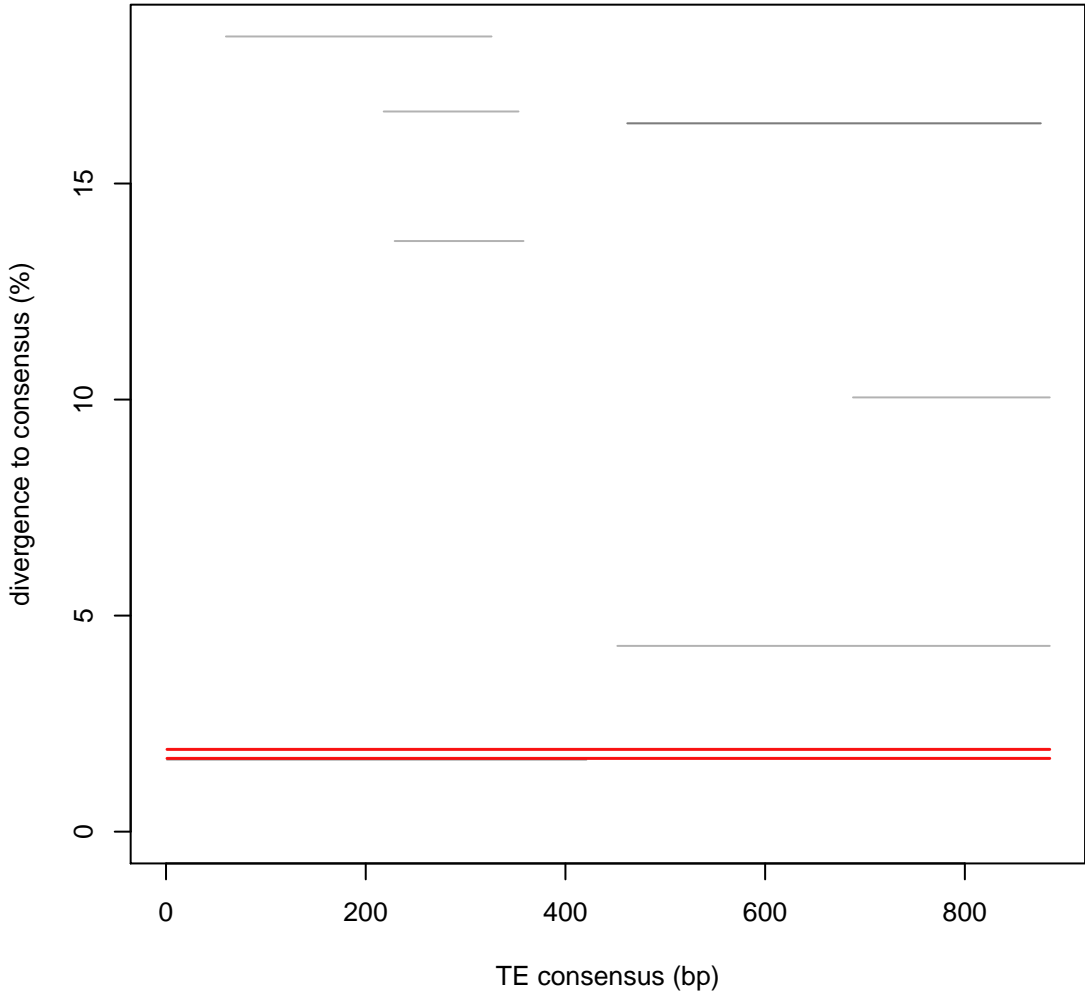
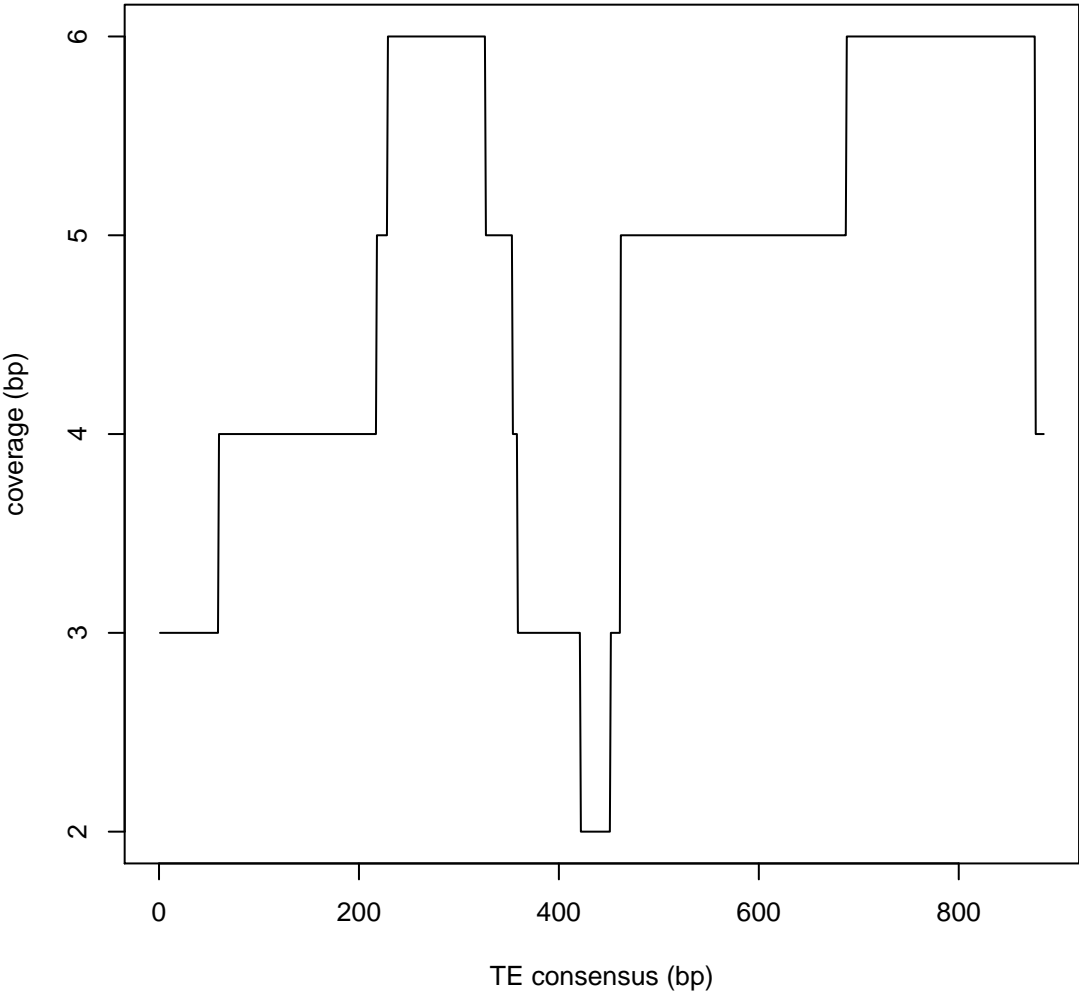


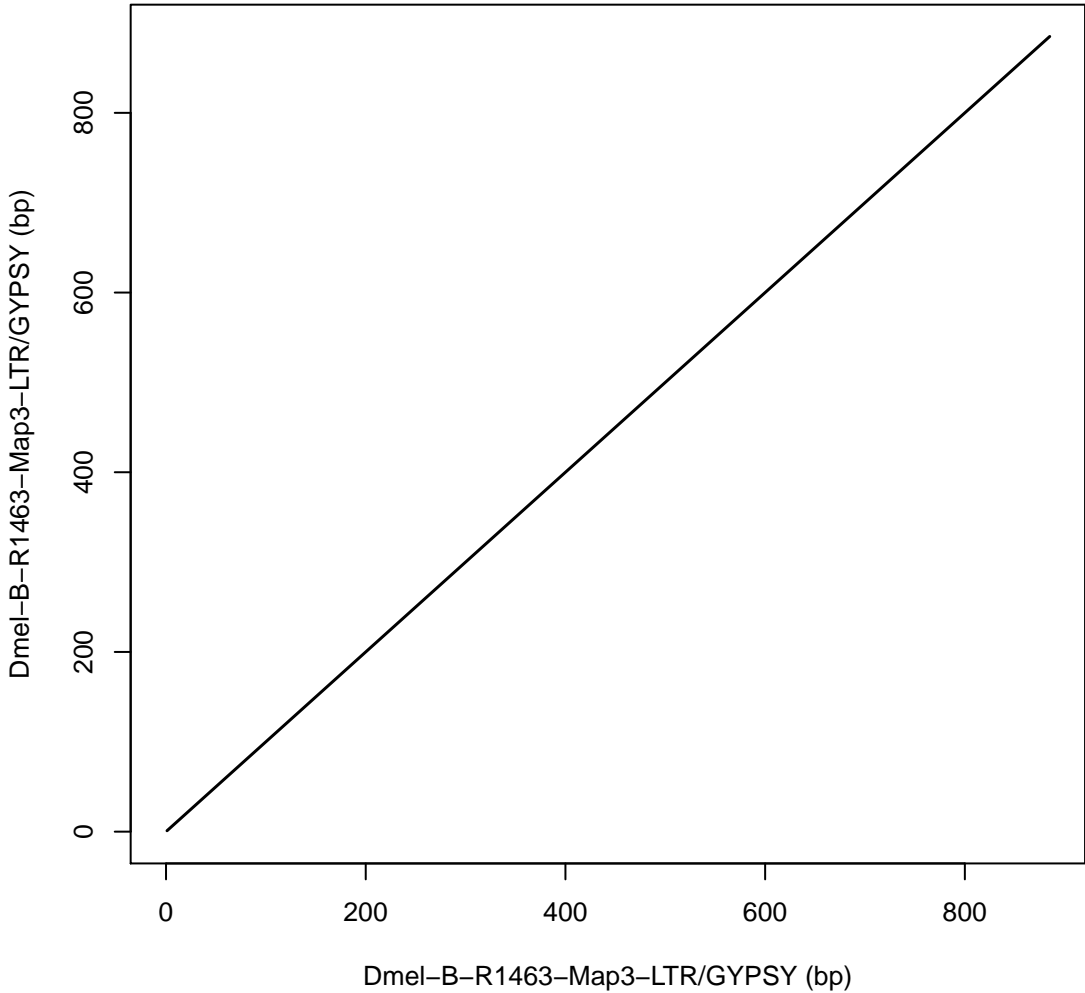
TE: Dmel-B-R1463-Map3-LTR/Gypsy  
consensus size: 885bp; fragments: 10; full length: 2 (>=796.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

