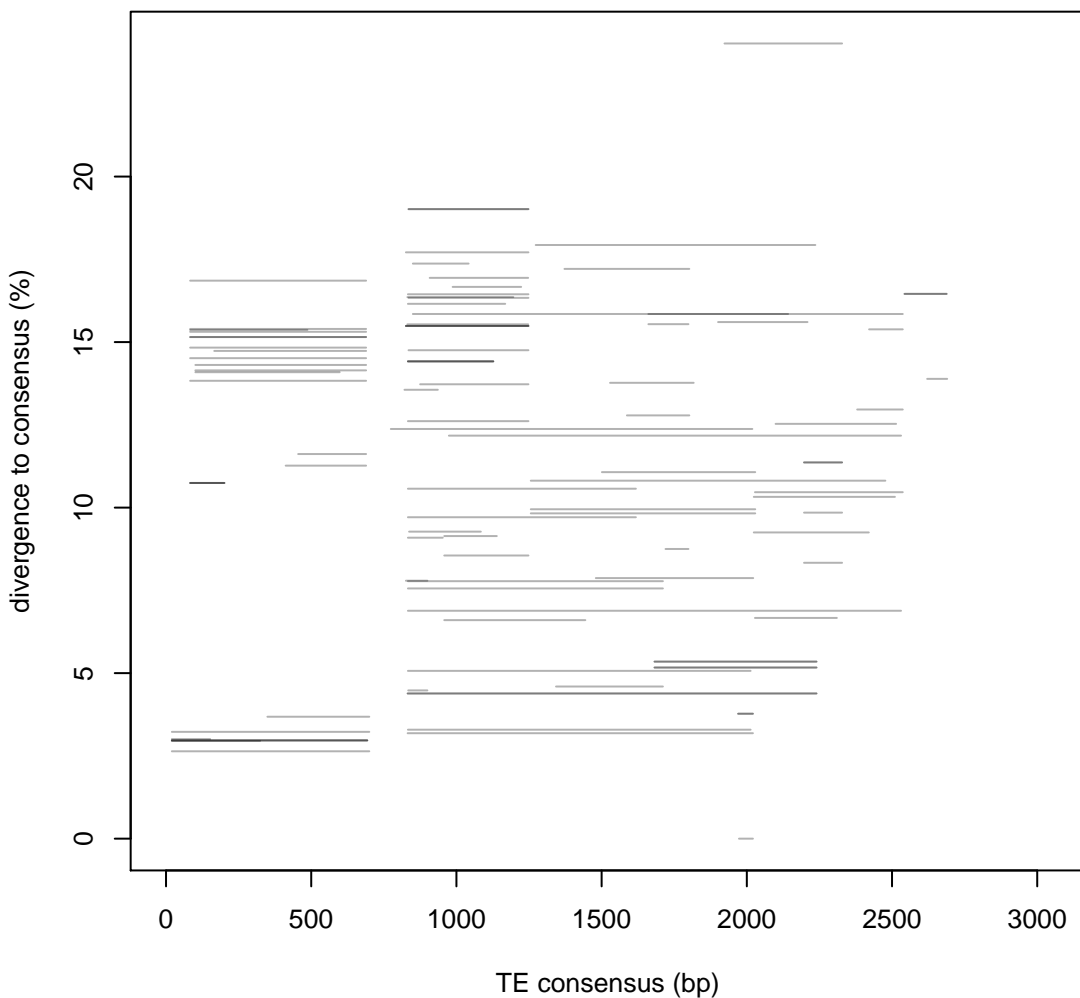
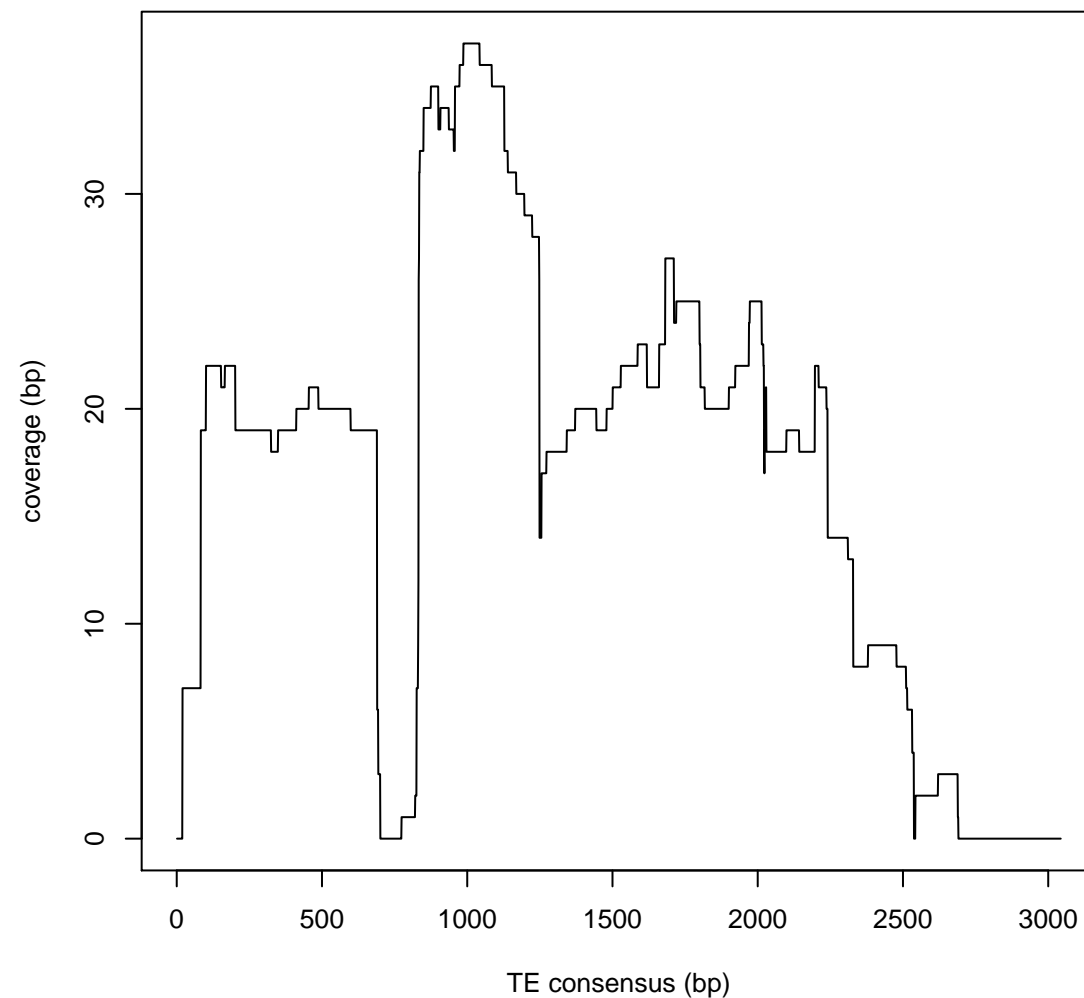


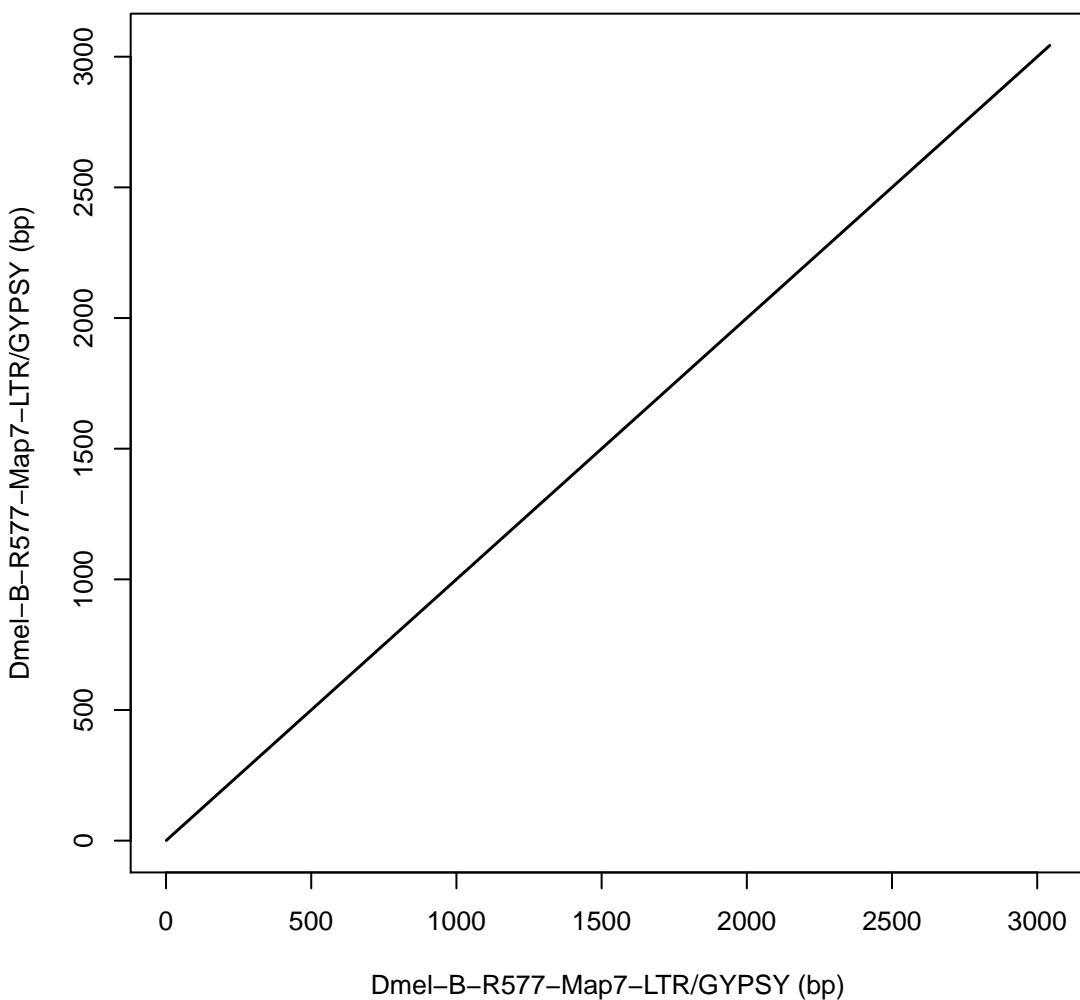
TE: Dmel-B-R577-Map7-LTR/GYPSY
consensus size: 3043bp; fragments: 103; full length: 0 (≥ 2738.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

