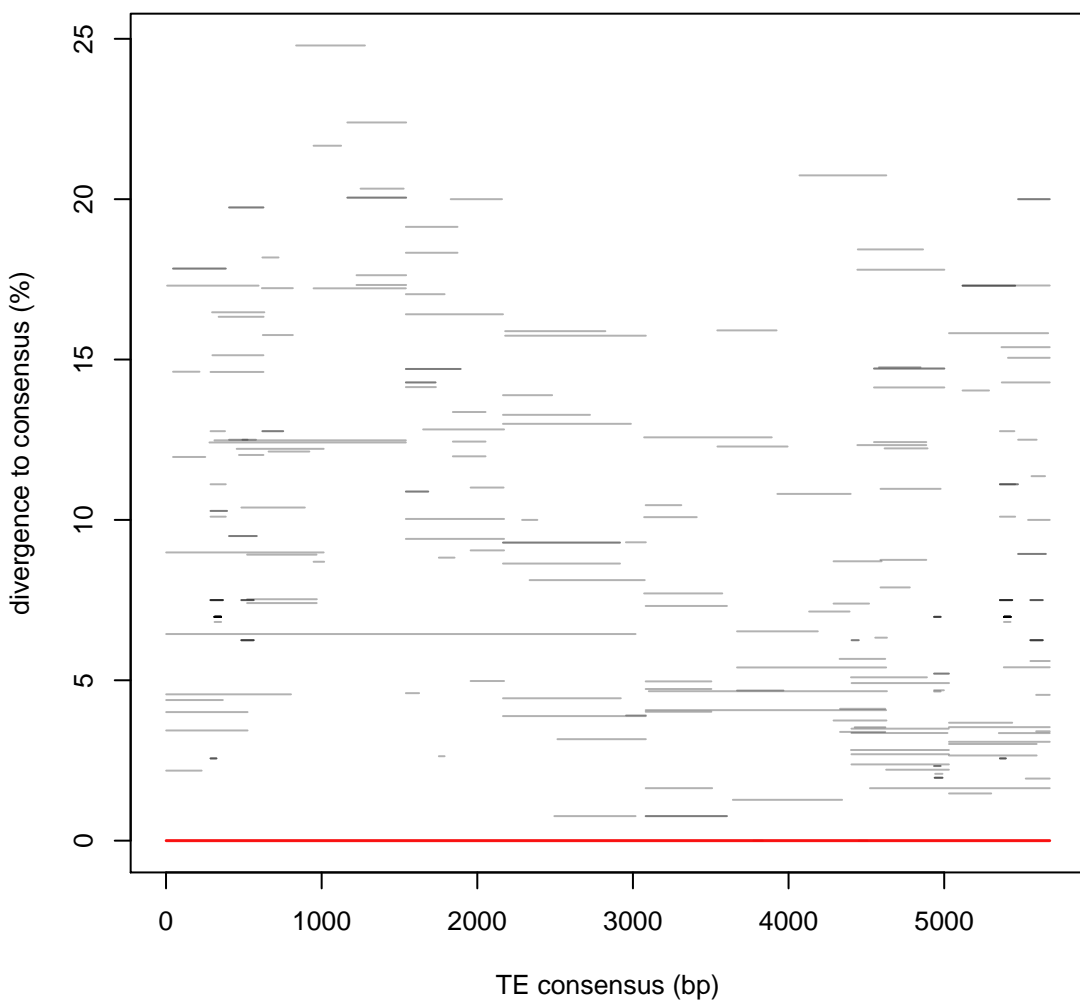
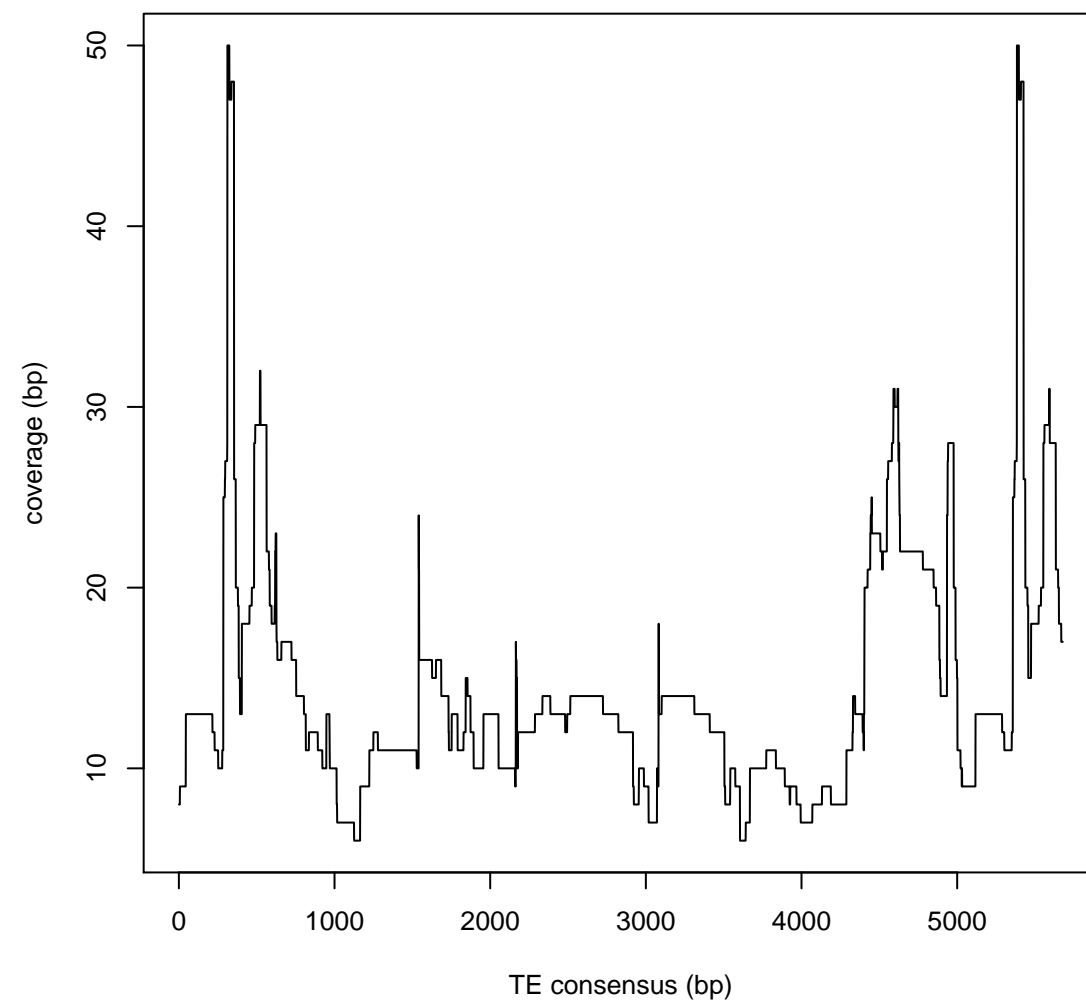


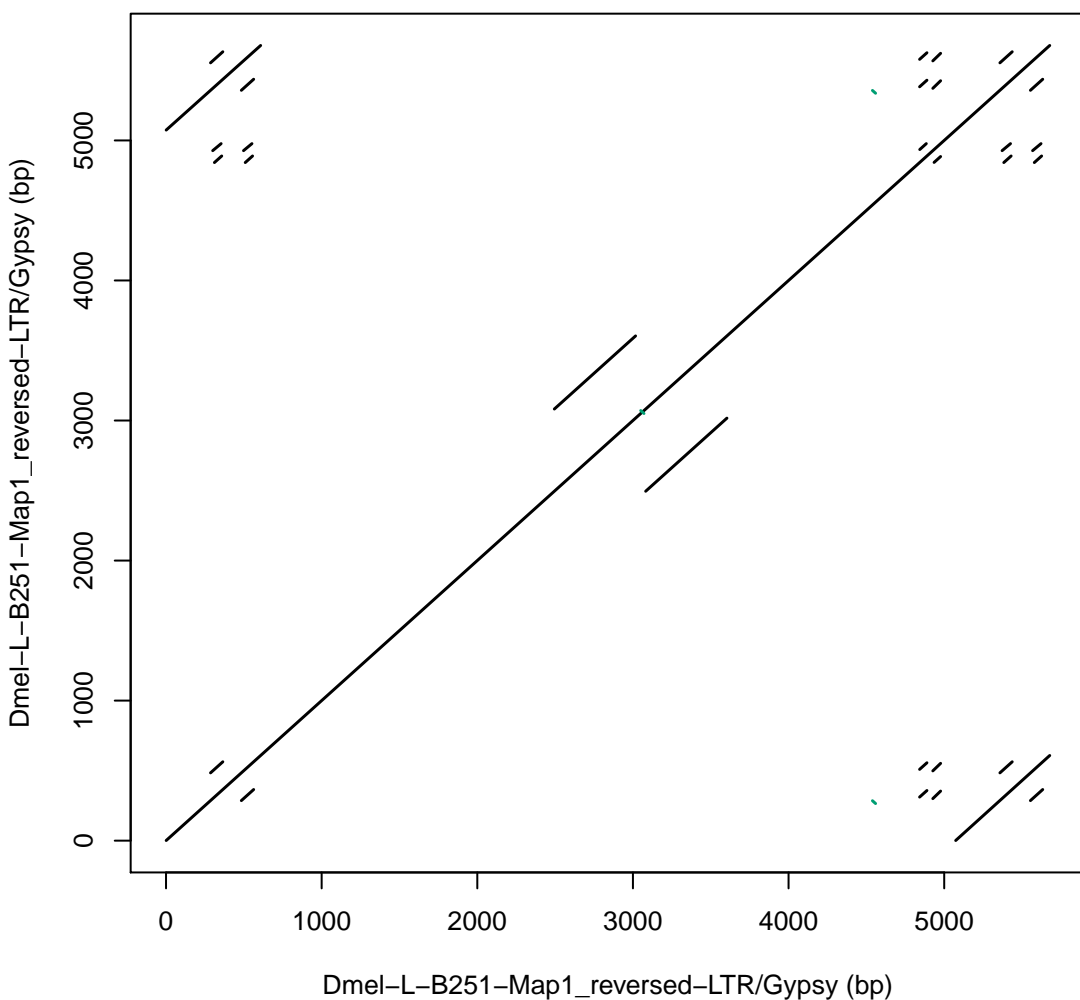
TE: Dmel-L-B251-Map1_reversed-LTR/Gypsy
consensus size: 5678bp; fragments: 267; full length: 1 (>=5110.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

