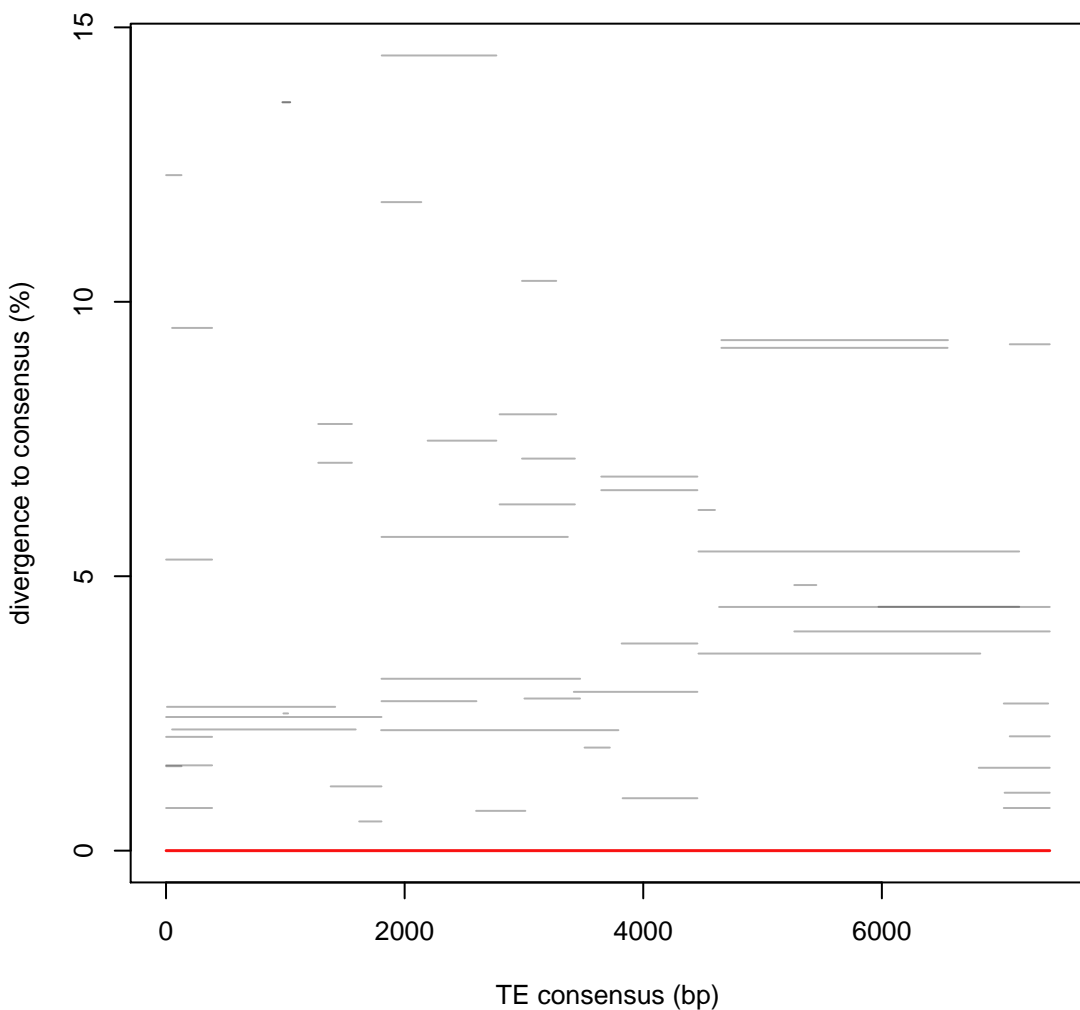
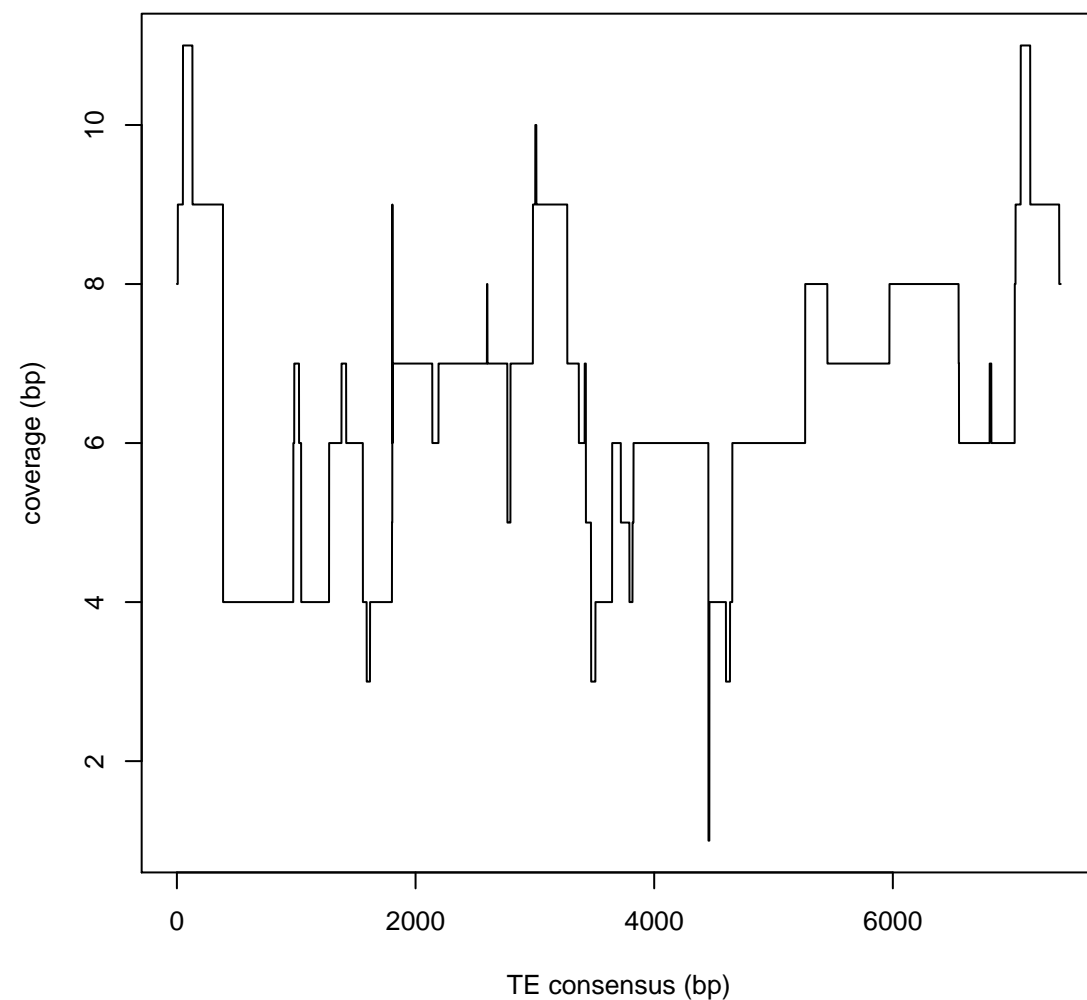


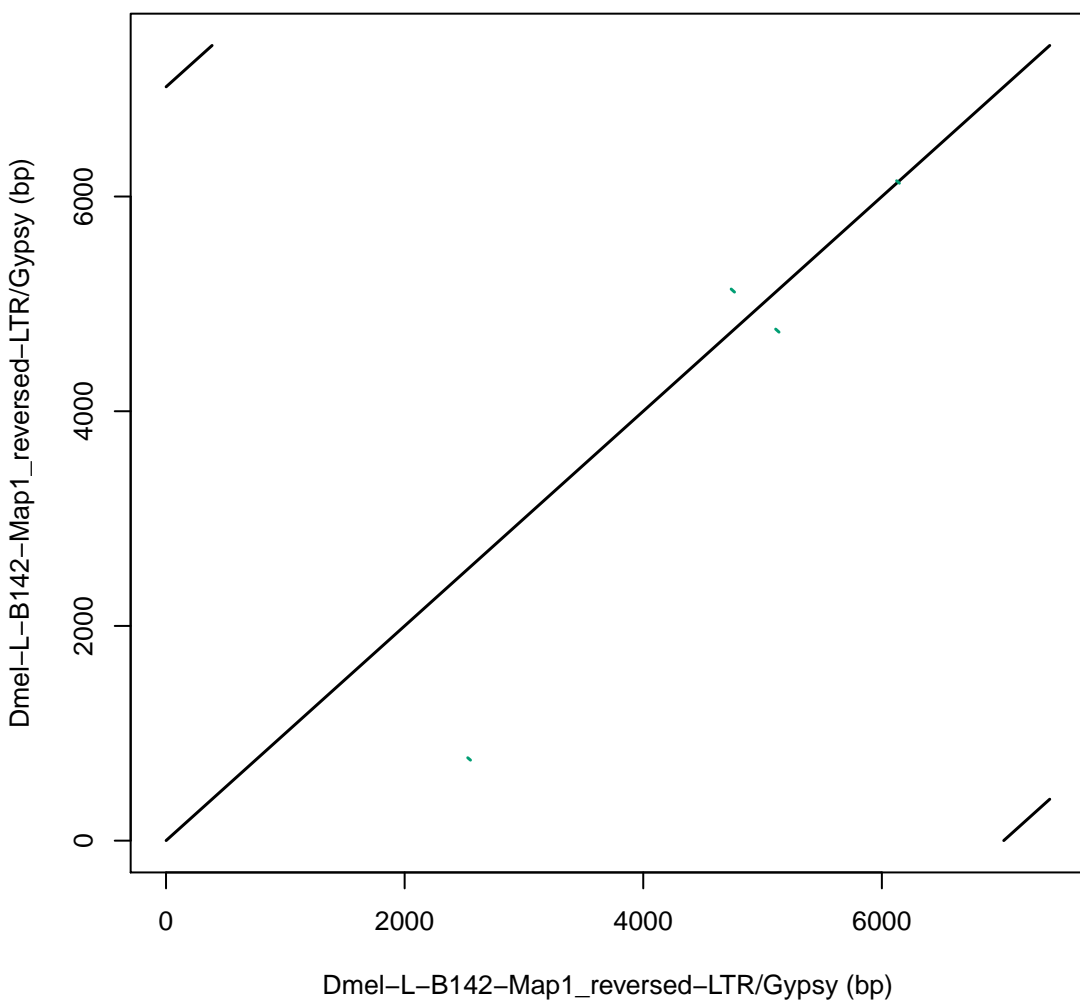
TE: Dmel-L-B142-Map1_reversed-LTR/Gypsy
consensus size: 7407bp; fragments: 52; full length: 1 (>=6666.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

