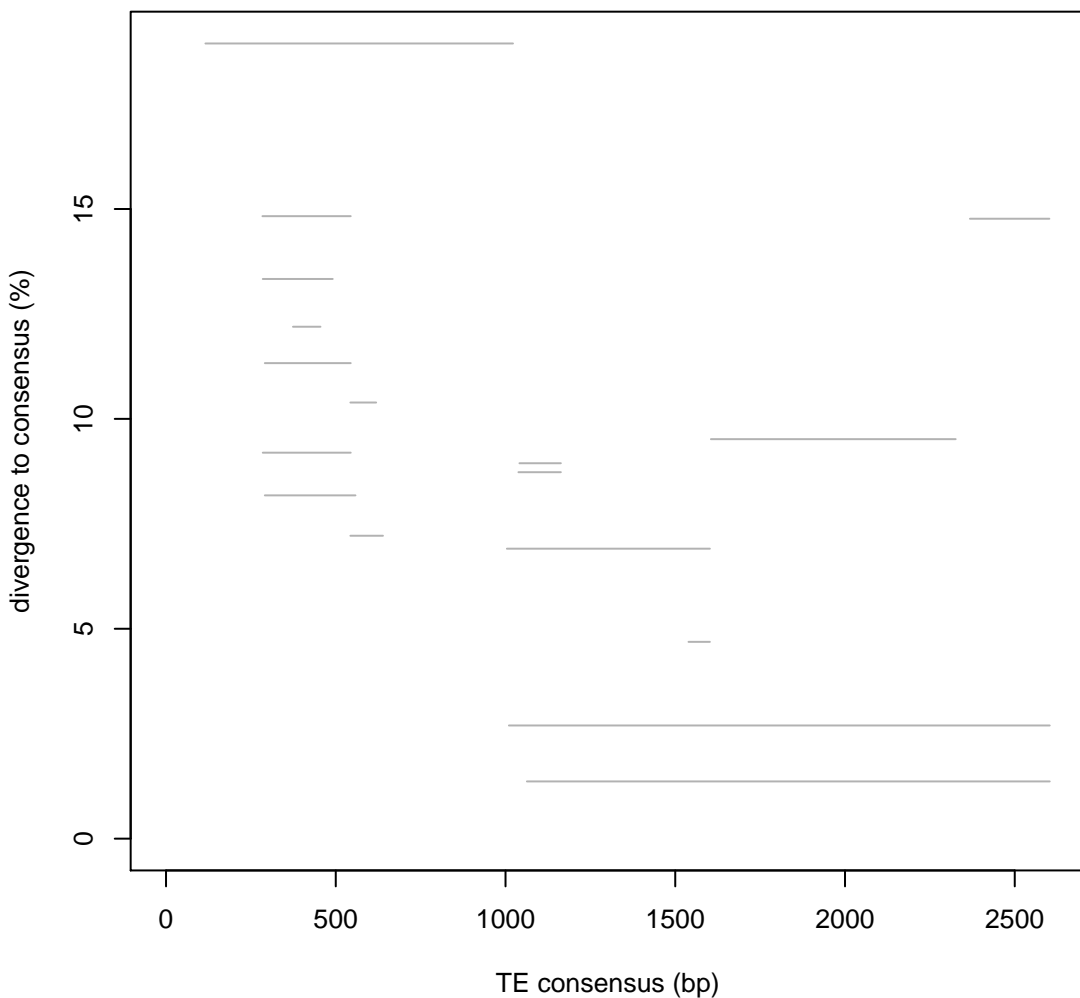
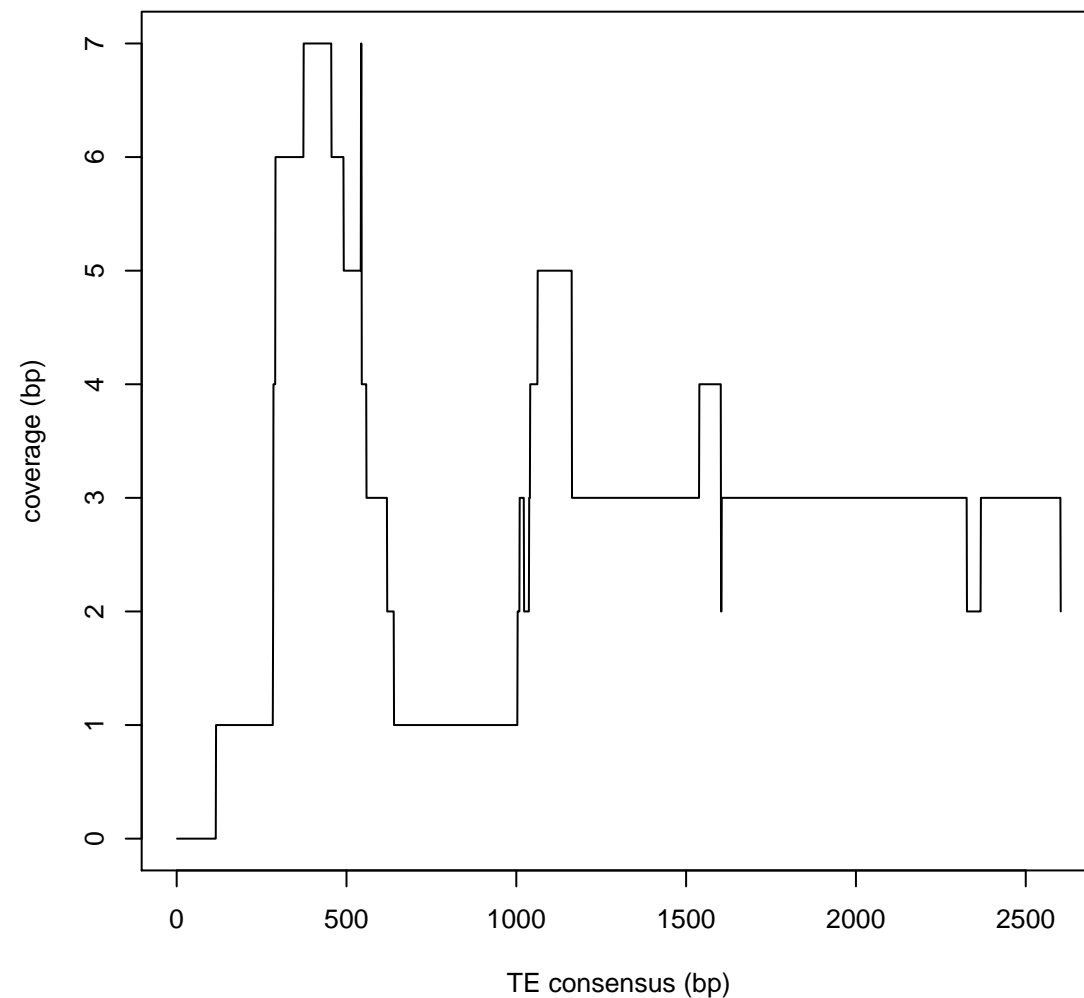


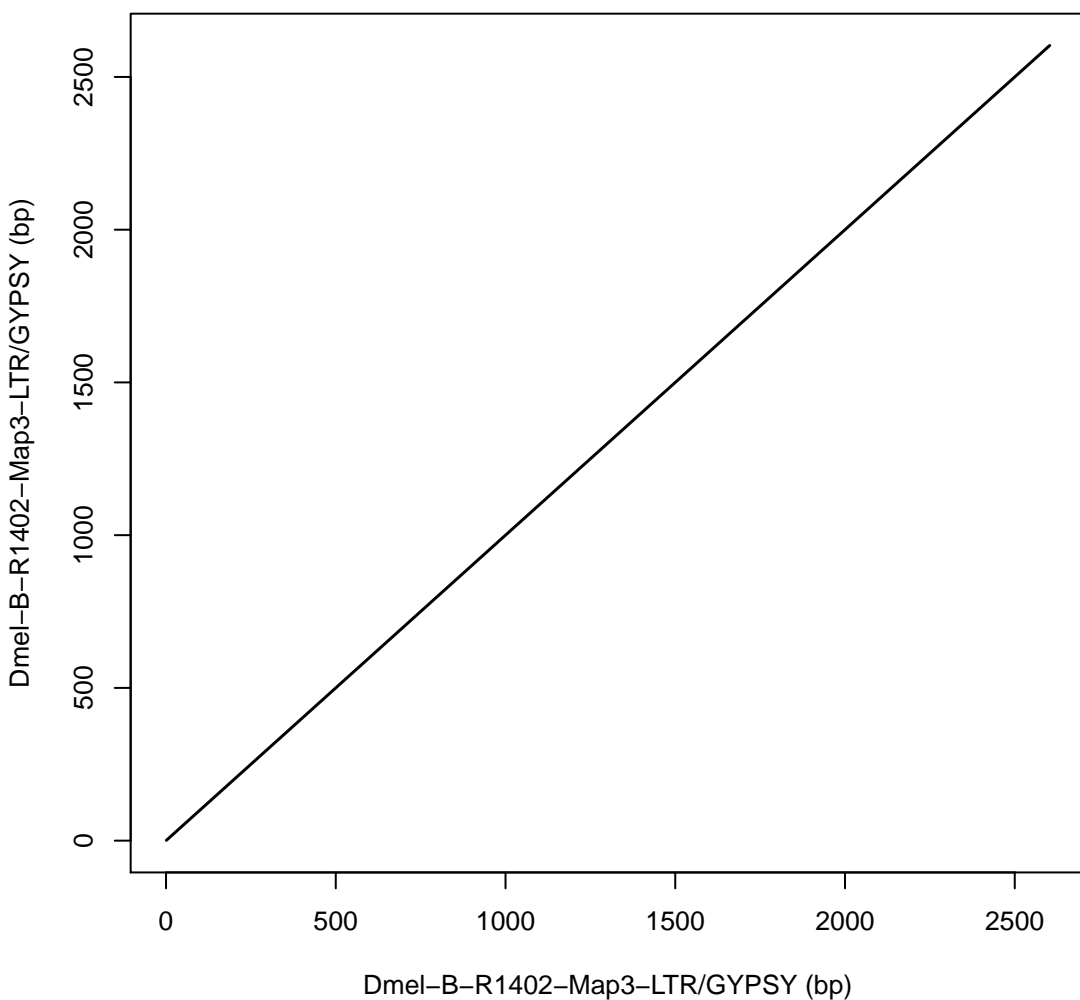
TE: Dmel-B-R1402-Map3-LTR/GYPSY
consensus size: 2603bp; fragments: 17; full length: 0 (>=2342.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

