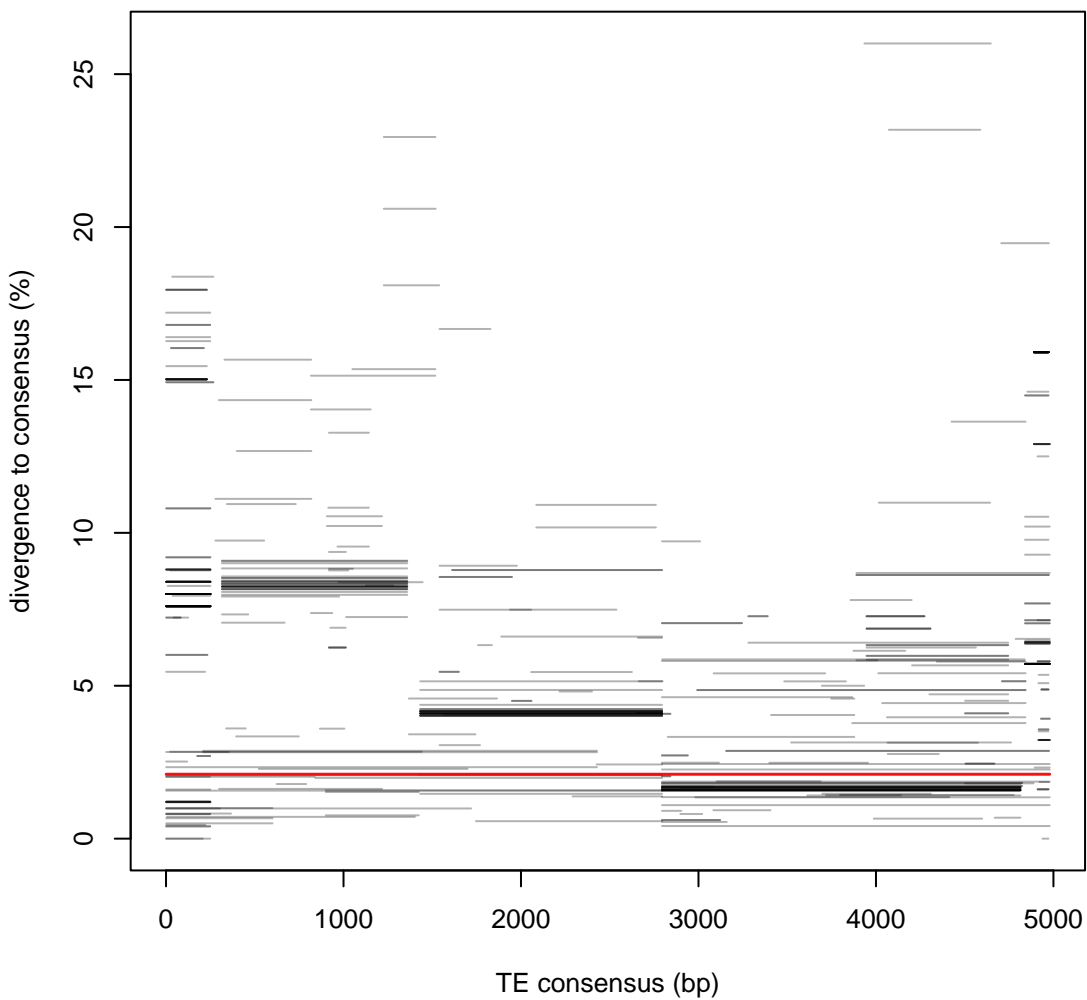
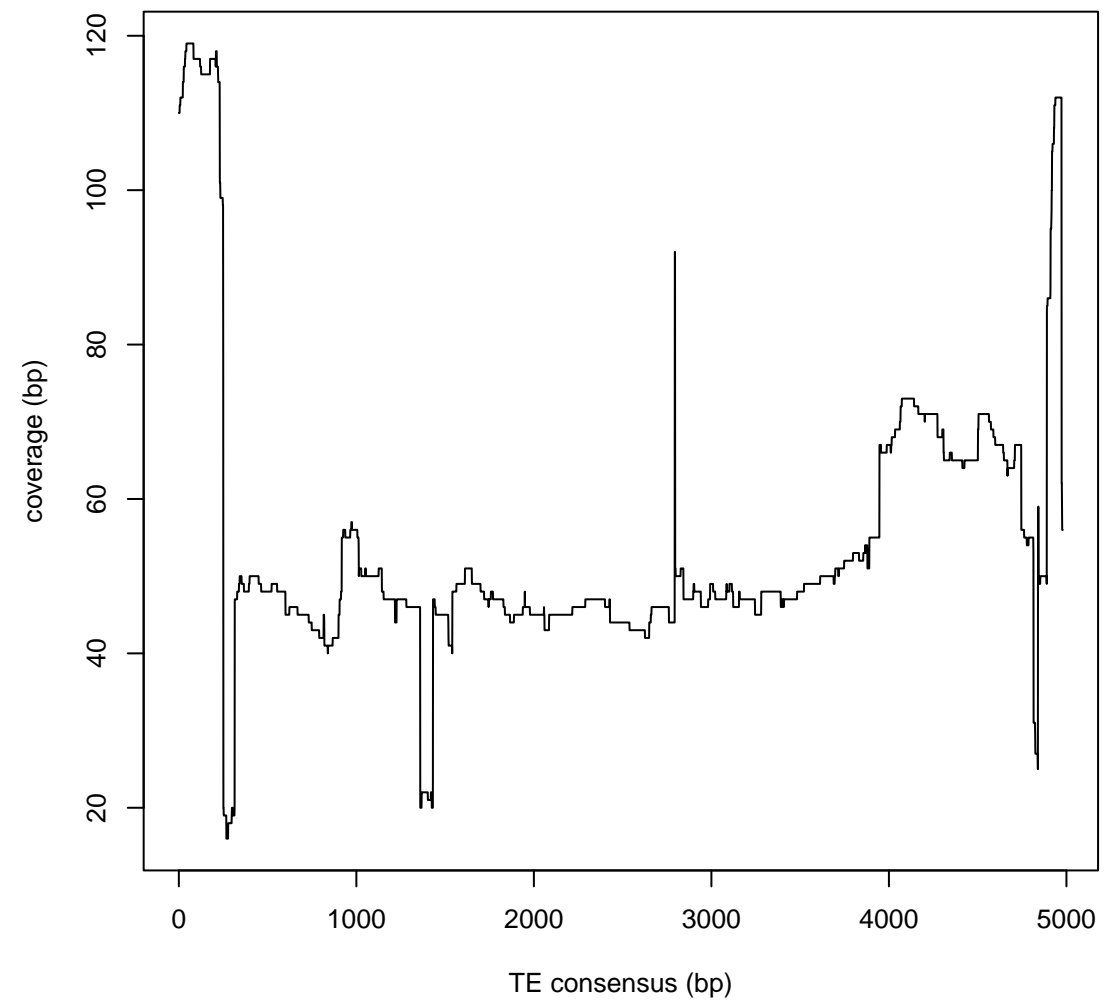


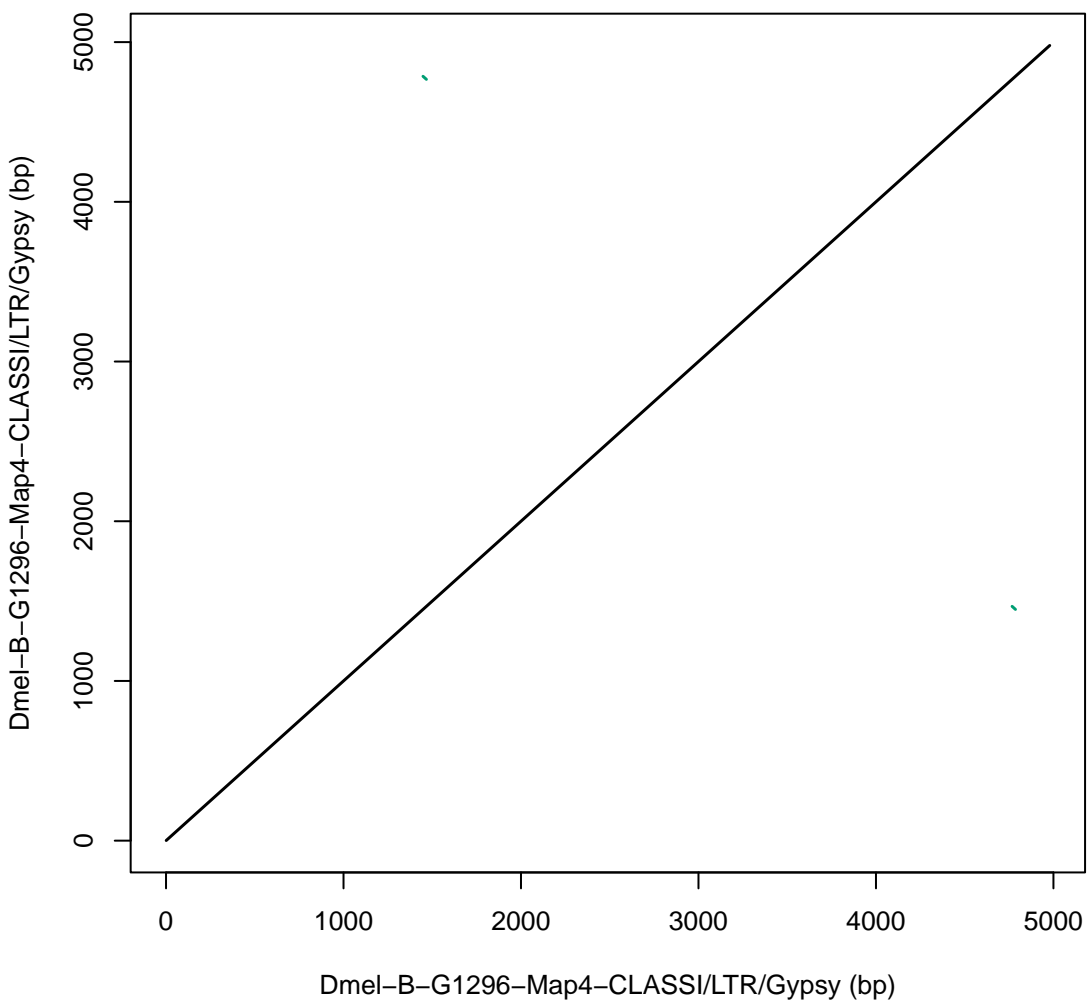
TE: Dmel-B-G1296-Map4-CLASSI/LTR/Gypsy
consensus size: 4979bp; fragments: 460; full length: 1 (>=4481.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

