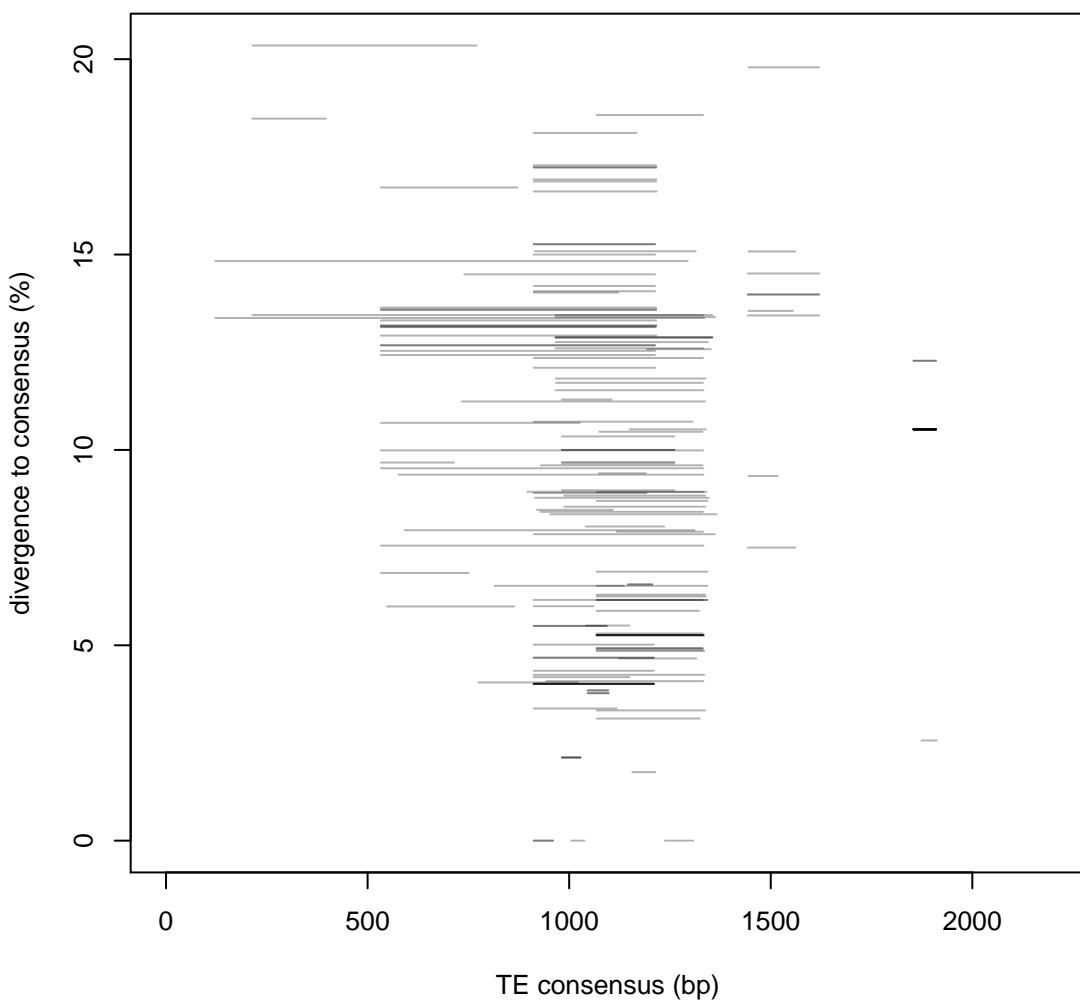
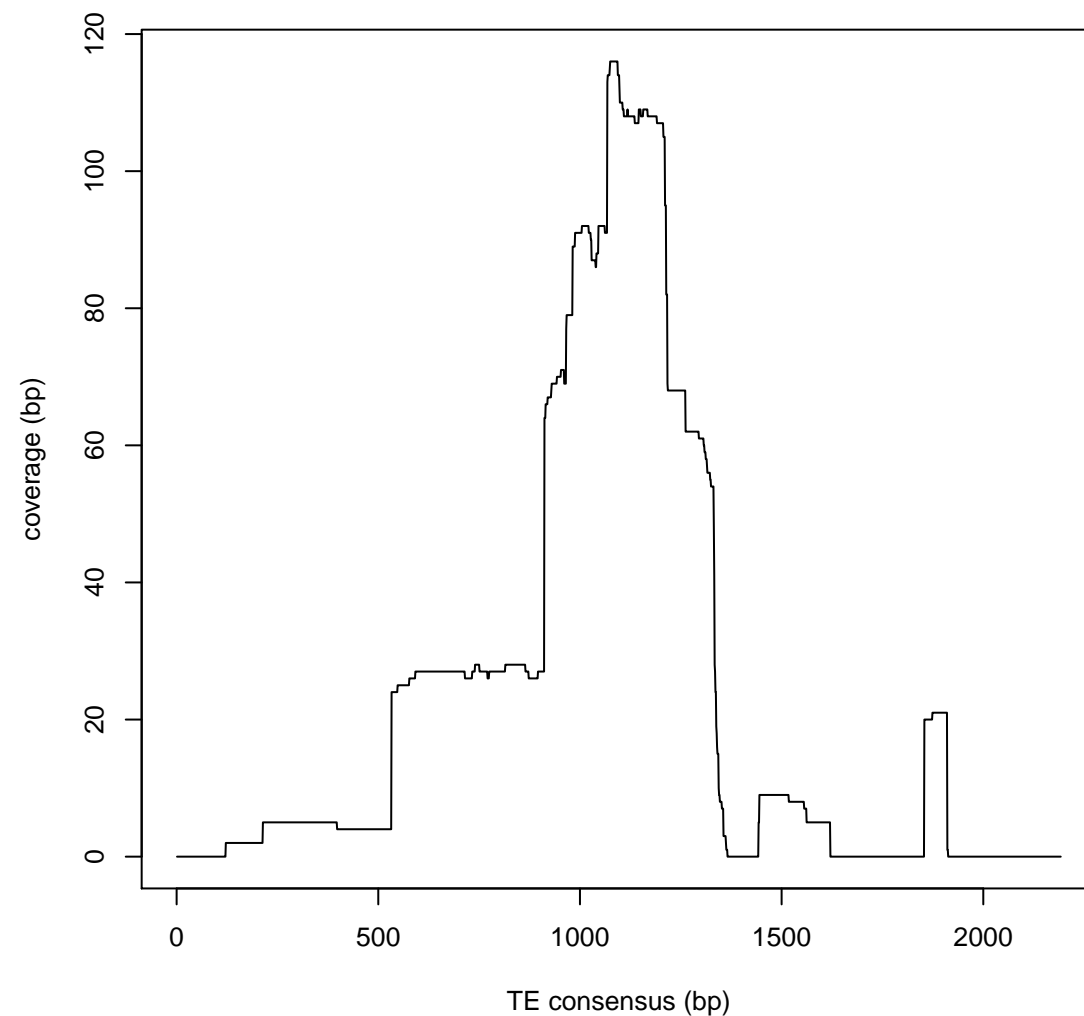


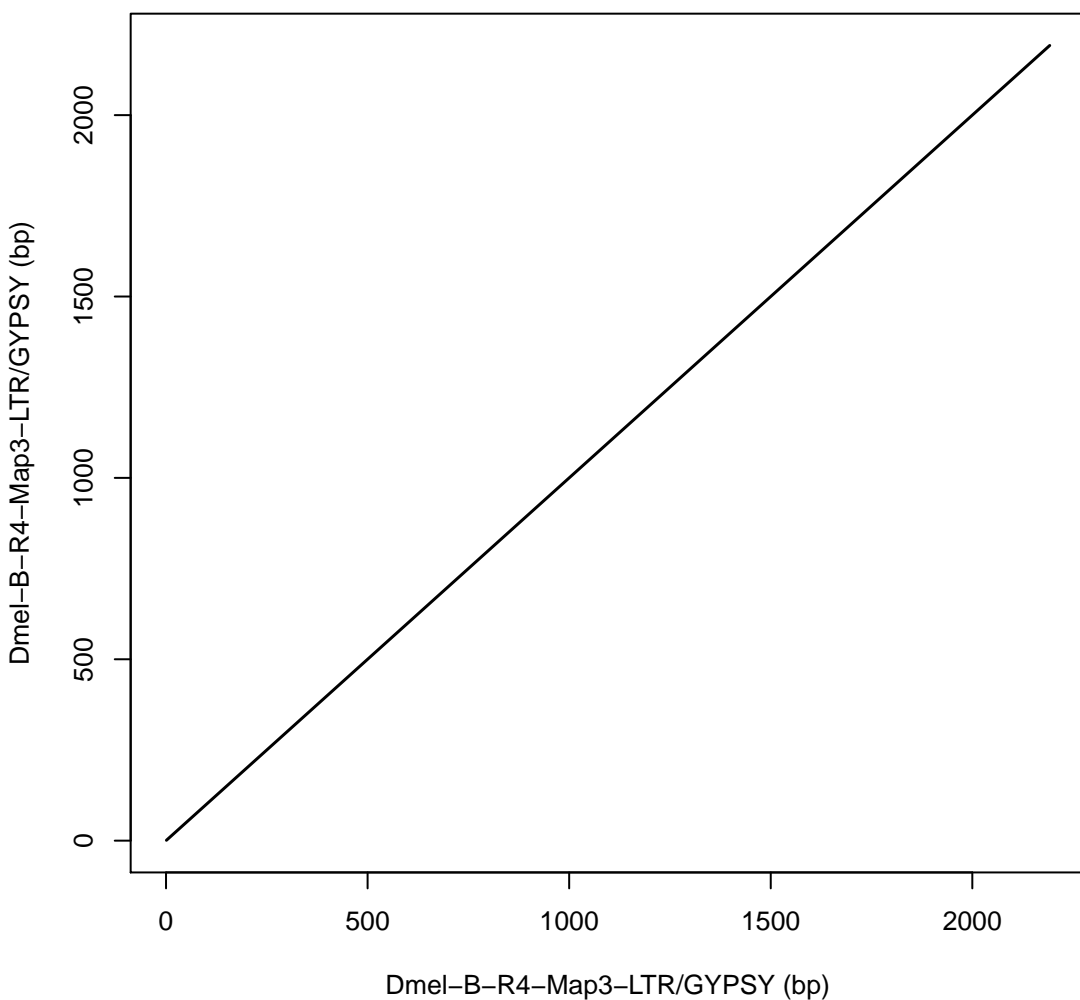
TE: Dmel-B-R4-Map3-LTR/Gypsy  
consensus size: 2192bp; fragments: 169; full length: 0 (>=1972.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

