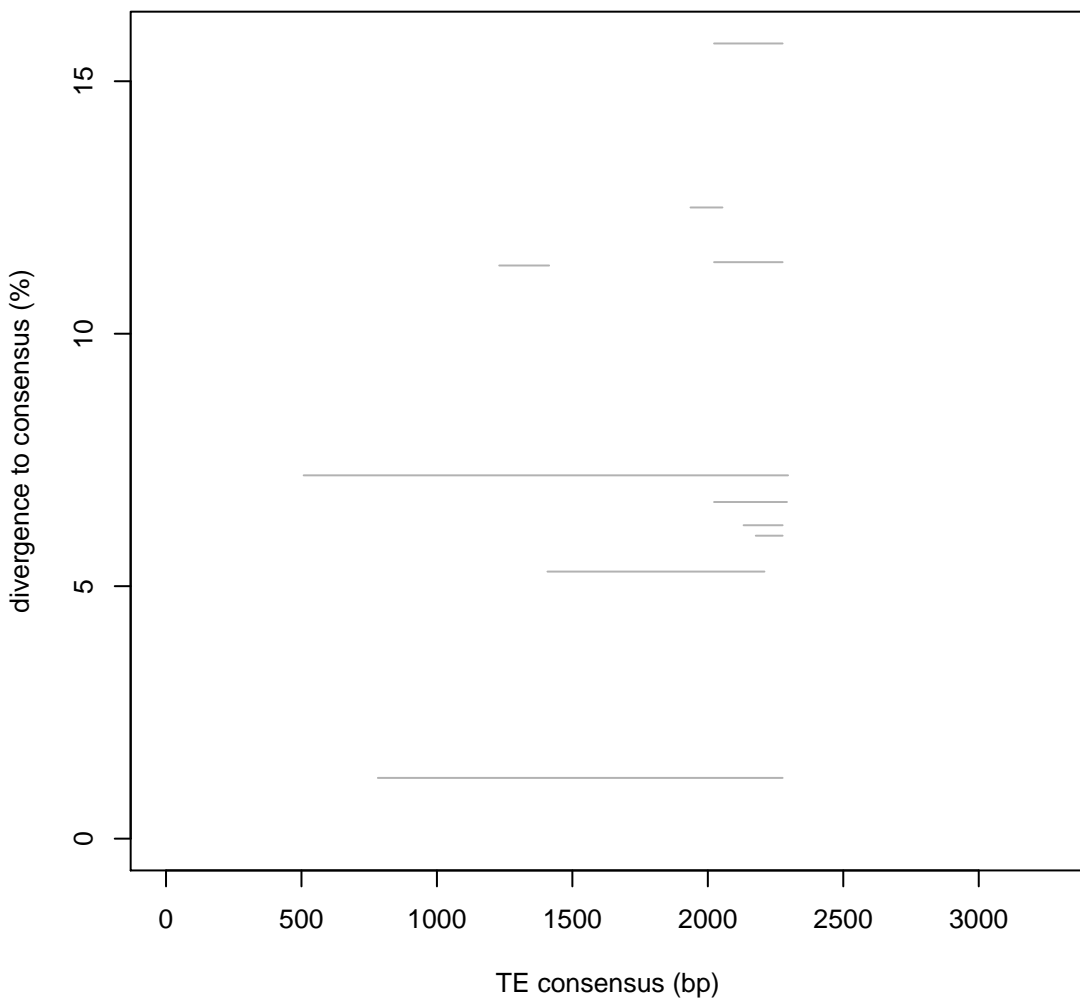
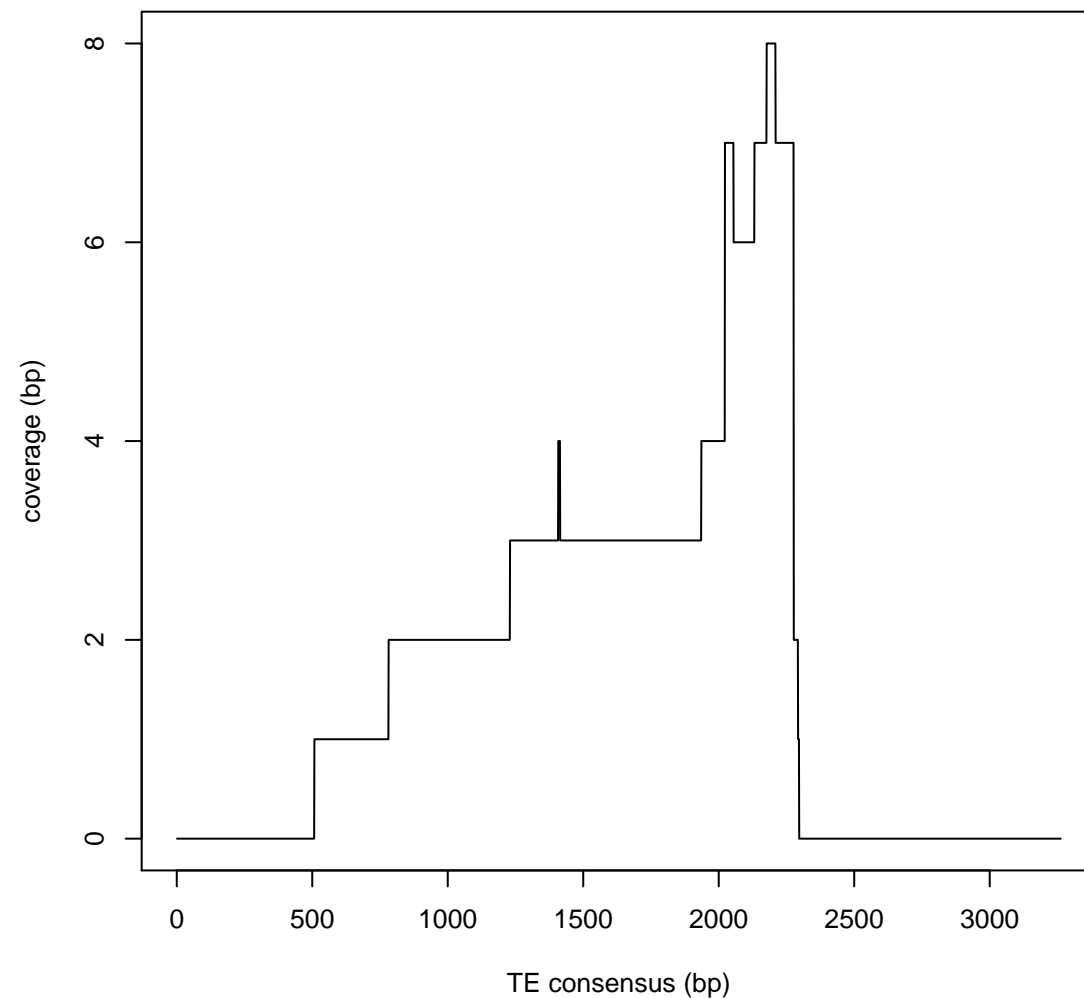


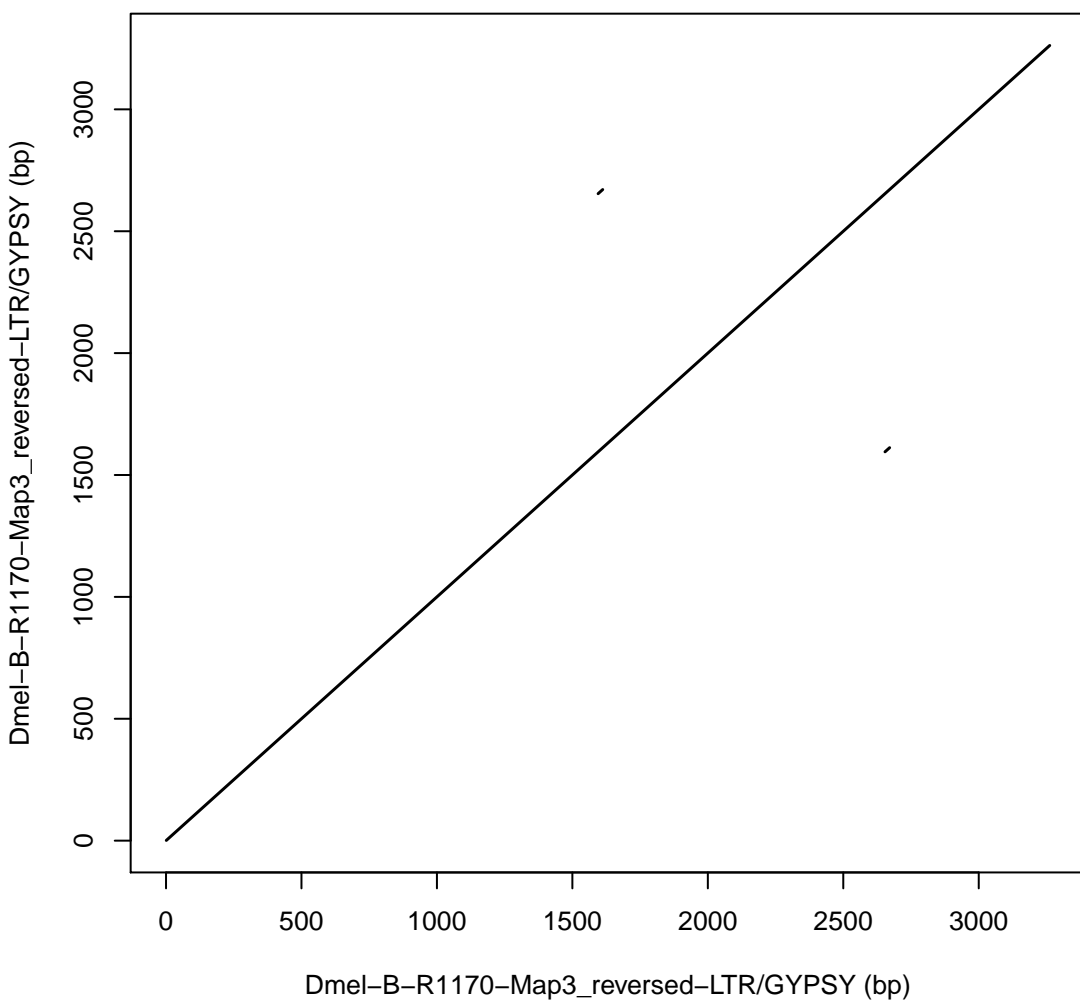
TE: Dmel-B-R1170-Map3_reversed-LTR/GYPSY
consensus size: 3262bp; fragments: 10; full length: 0 (>=2935.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

