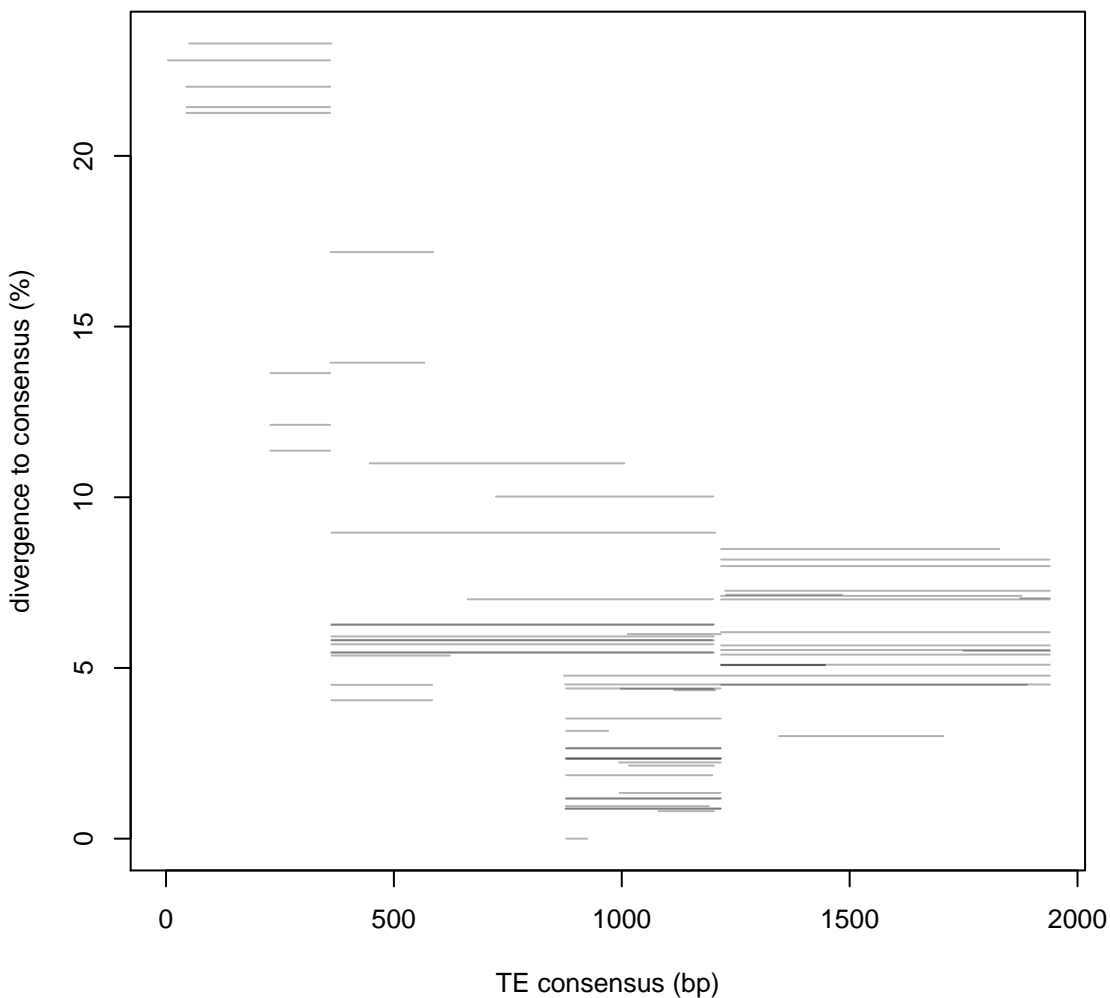
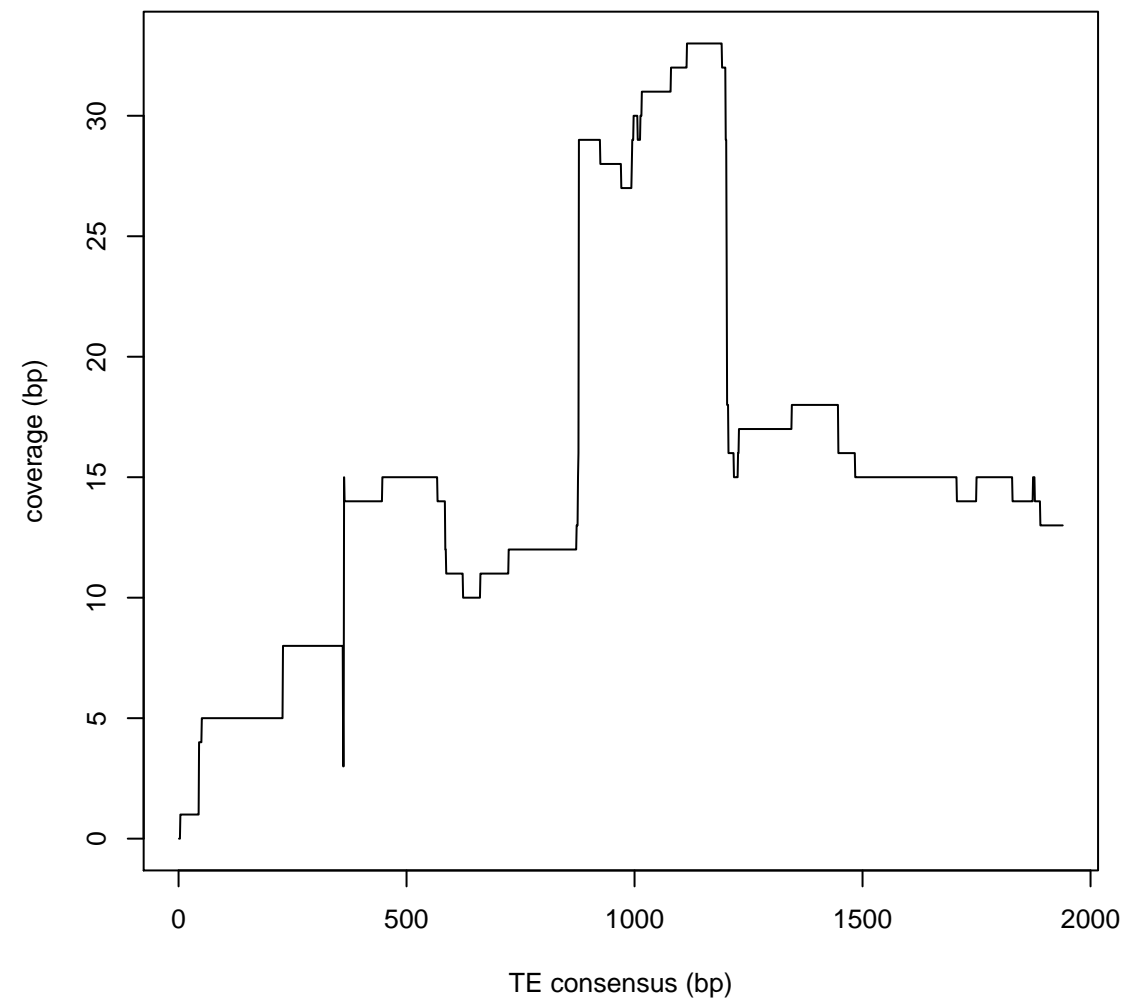


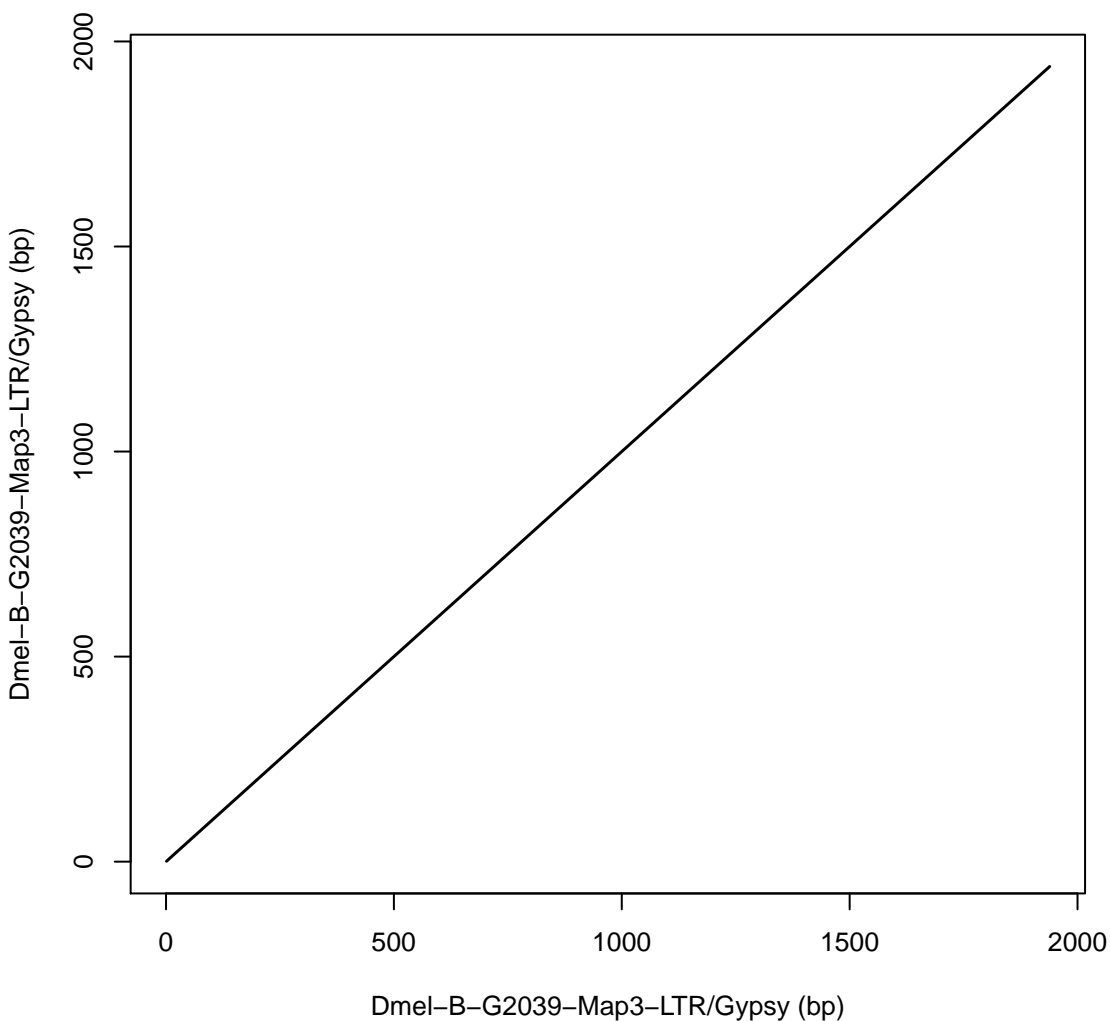
TE: Dmel-B-G2039-Map3-LTR/Gypsy  
consensus size: 1939bp; fragments: 67; full length: 0 (>=1745.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

