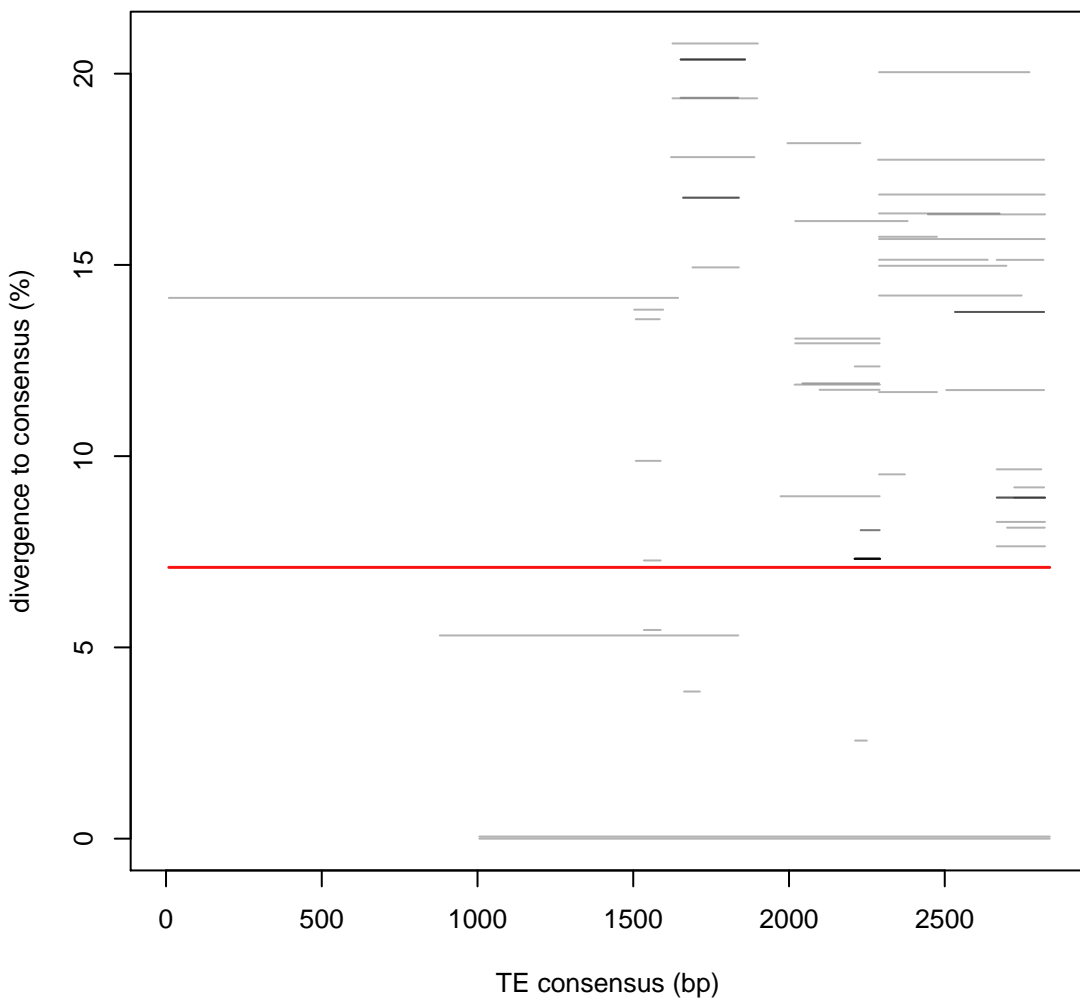
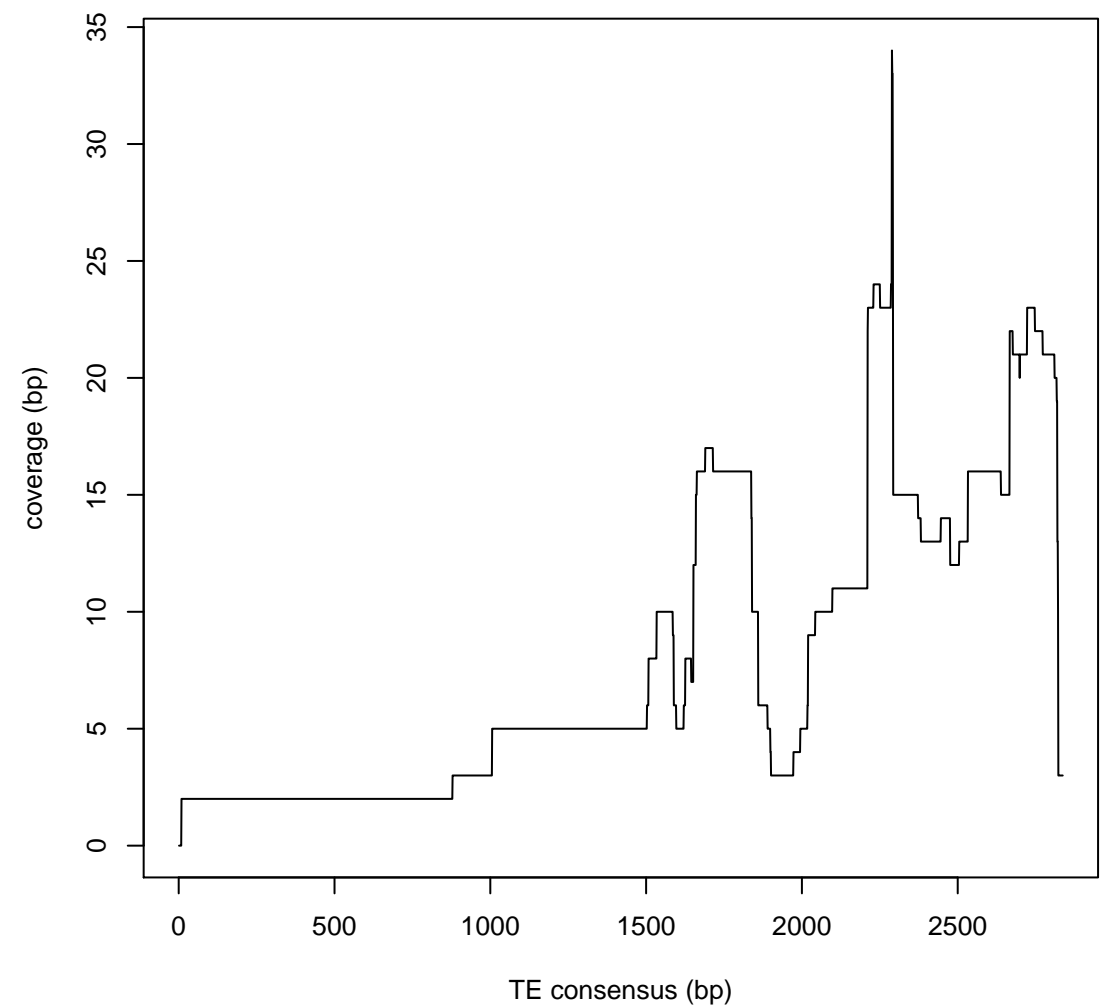


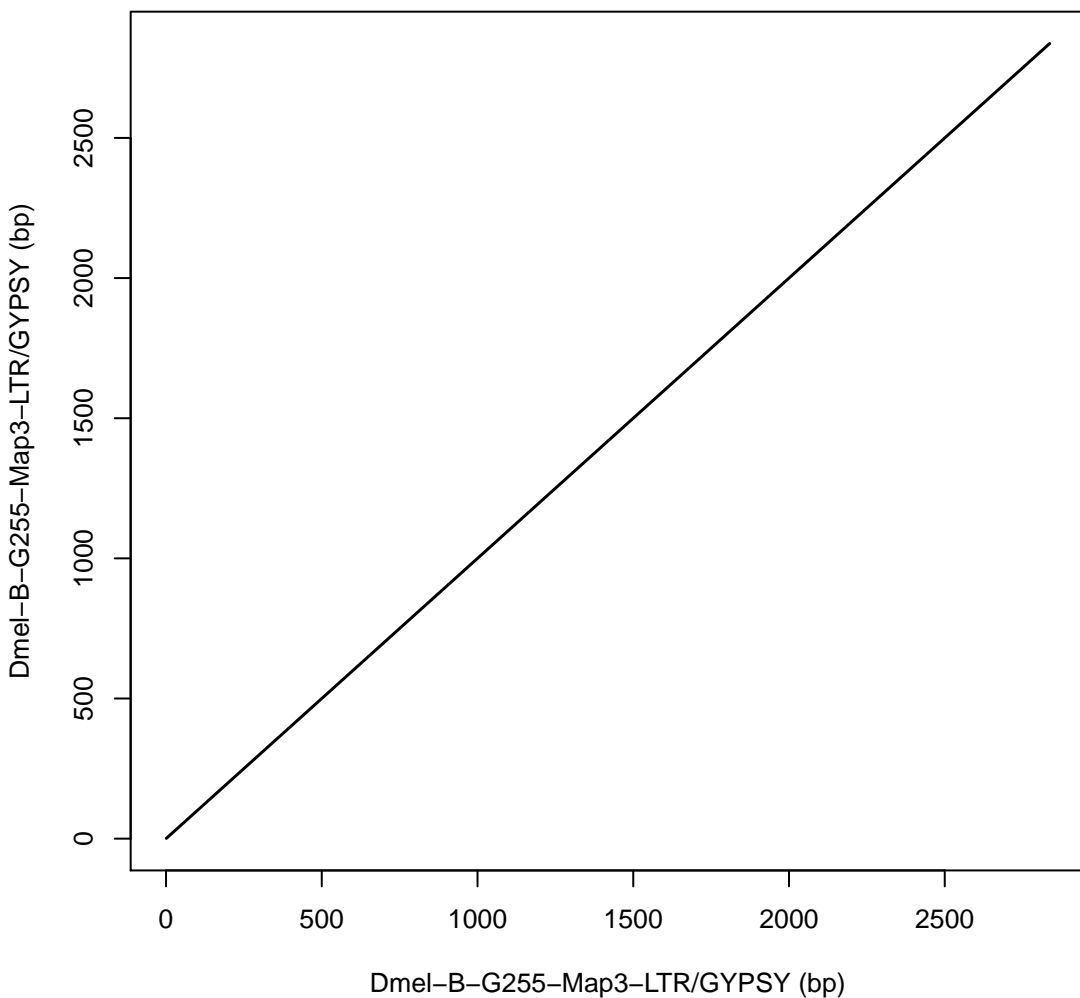
TE: Dmel-B-G255-Map3-LTR/GYPSY
consensus size: 2837bp; fragments: 71; full length: 1 (>=2553.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

