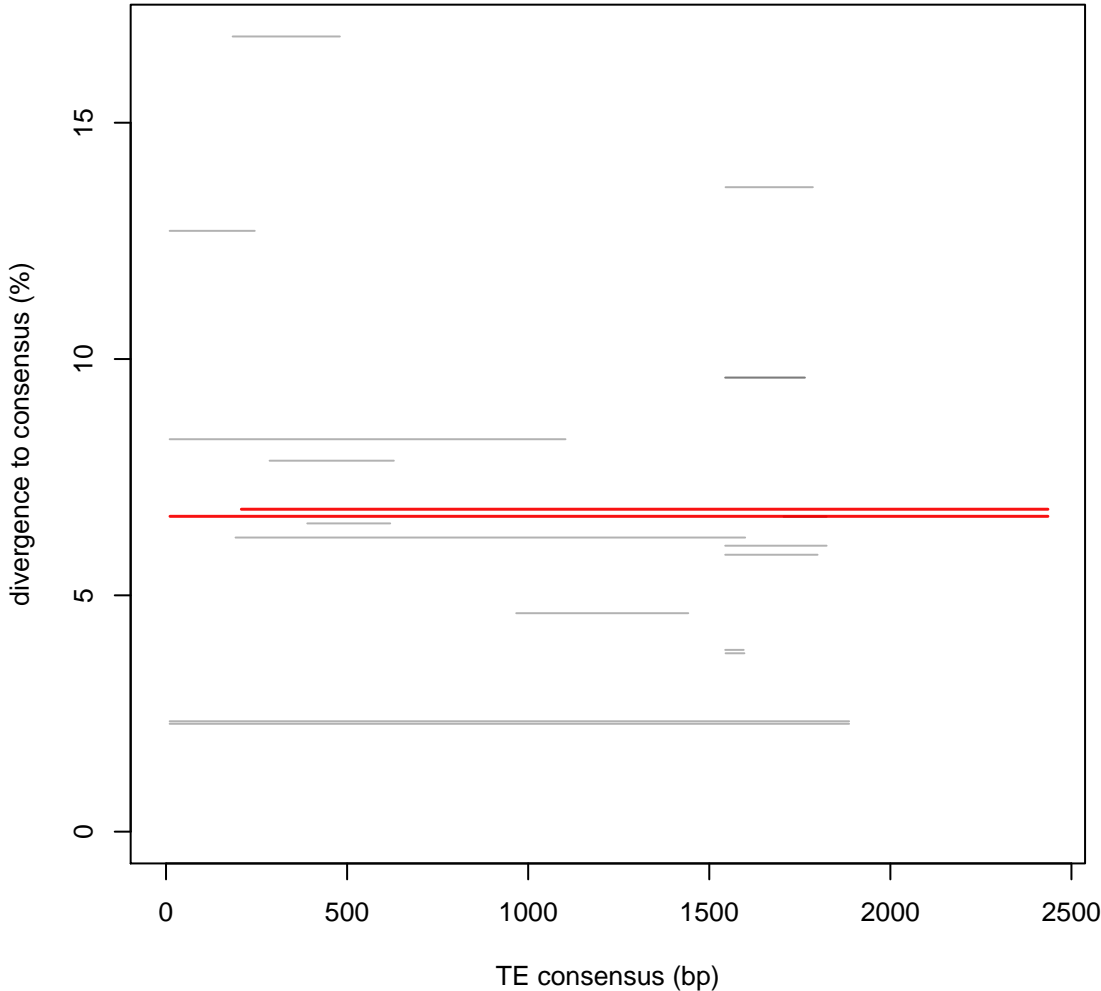
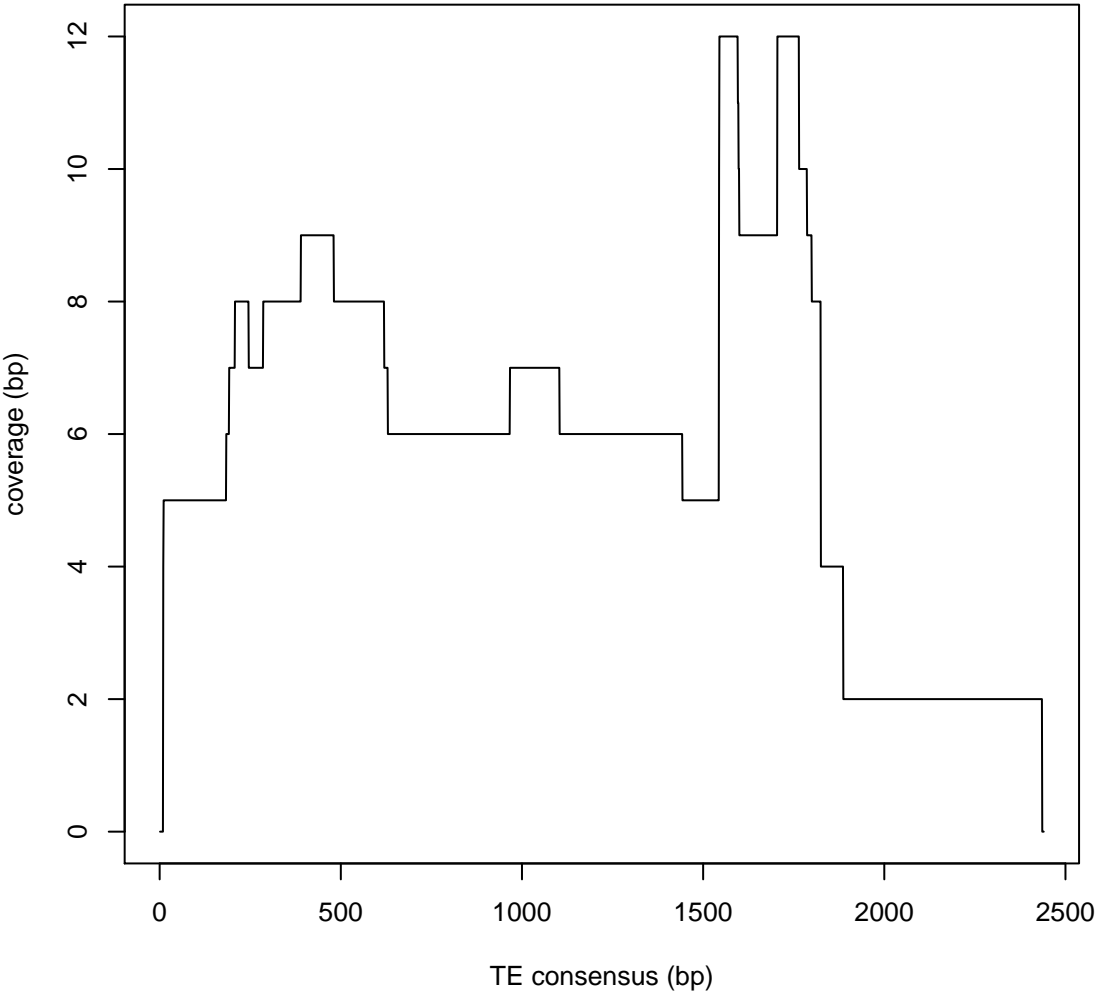


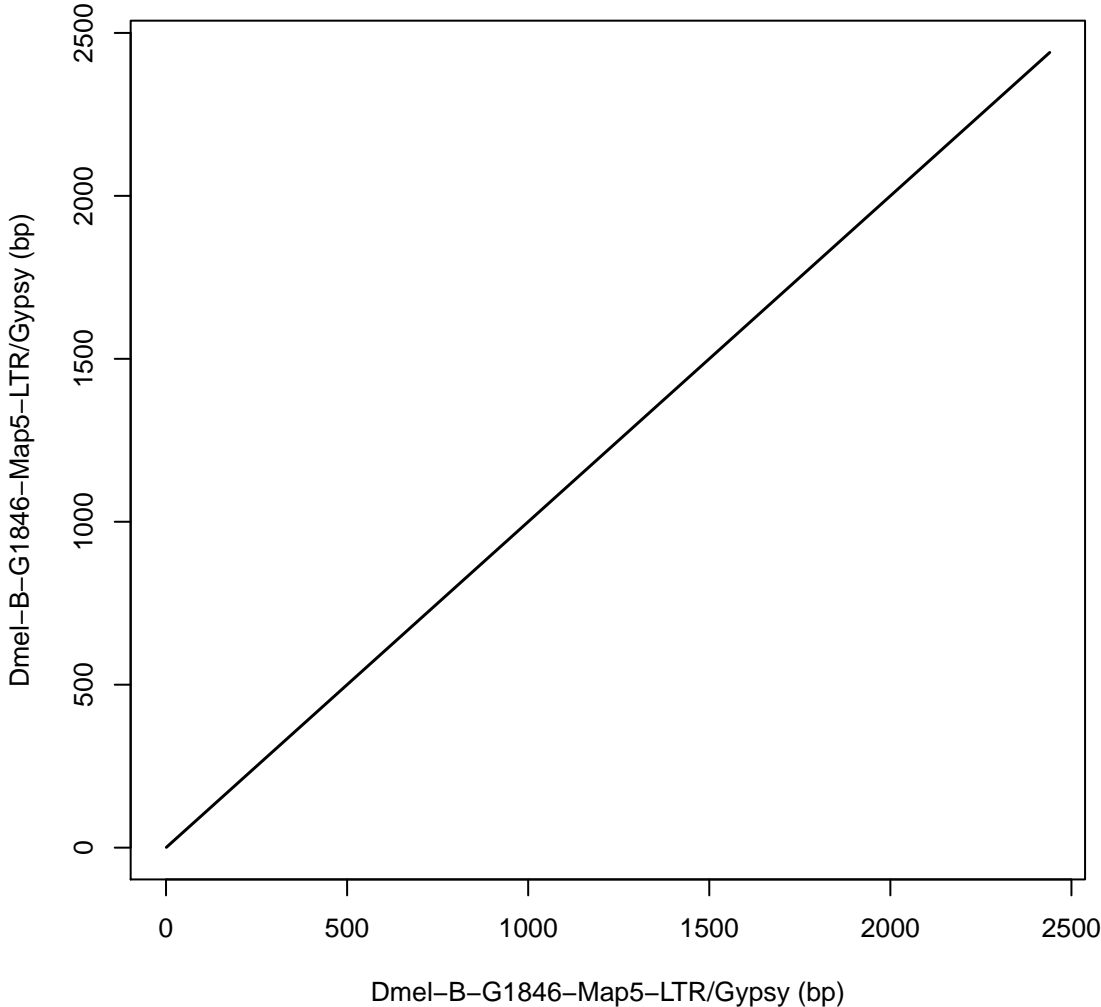
TE: Dmel-B-G1846-Map5-LTR/Gypsy
consensus size: 2440bp; fragments: 21; full length: 2 (>=2196bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

