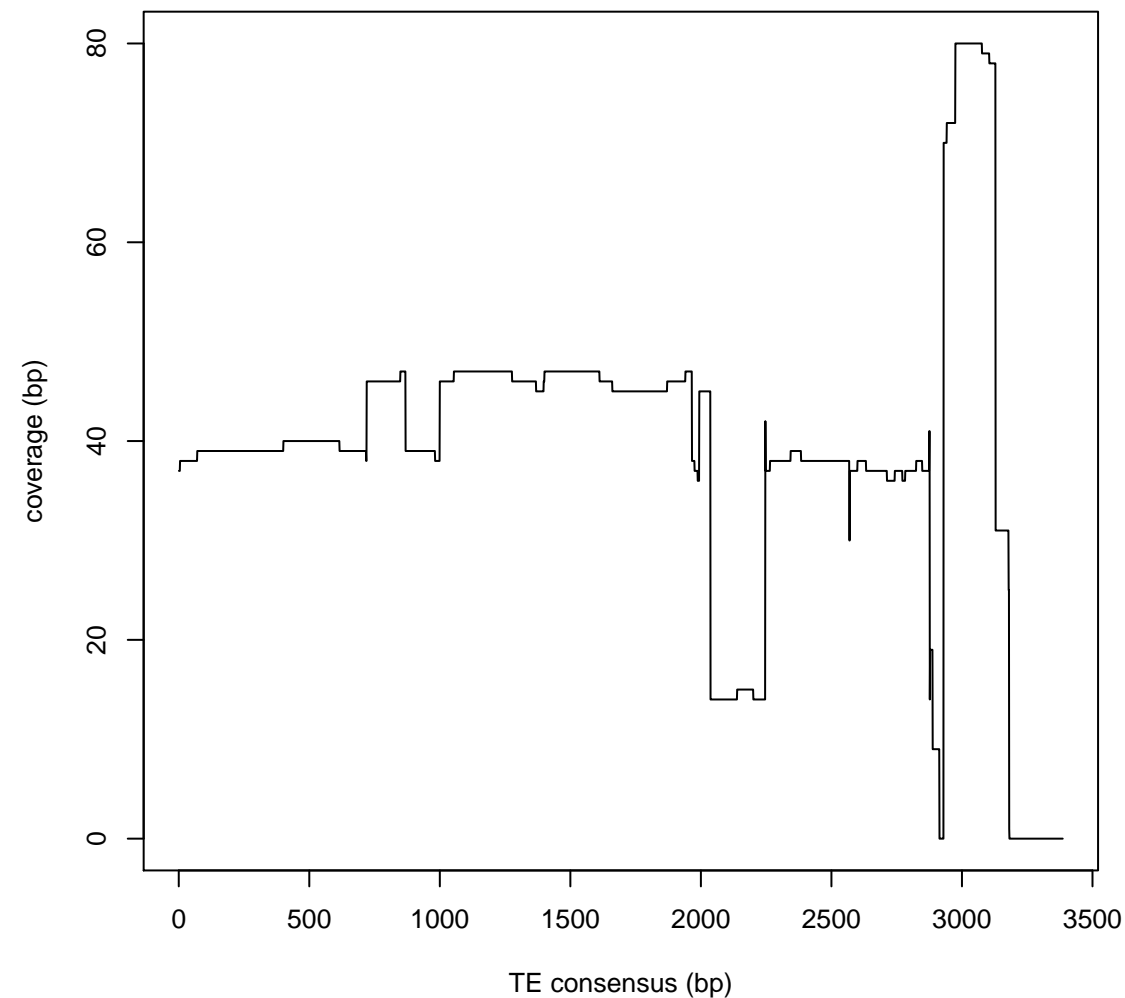


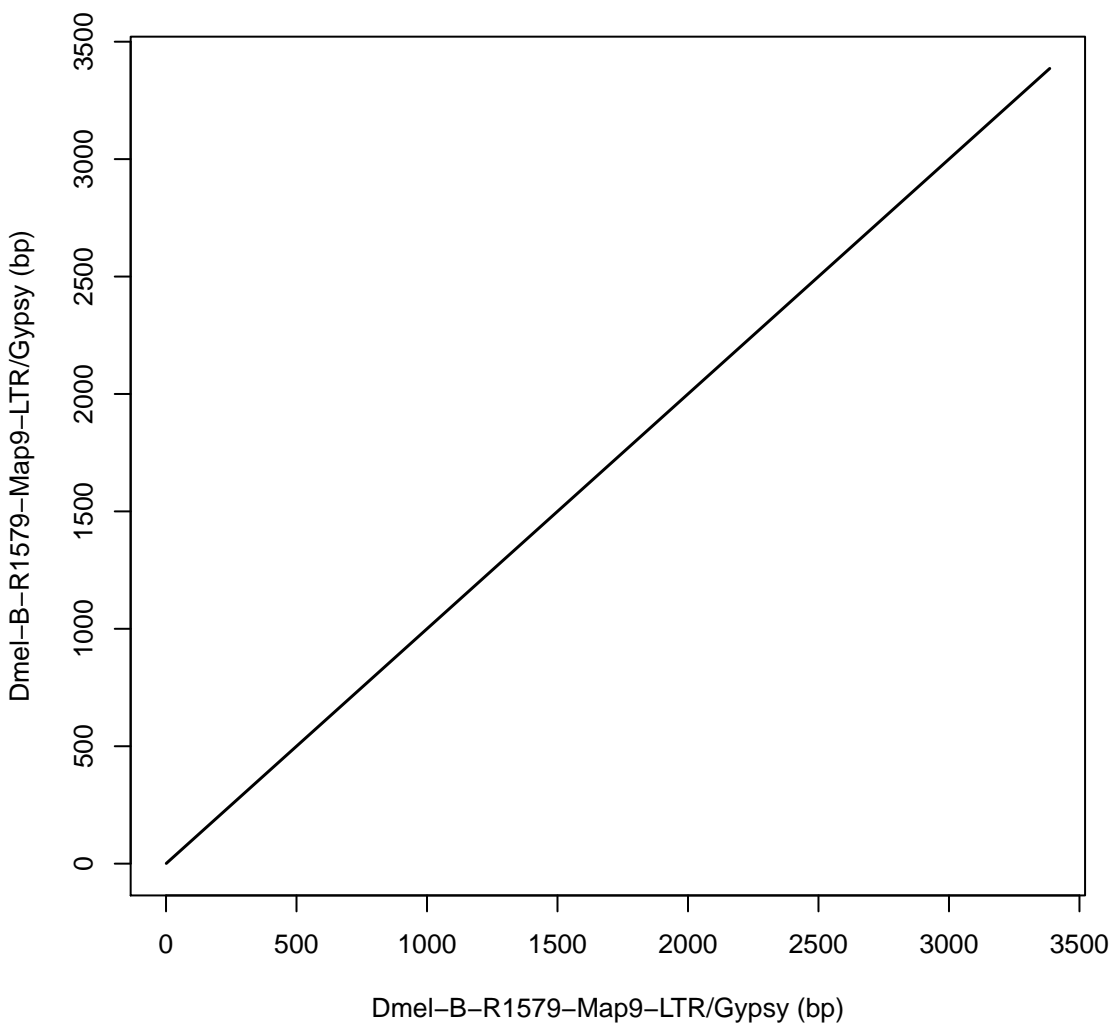
TE: Dmel-B-R1579-Map9-LTR/Gypsy
consensus size: 3386bp; fragments: 202; full length: 0 (≥ 3047.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

