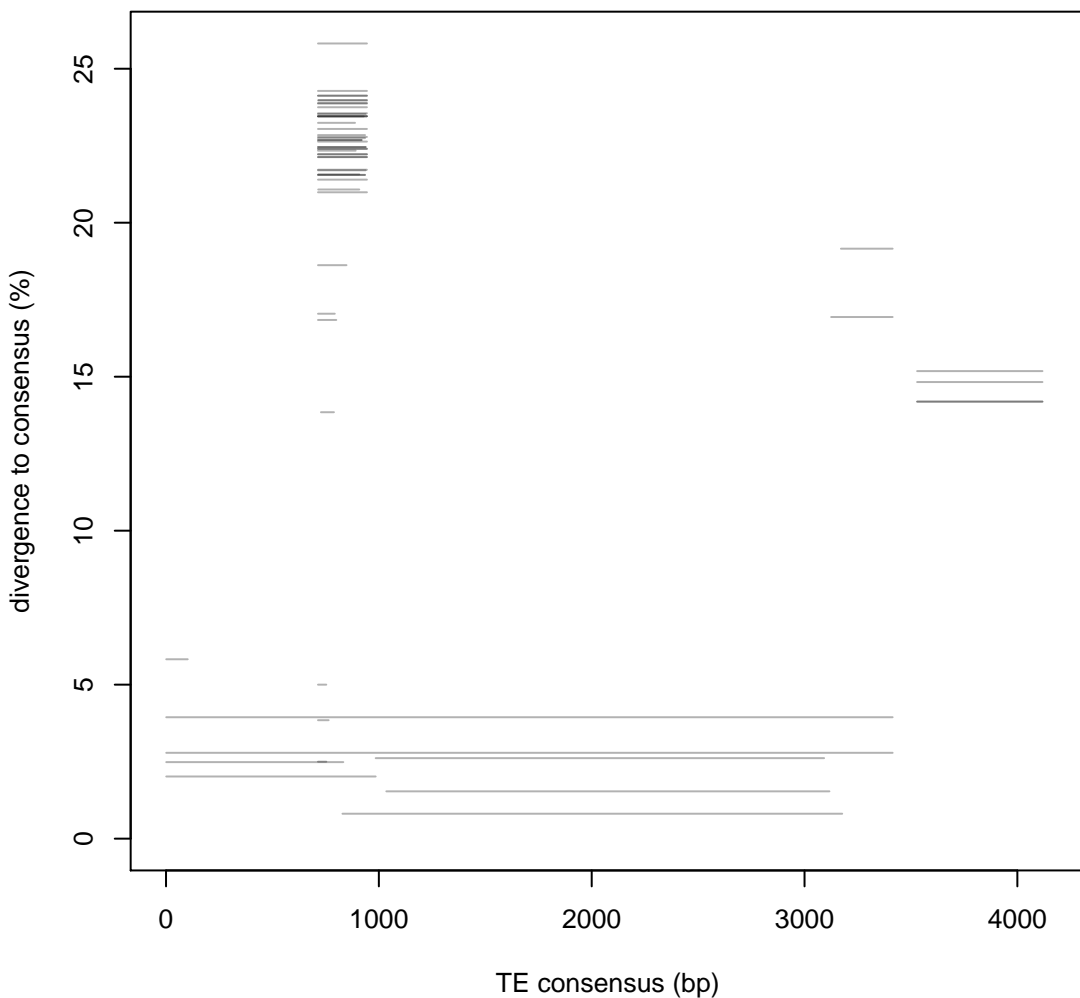
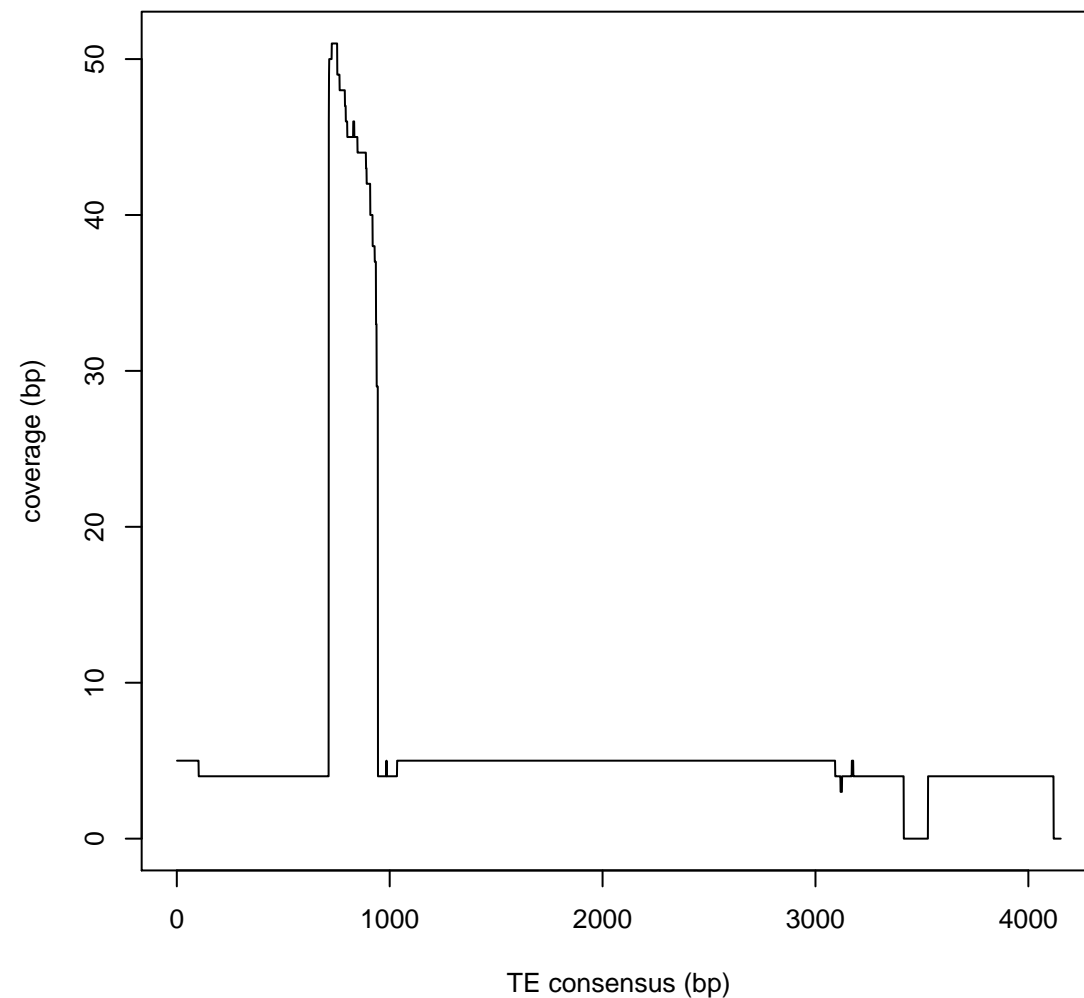


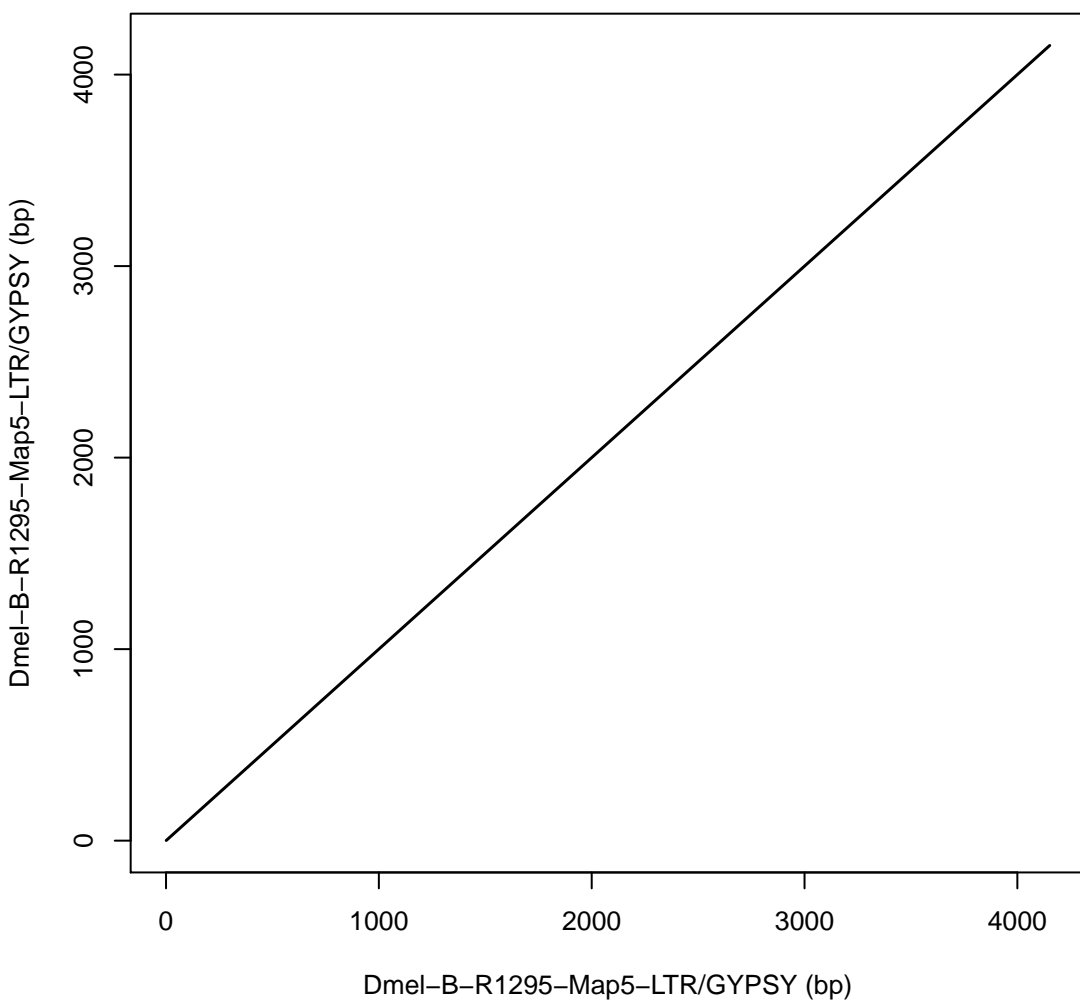
TE: Dmel-B-R1295-Map5-LTR/GYPSY
consensus size: 4152bp; fragments: 61; full length: 0 (>=3736.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

