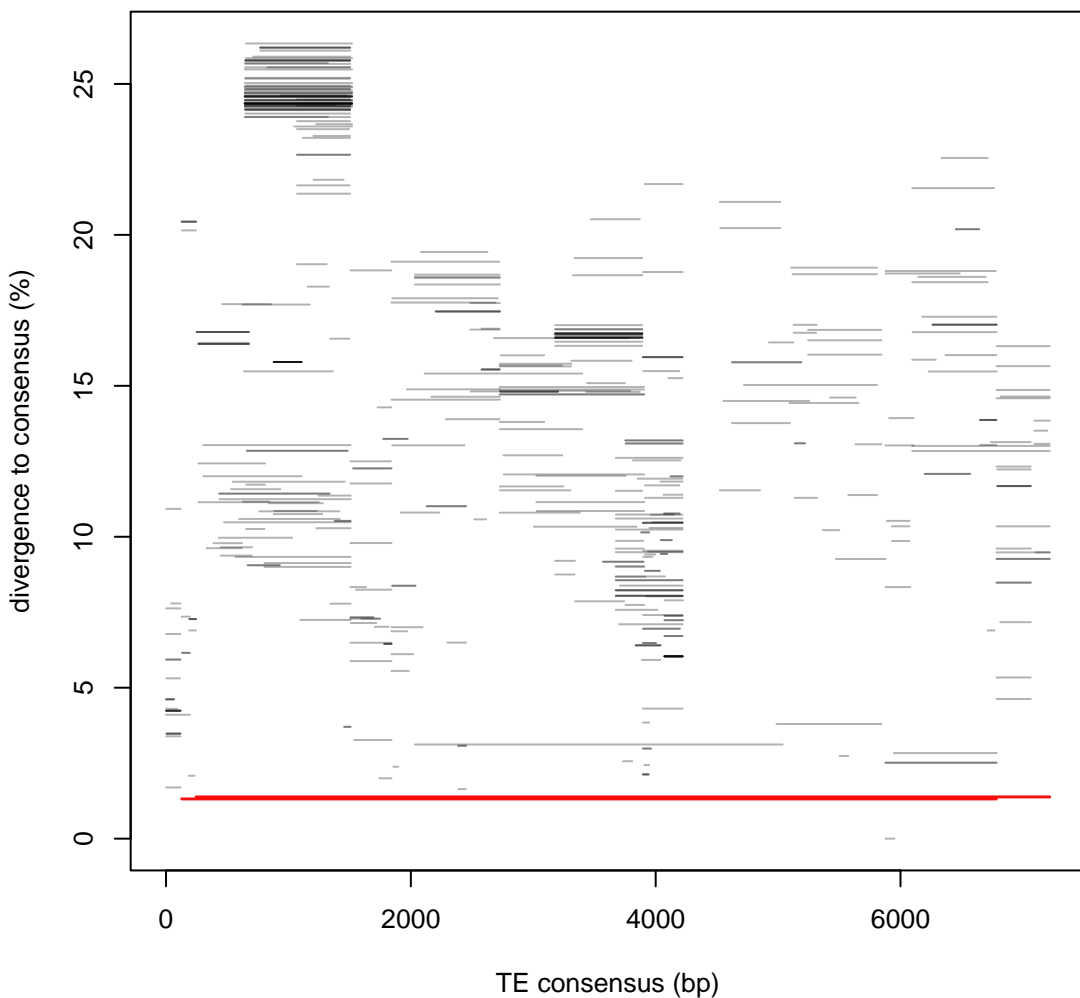
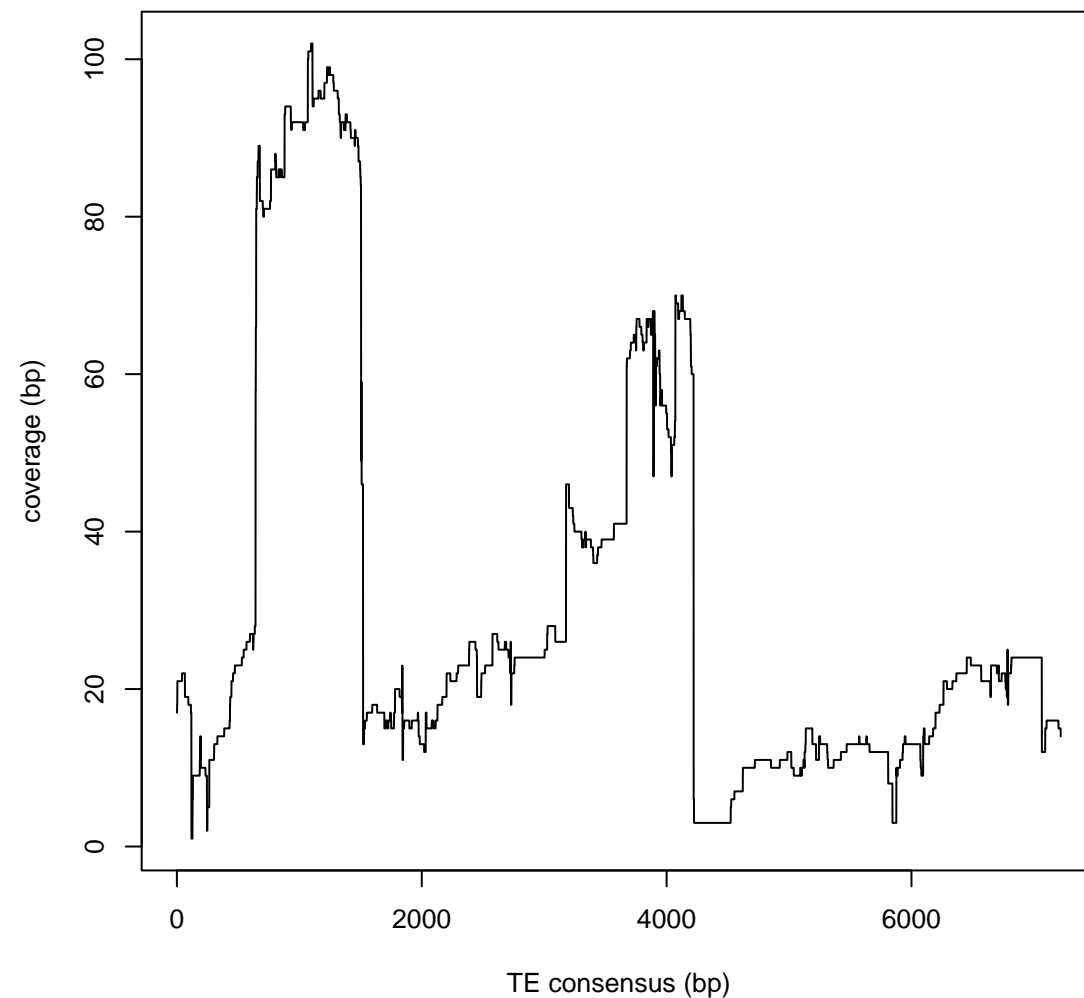


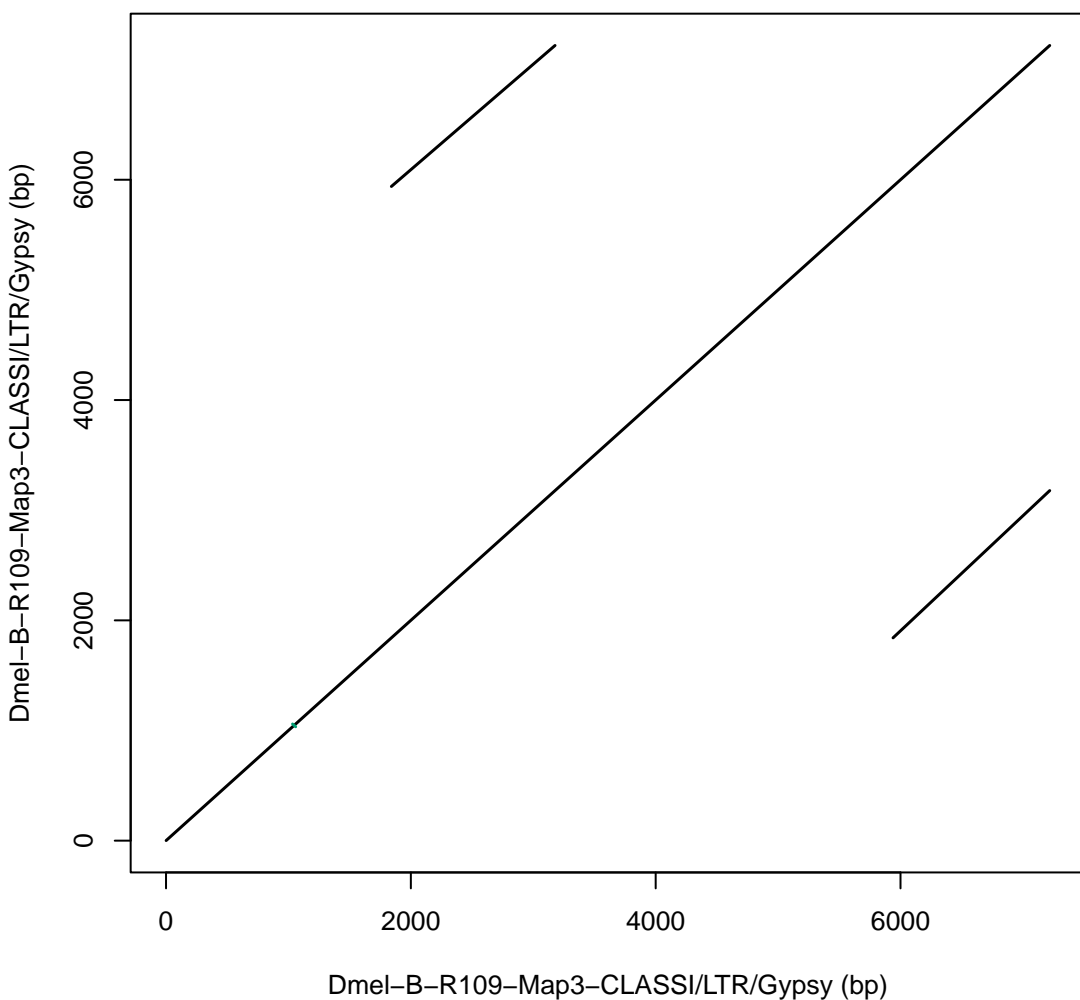
TE: Dmel-B-R109-Map3-CLASSI/LTR/Gypsy
consensus size: 7219bp; fragments: 471; full length: 2 (>=6497.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

