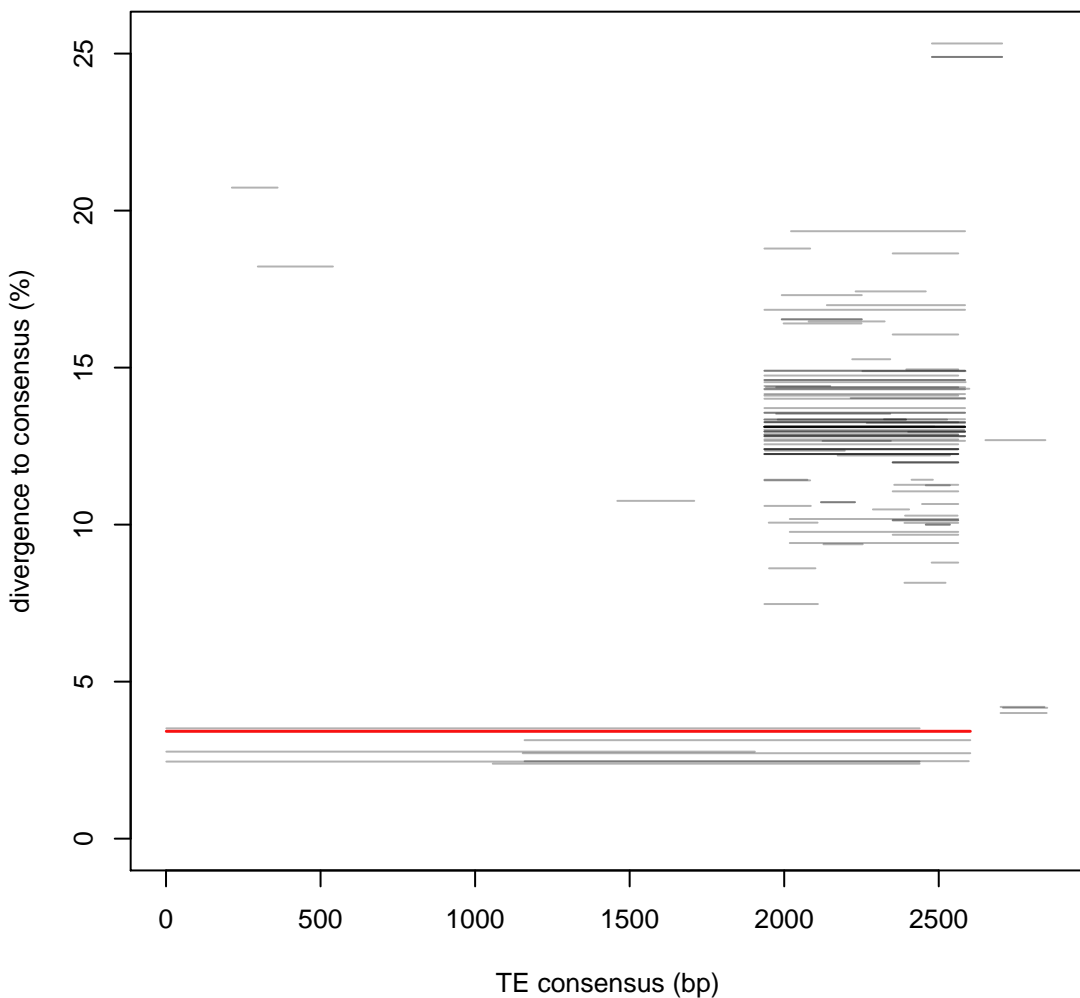
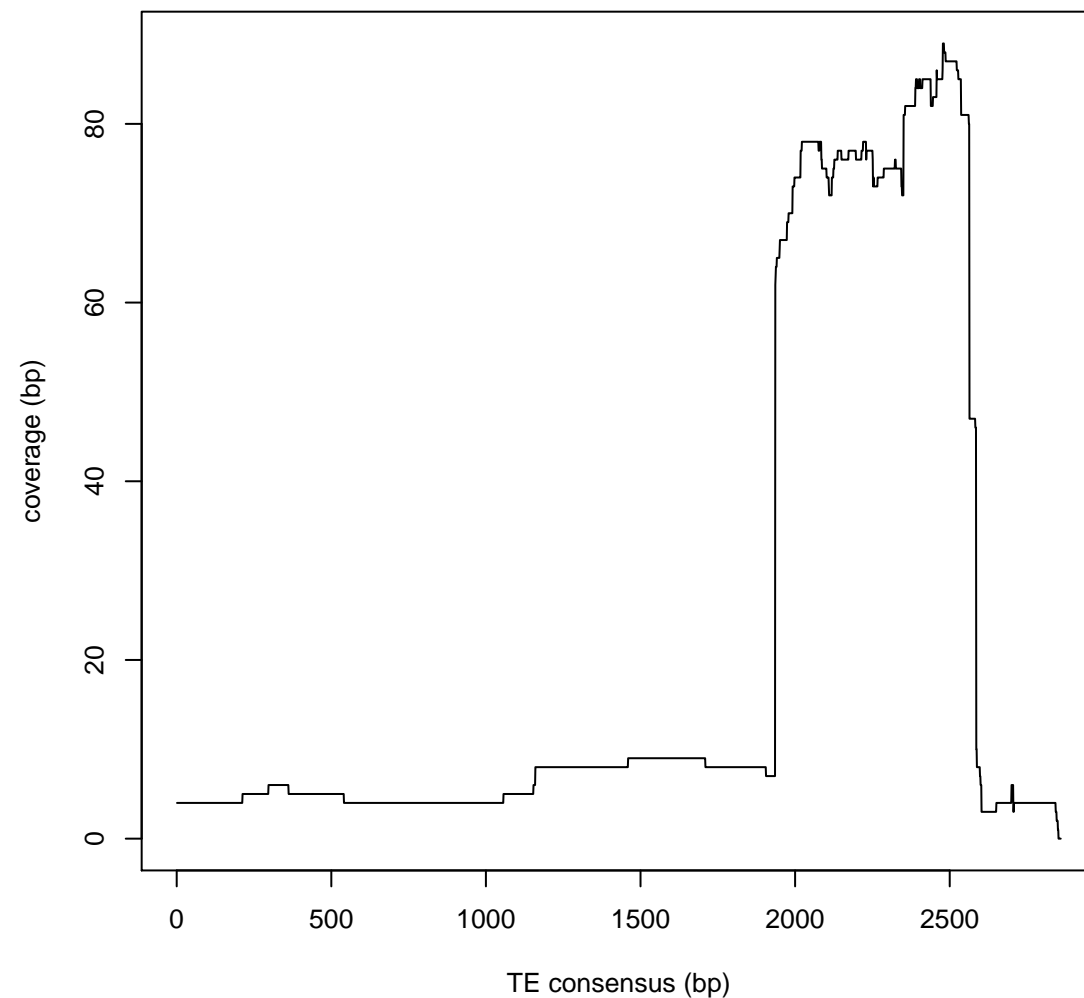


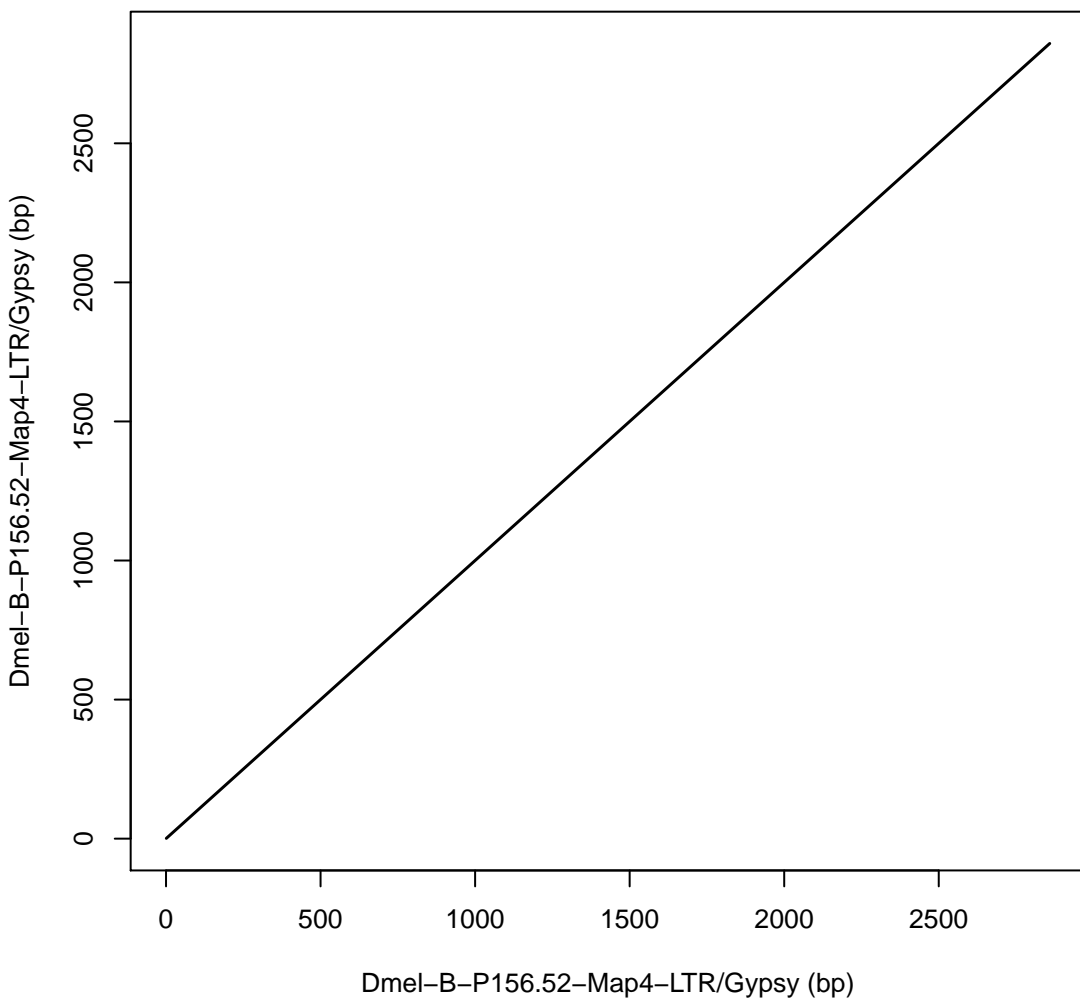
TE: Dmel-B-P156.52-Map4-LTR/Gypsy  
consensus size: 2859bp; fragments: 124; full length: 1 (>=2573.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

