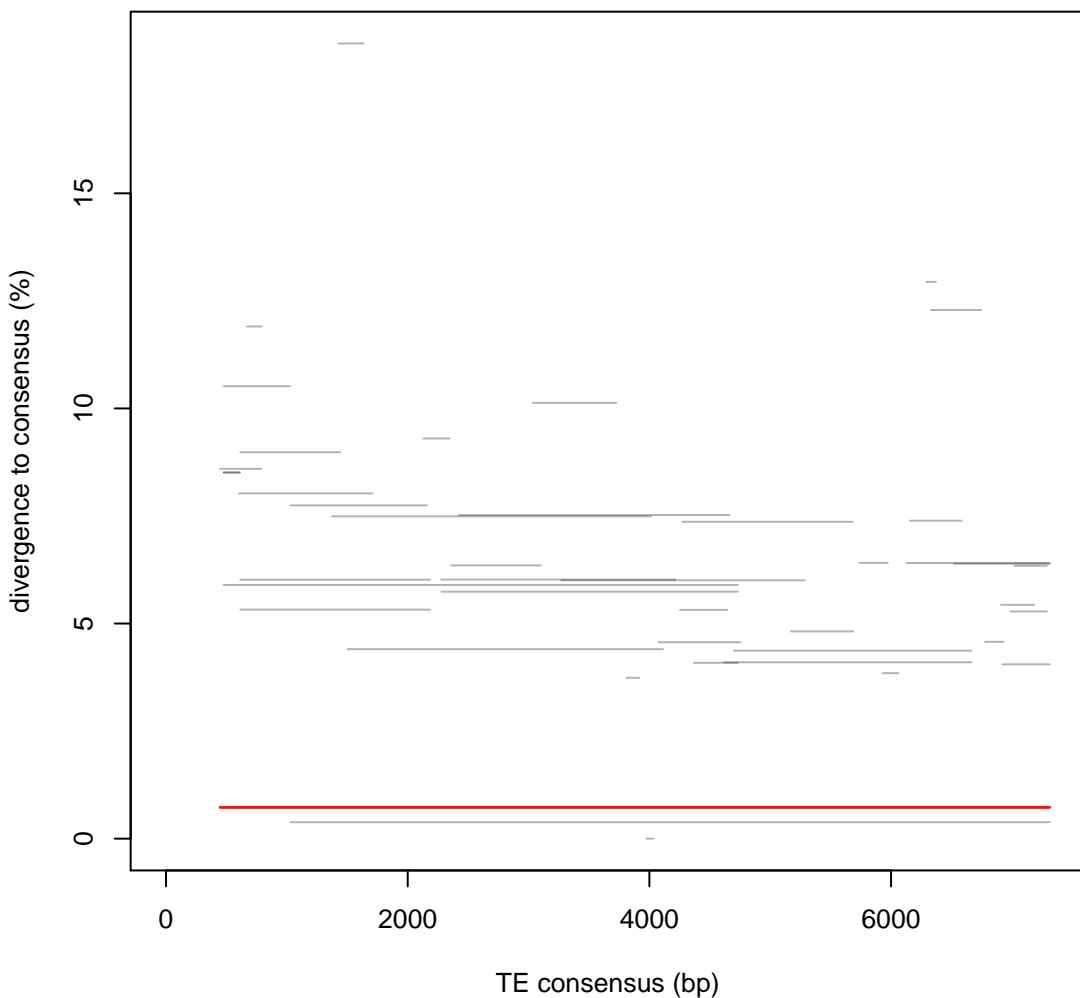
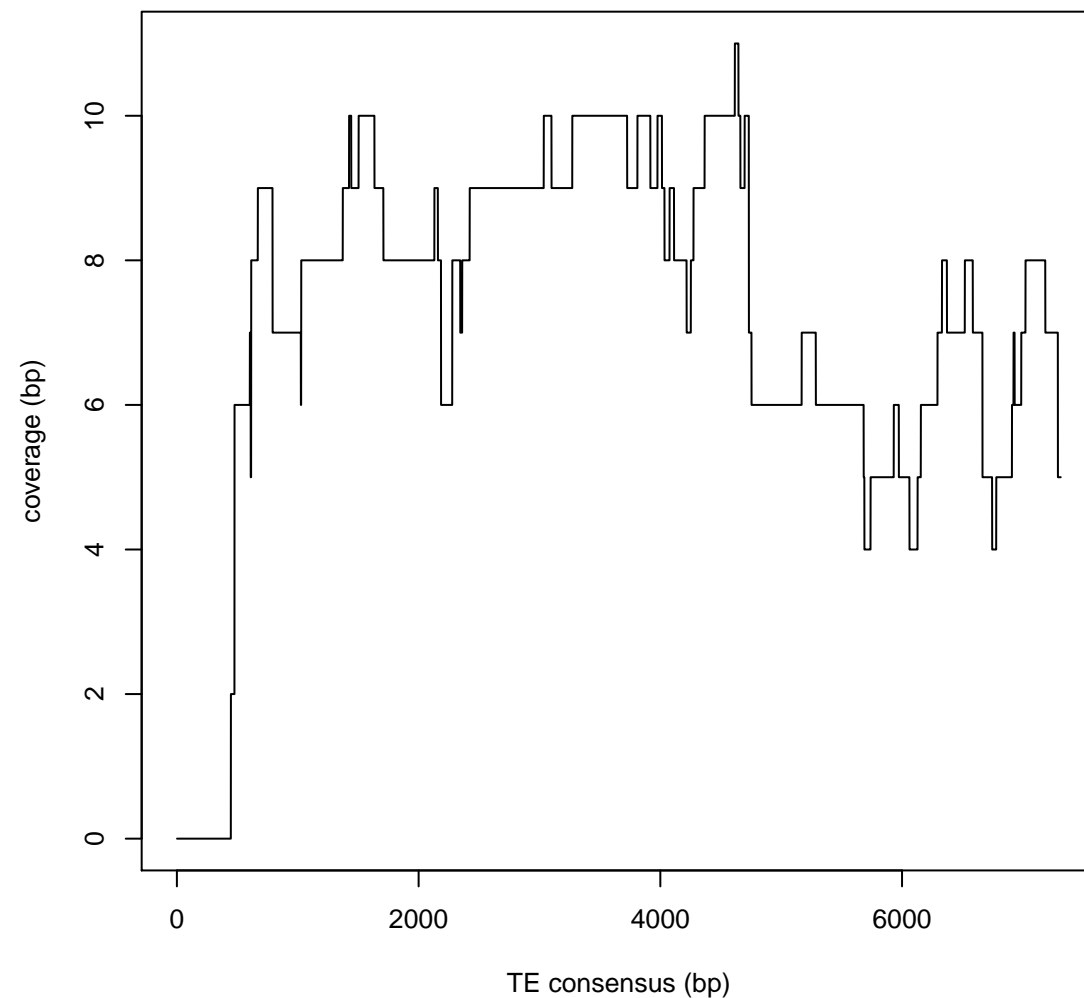


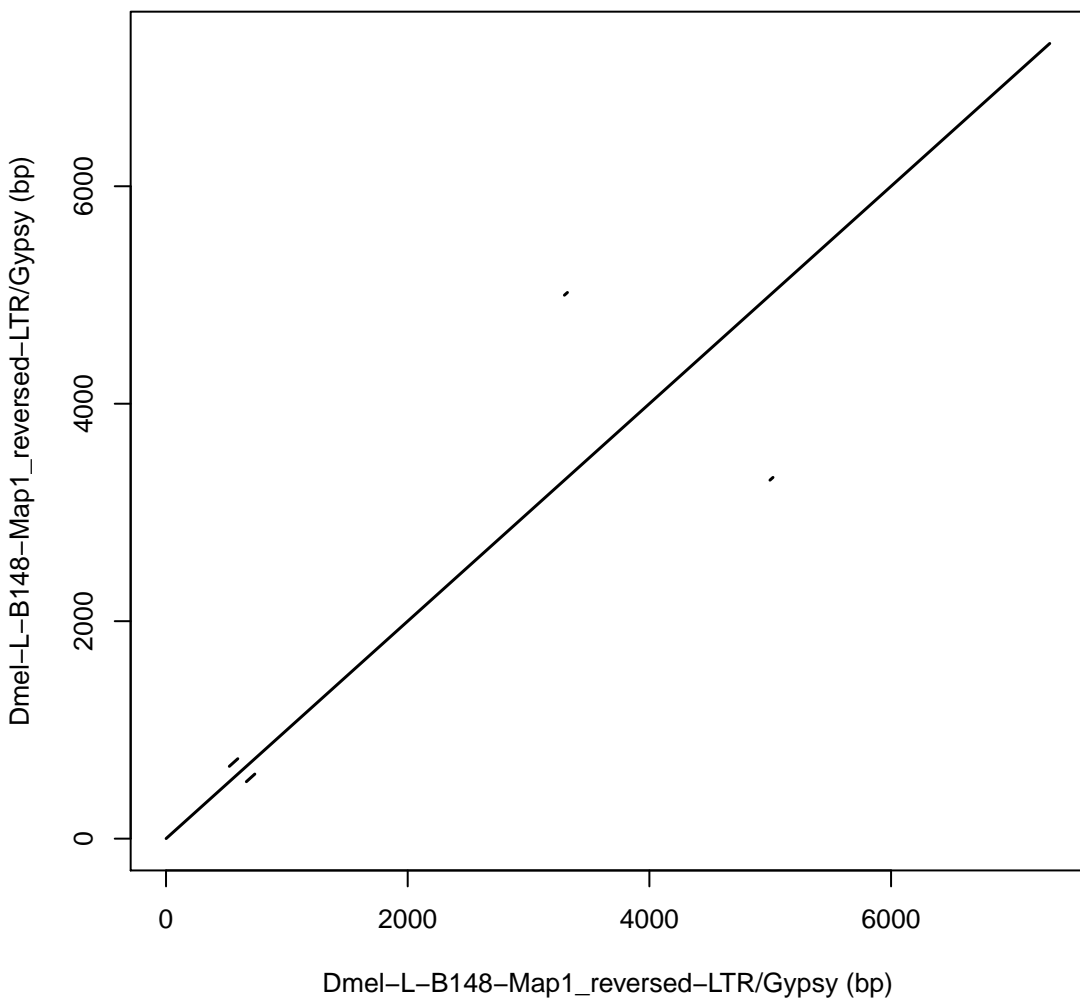
TE: Dmel-L-B148-Map1_reversed-LTR/Gypsy
consensus size: 7313bp; fragments: 44; full length: 1 (>=6581.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

