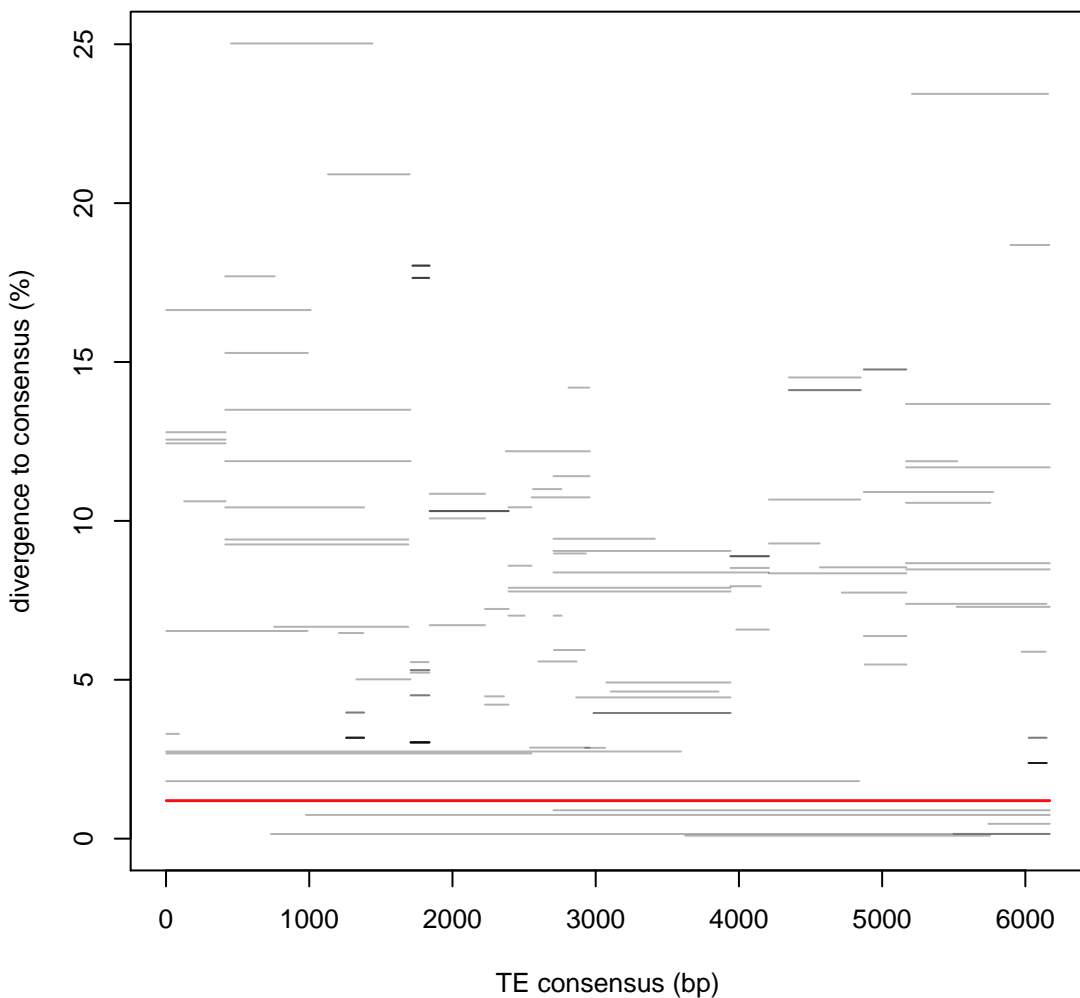
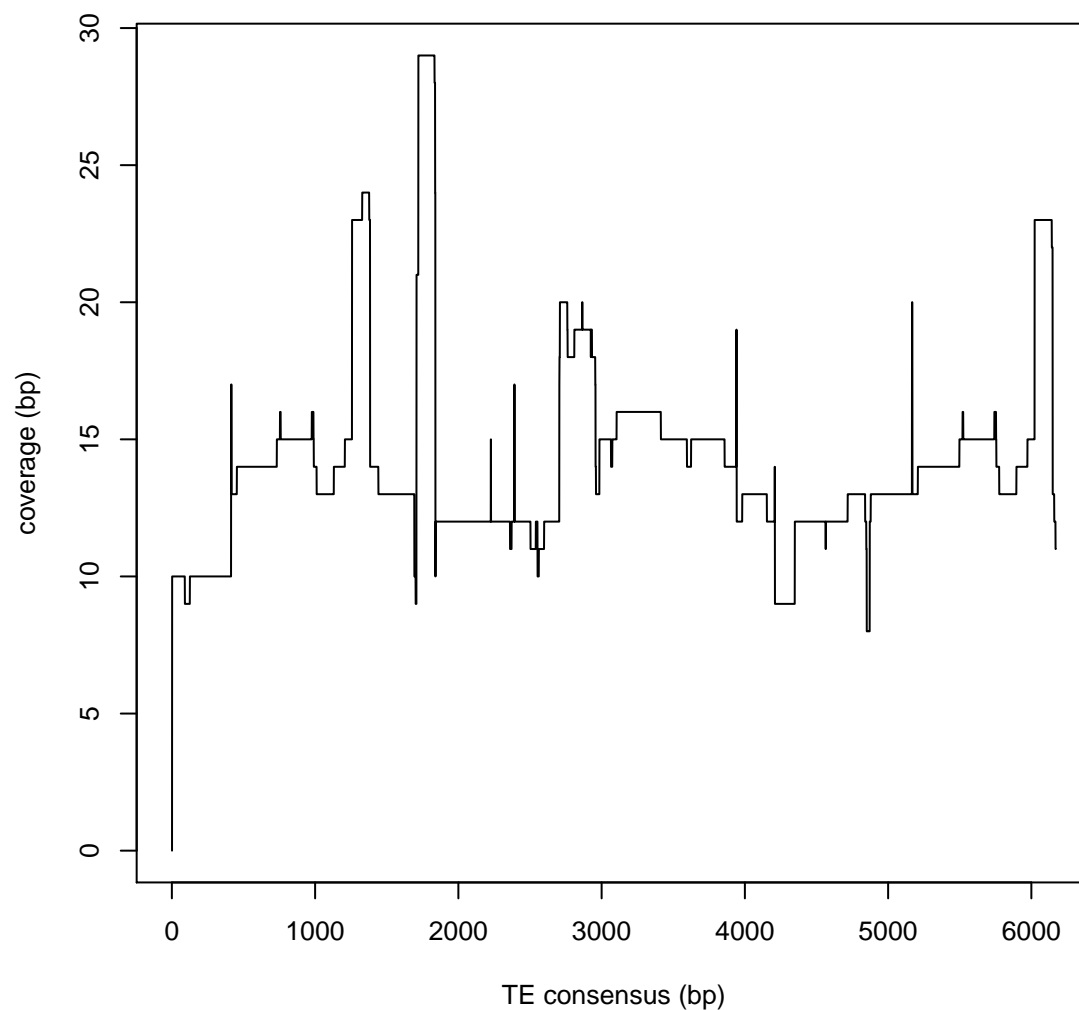


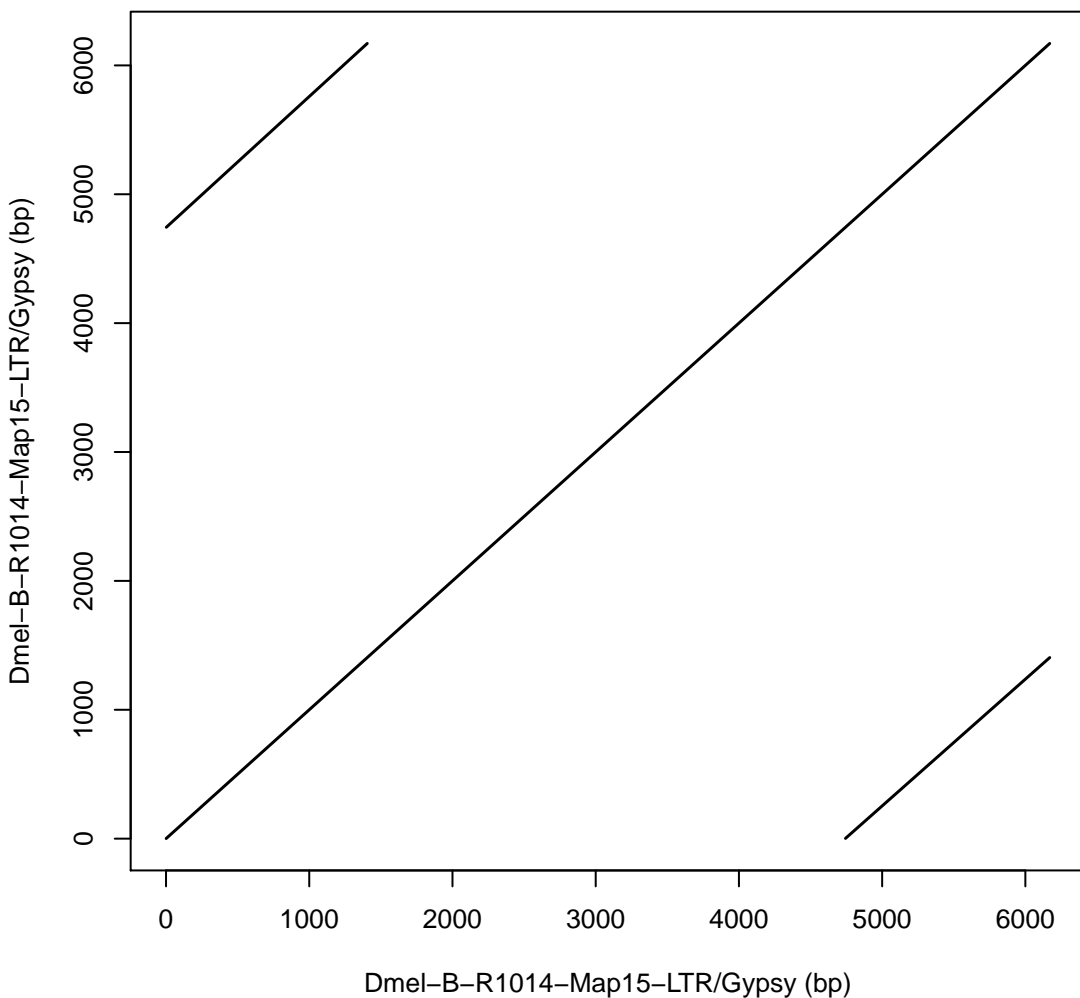
TE: Dmel-B-R1014-Map15-LTR/Gypsy  
consensus size: 6170bp; fragments: 131; full length: 1 (>=5553bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

