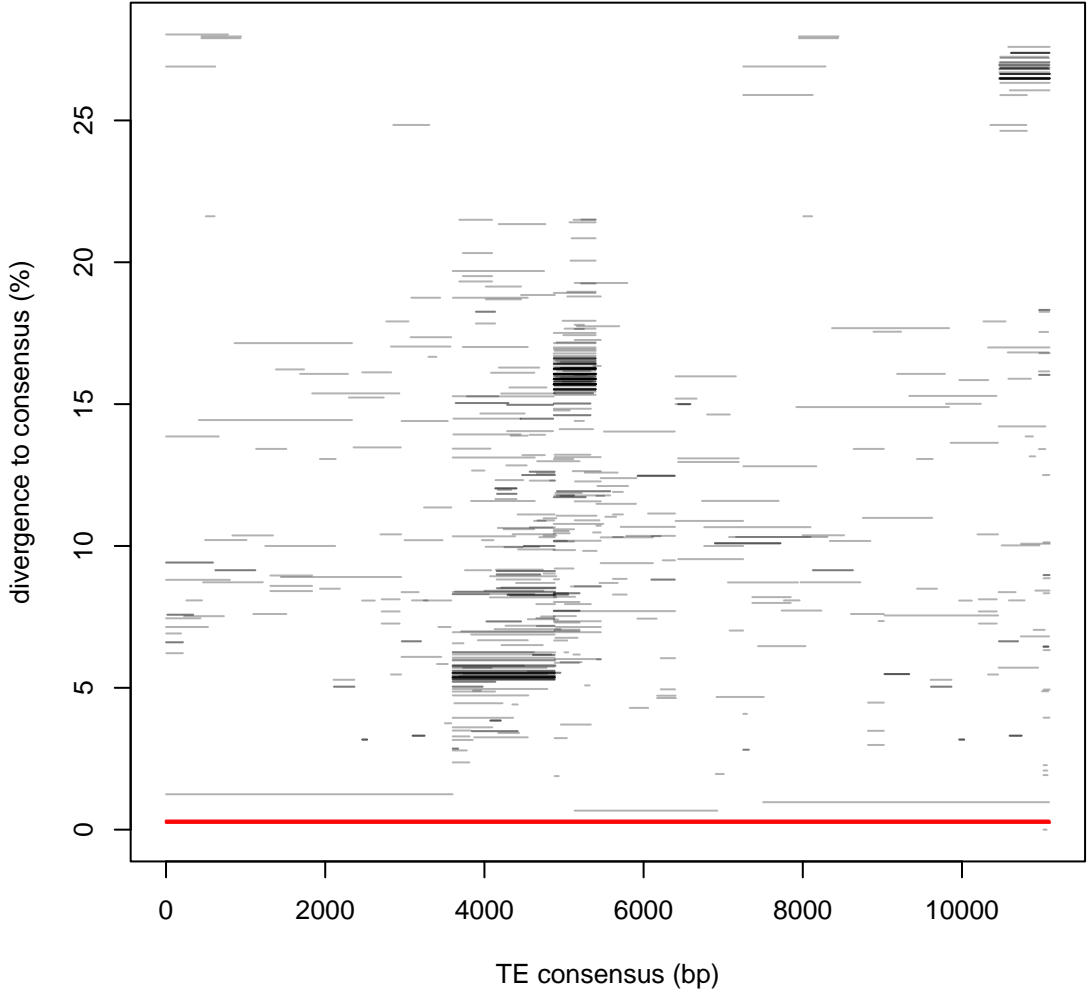
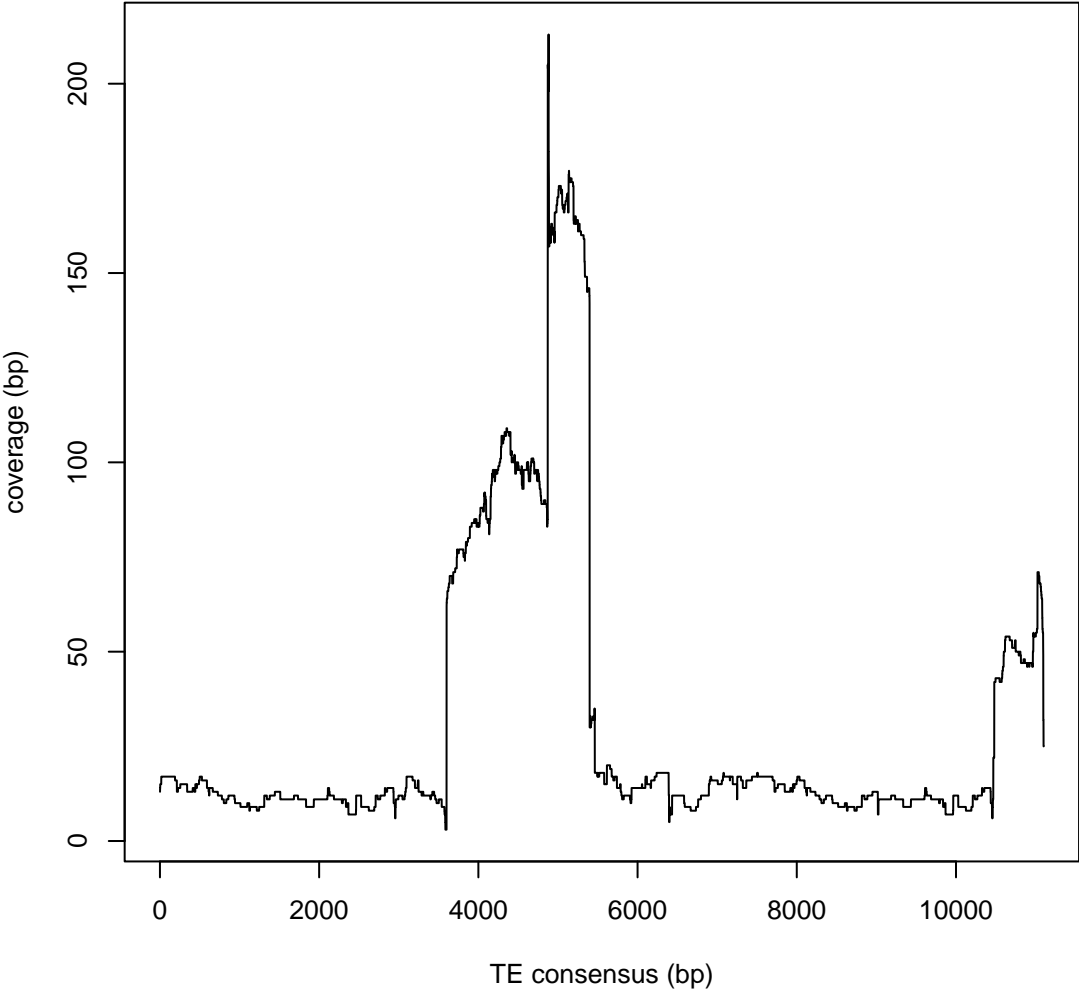


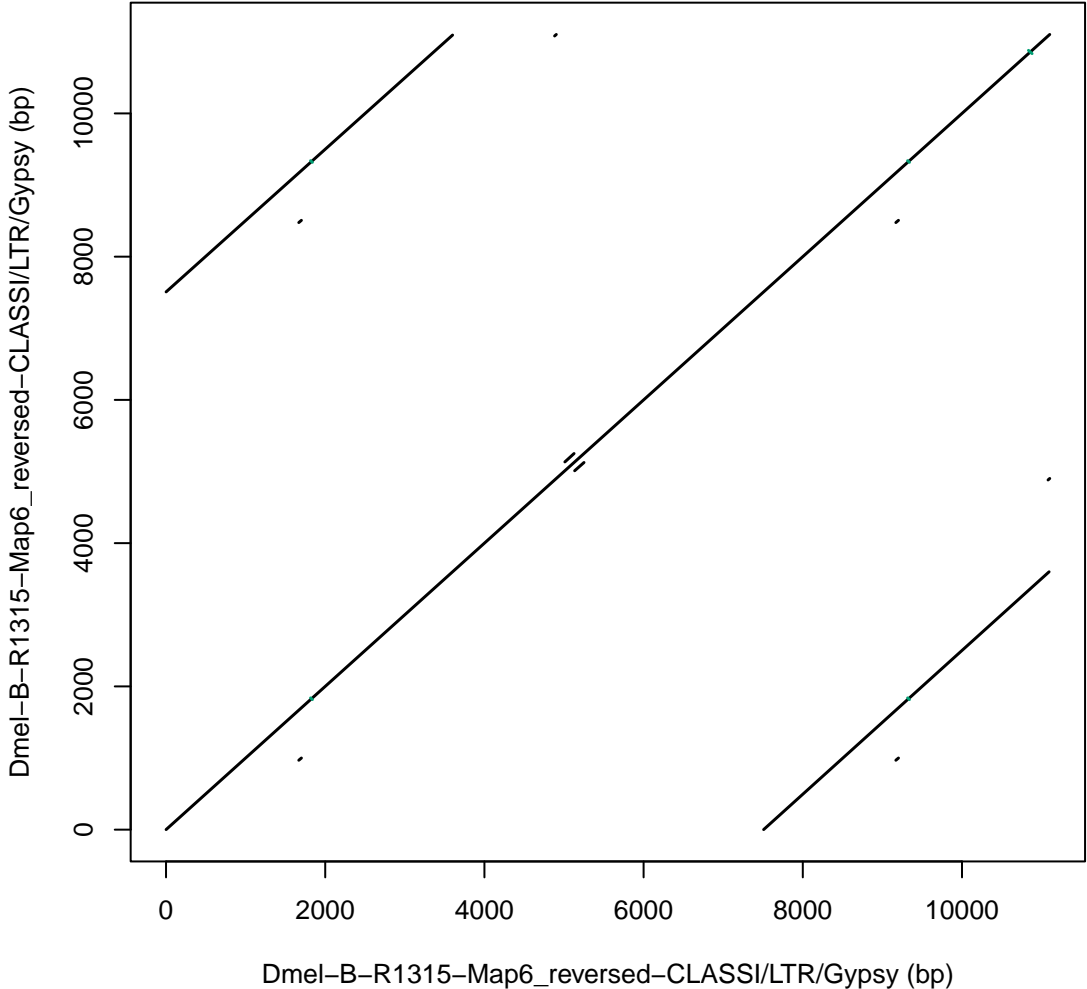
TE: Dmel-B-R1315-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 11102bp; fragments: 644; full length: 2 (>=9991.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

