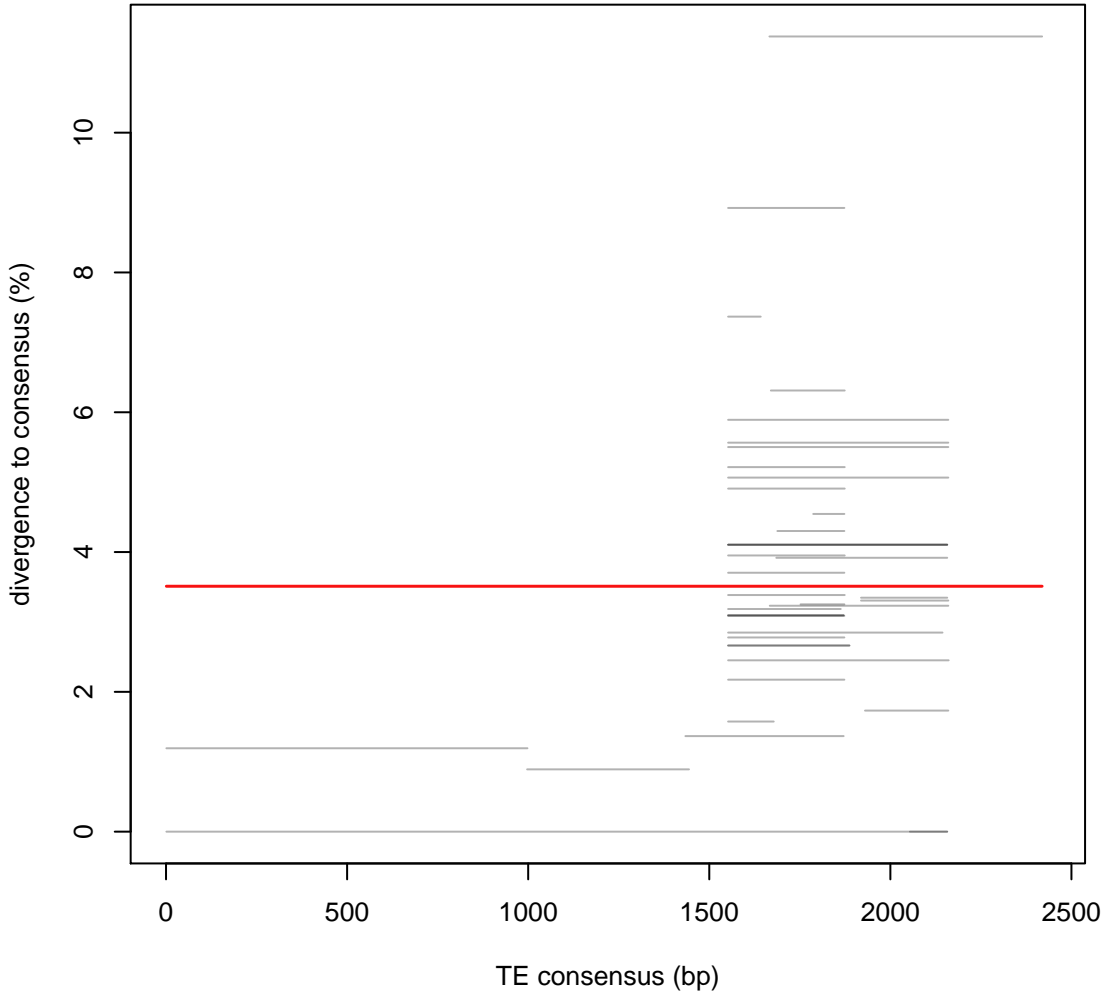
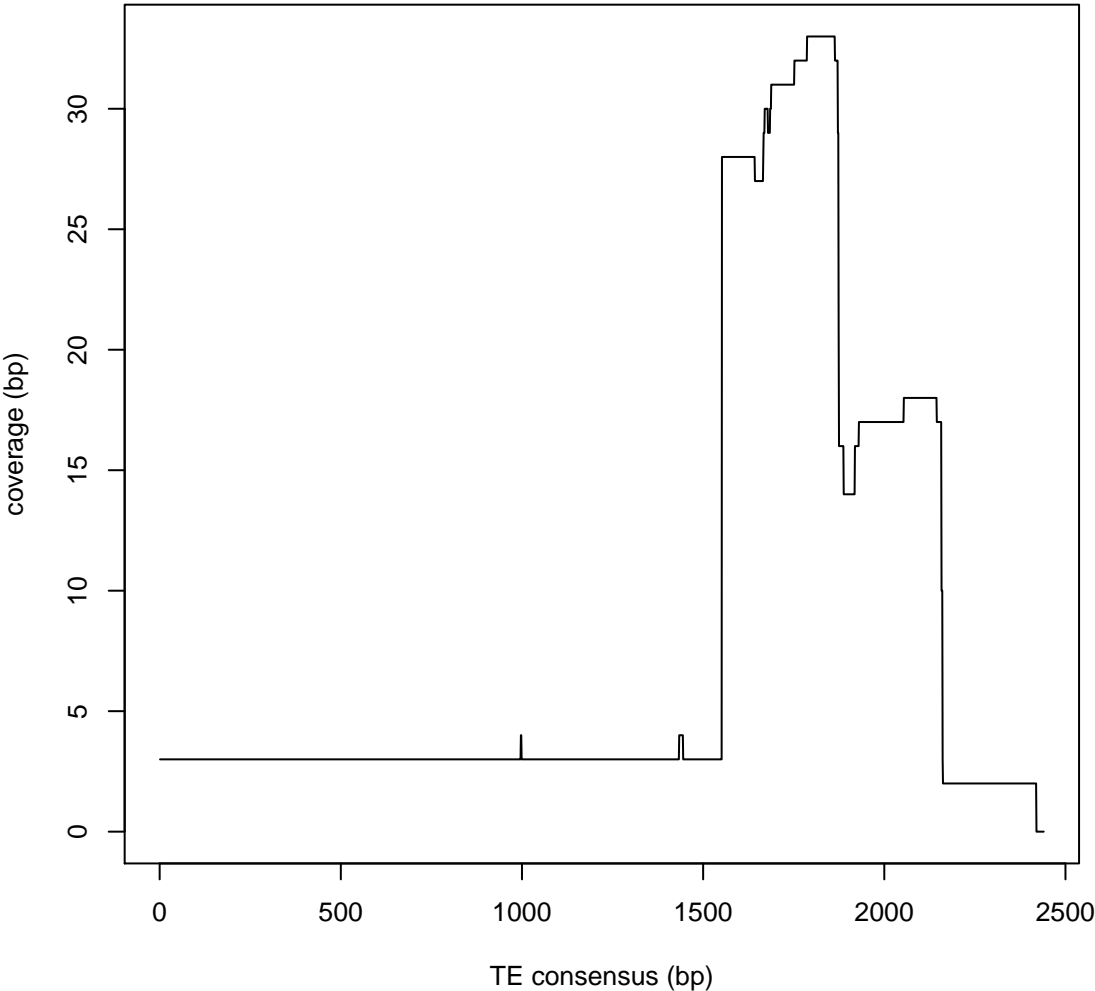


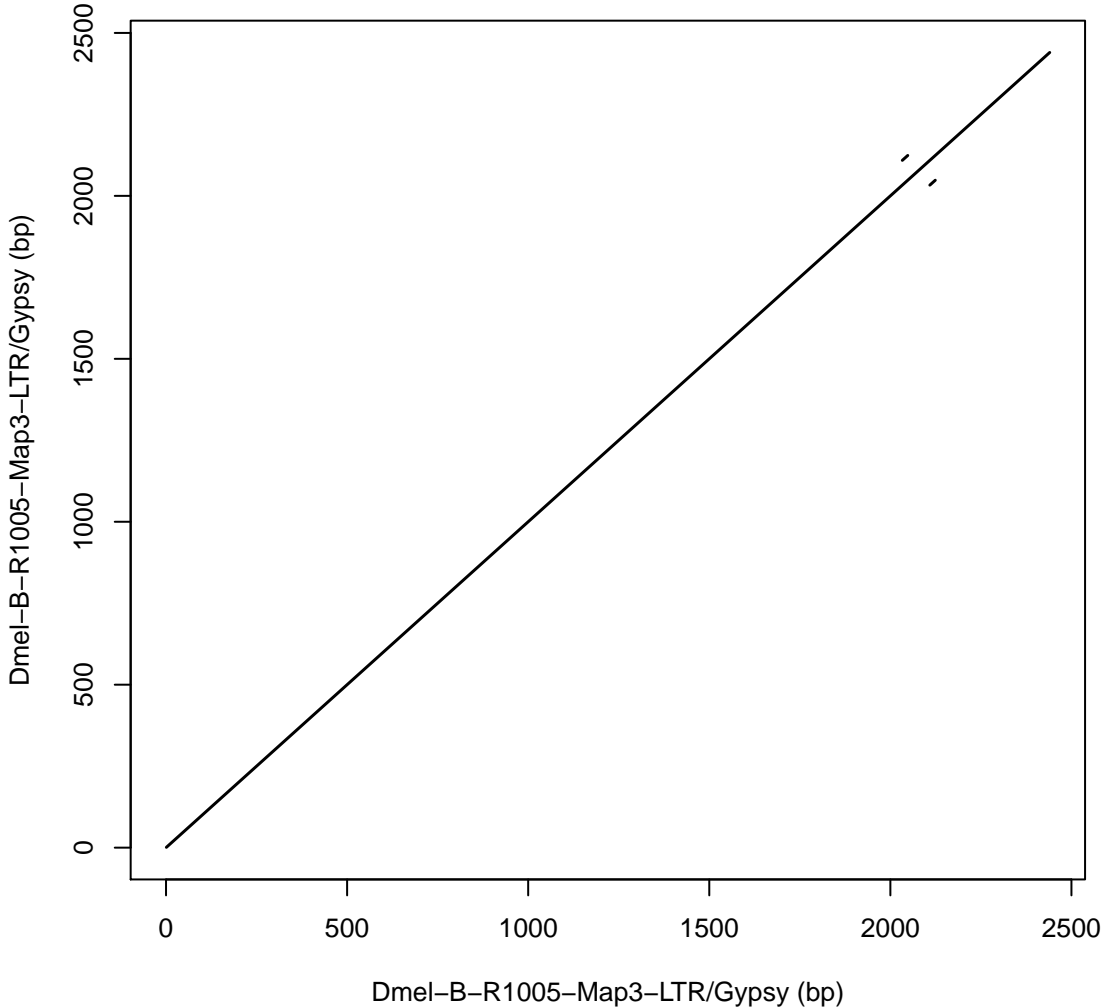
TE: Dmel-B-R1005-Map3-LTR/Gypsy  
consensus size: 2440bp; fragments: 41; full length: 1 (>=2196bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

