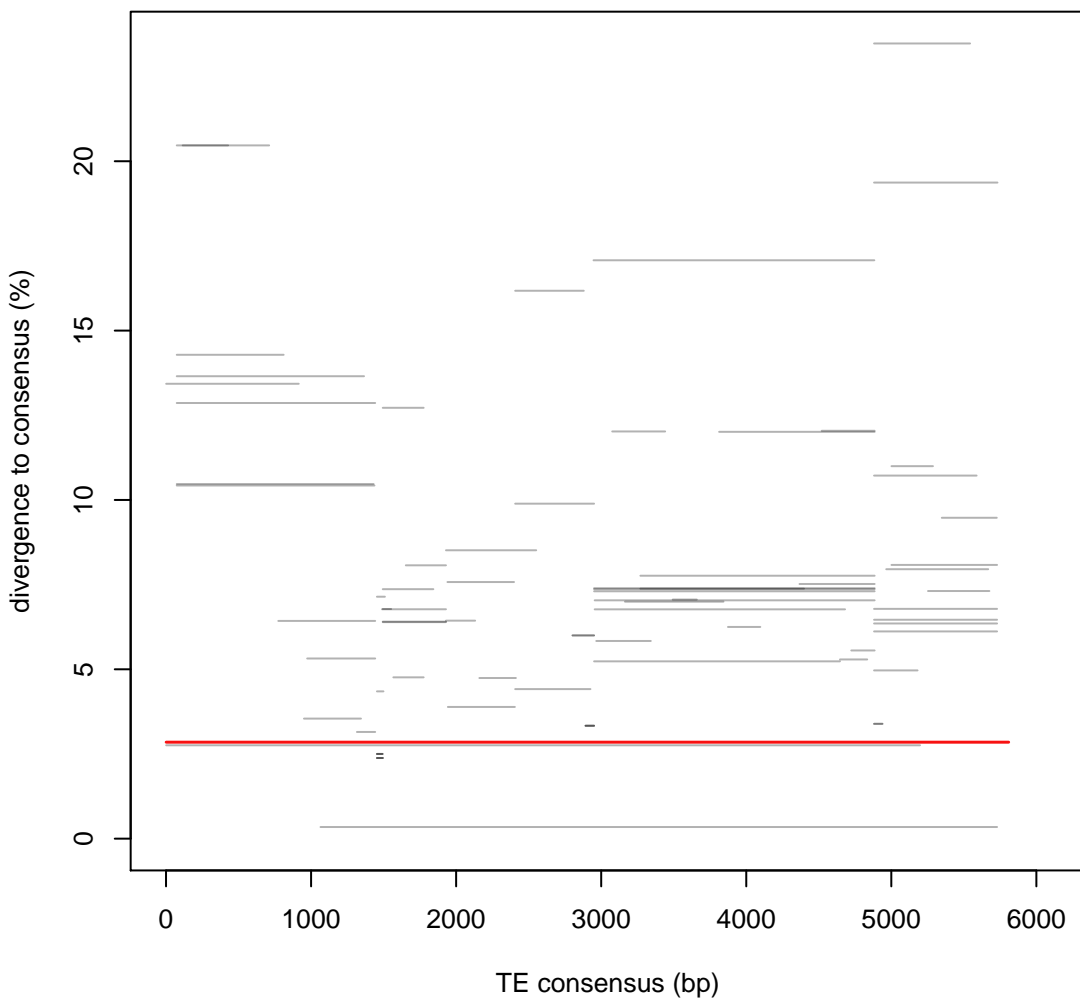
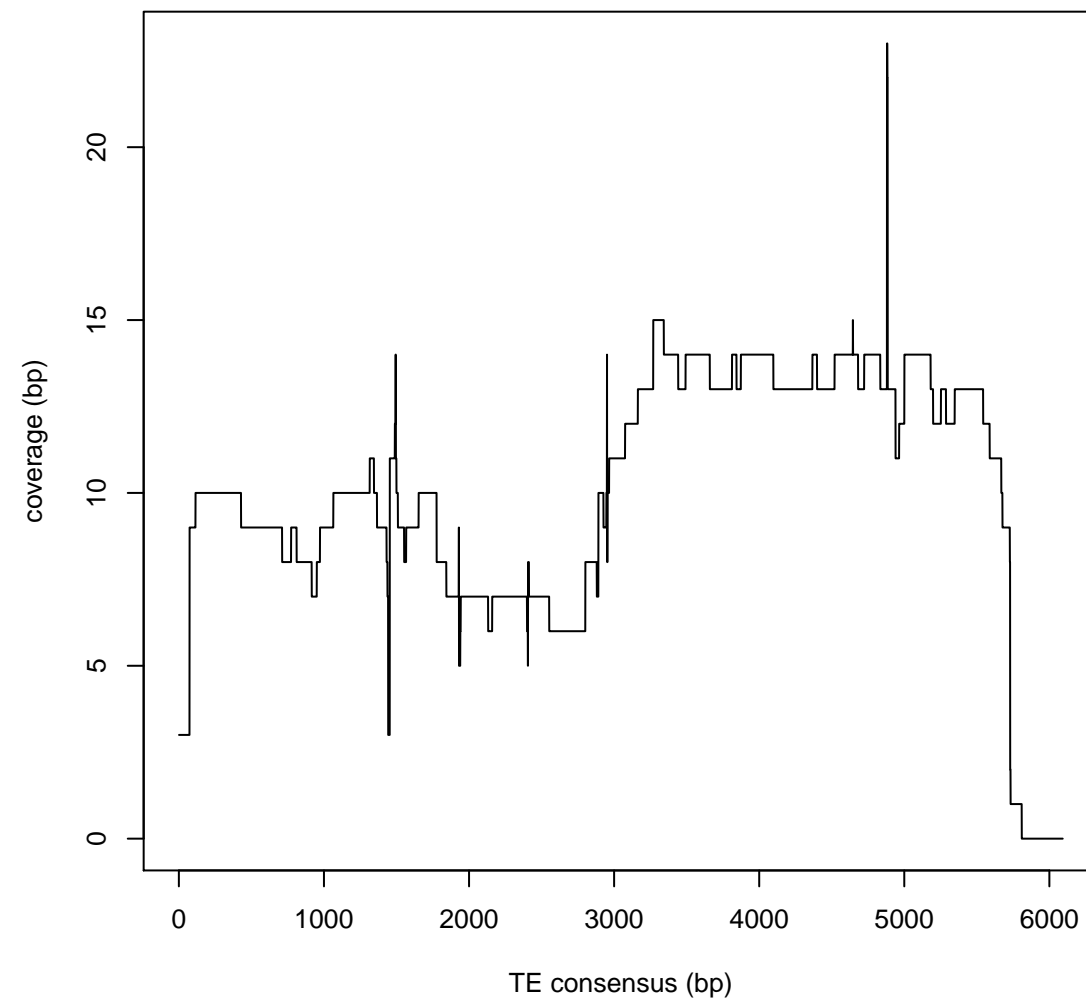


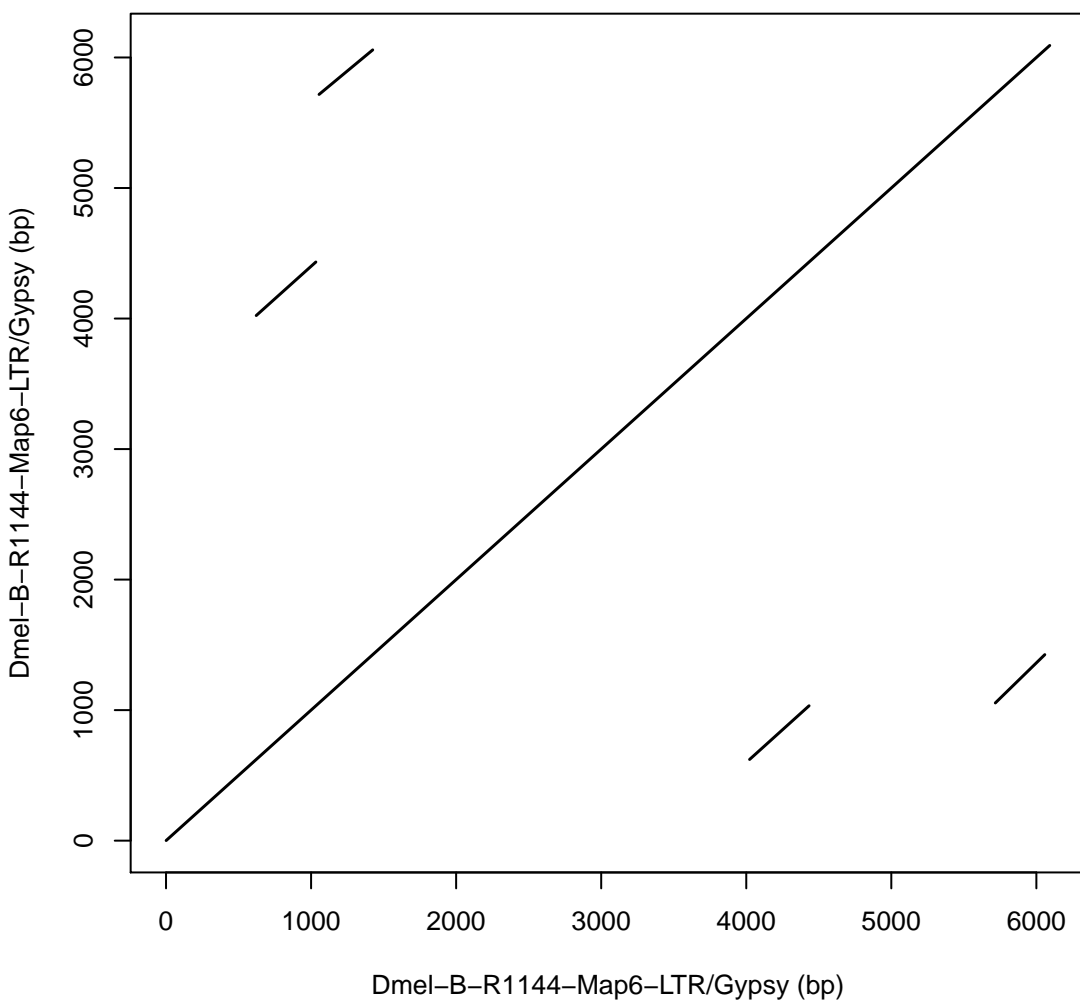
TE: Dmel-B-R1144-Map6-LTR/Gypsy  
consensus size: 6092bp; fragments: 78; full length: 1 (>=5482.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

