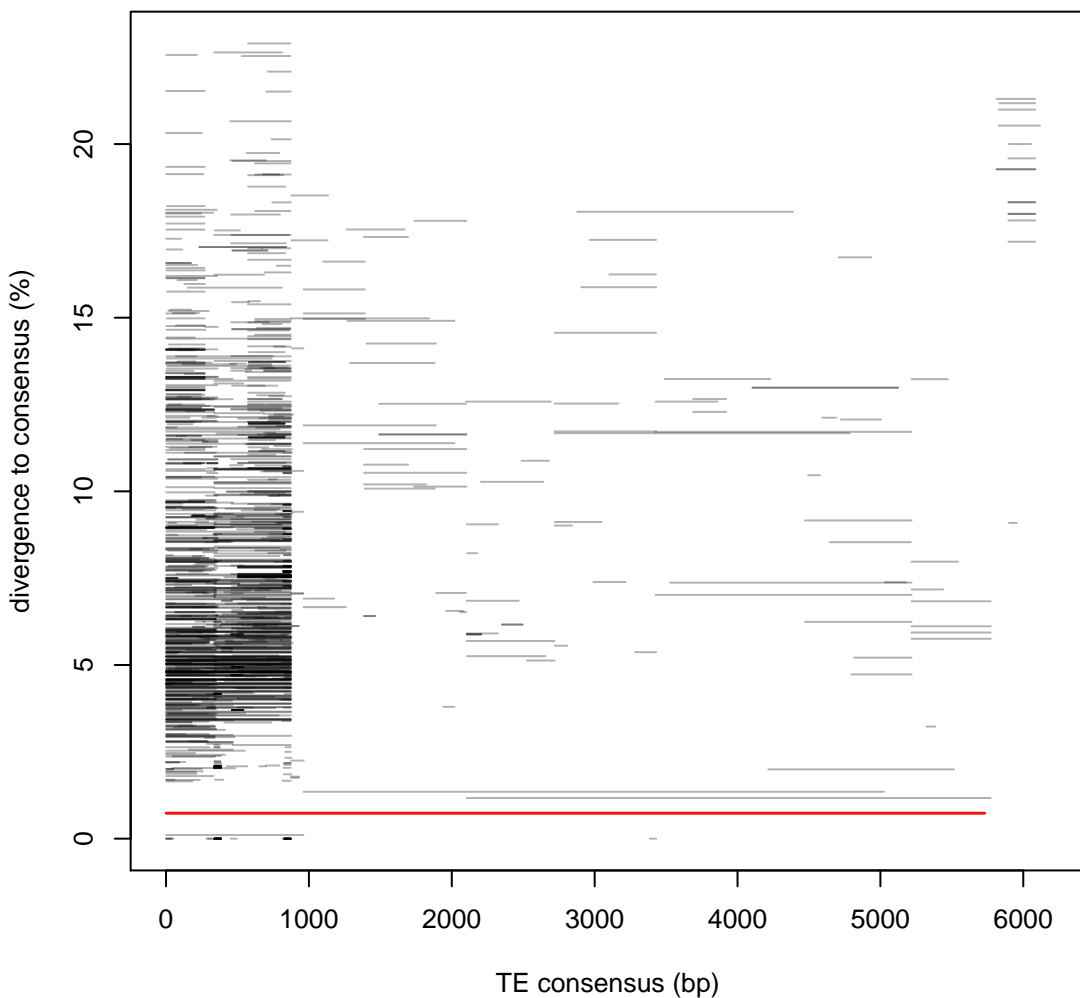
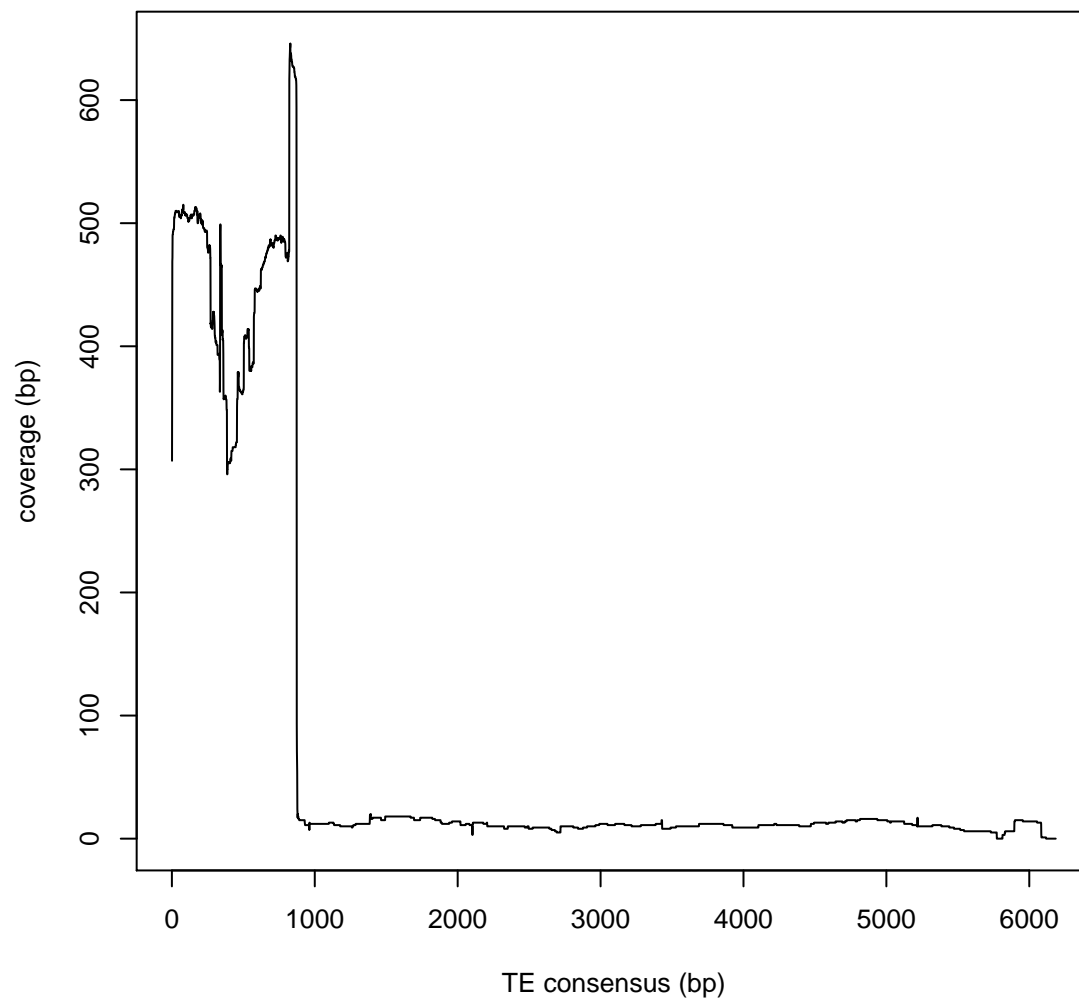


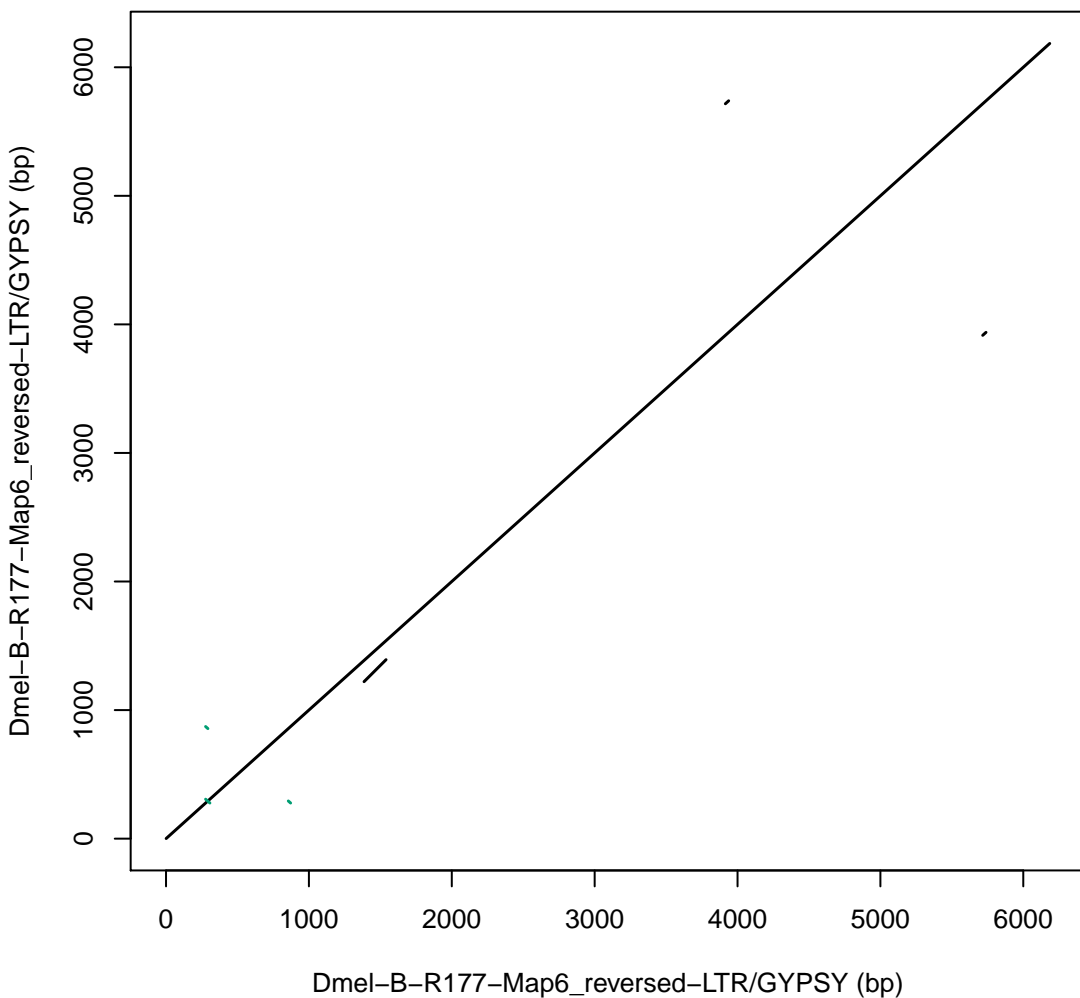
TE: Dmel-B-R177-Map6_reversed-LTR/GYPSY
consensus size: 6185bp; fragments: 1409; full length: 1 (>=5566.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

