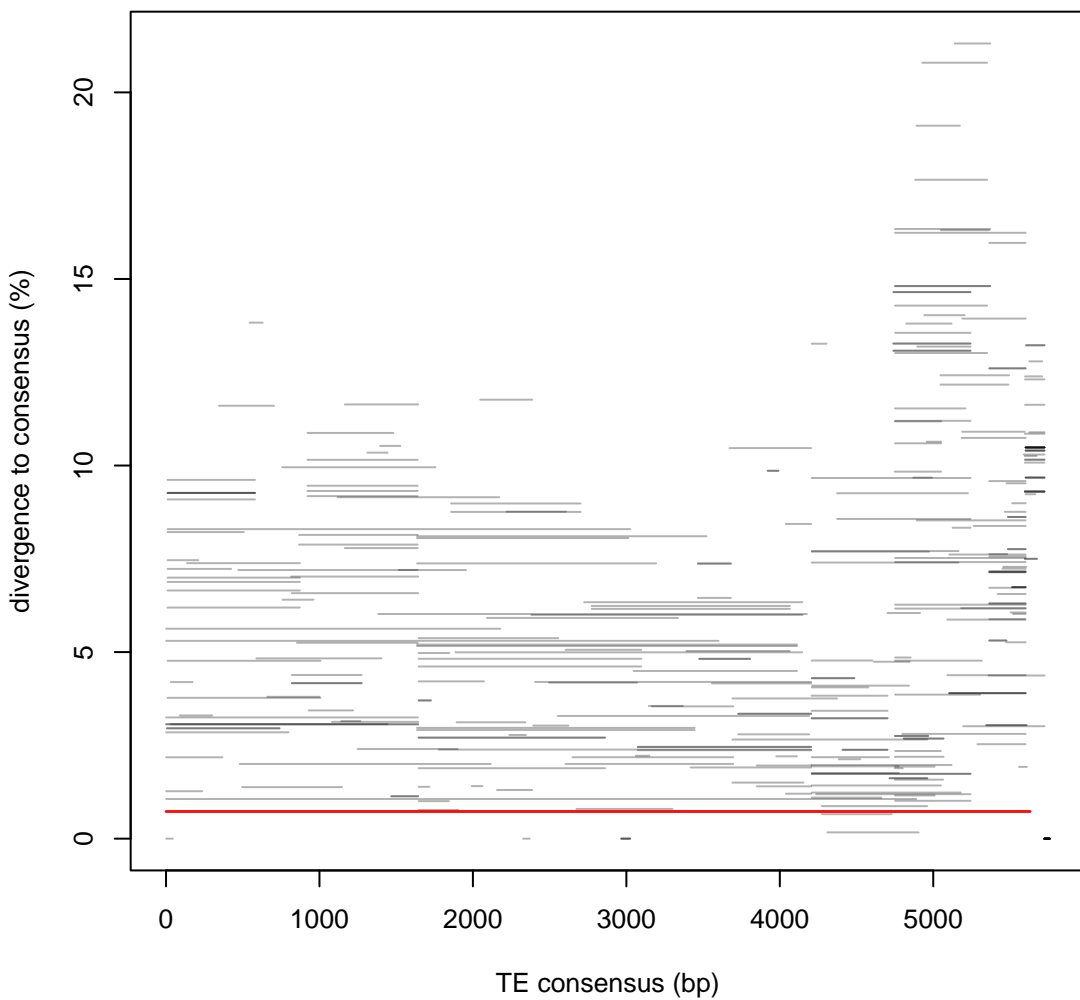
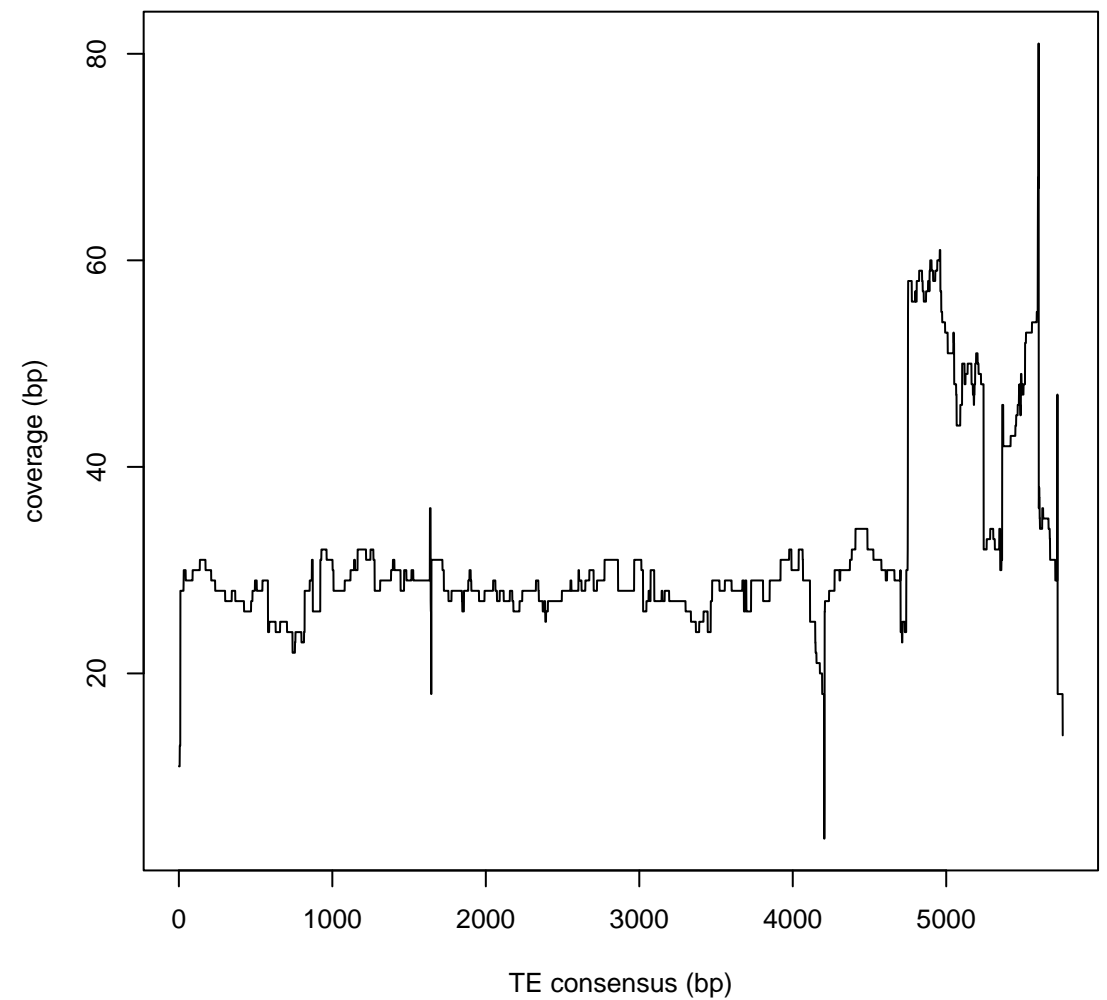


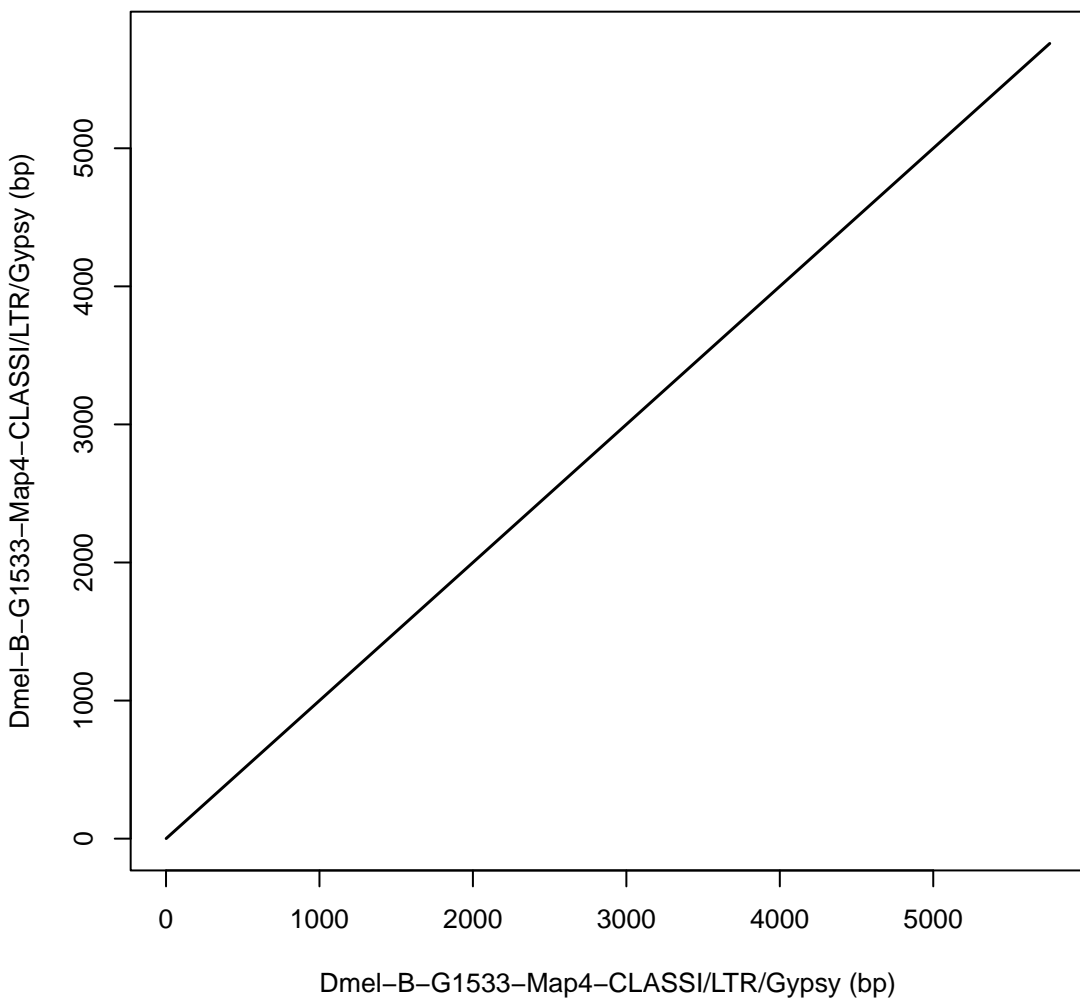
TE: Dmel-B-G1533-Map4-CLASSI/LTR/Gypsy
consensus size: 5759bp; fragments: 330; full length: 1 (>=5183.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

