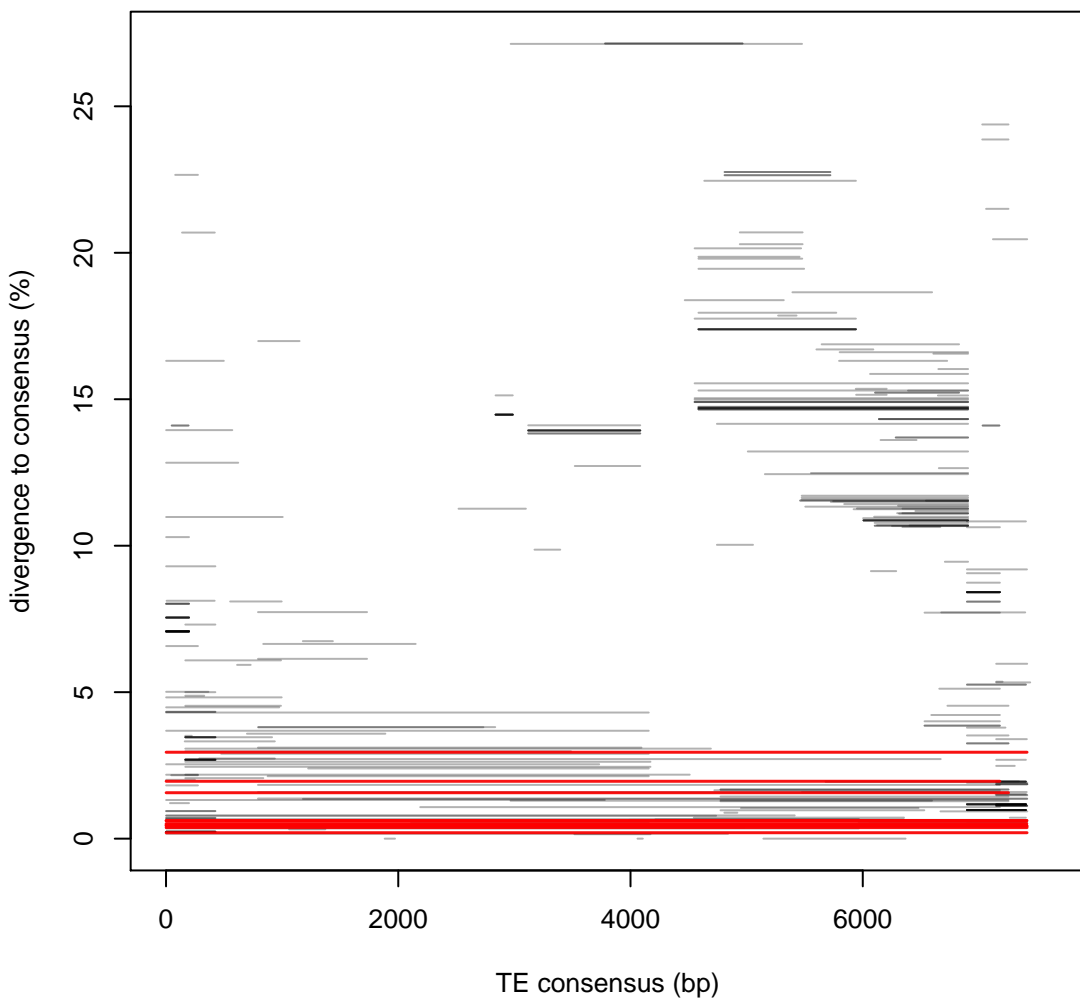
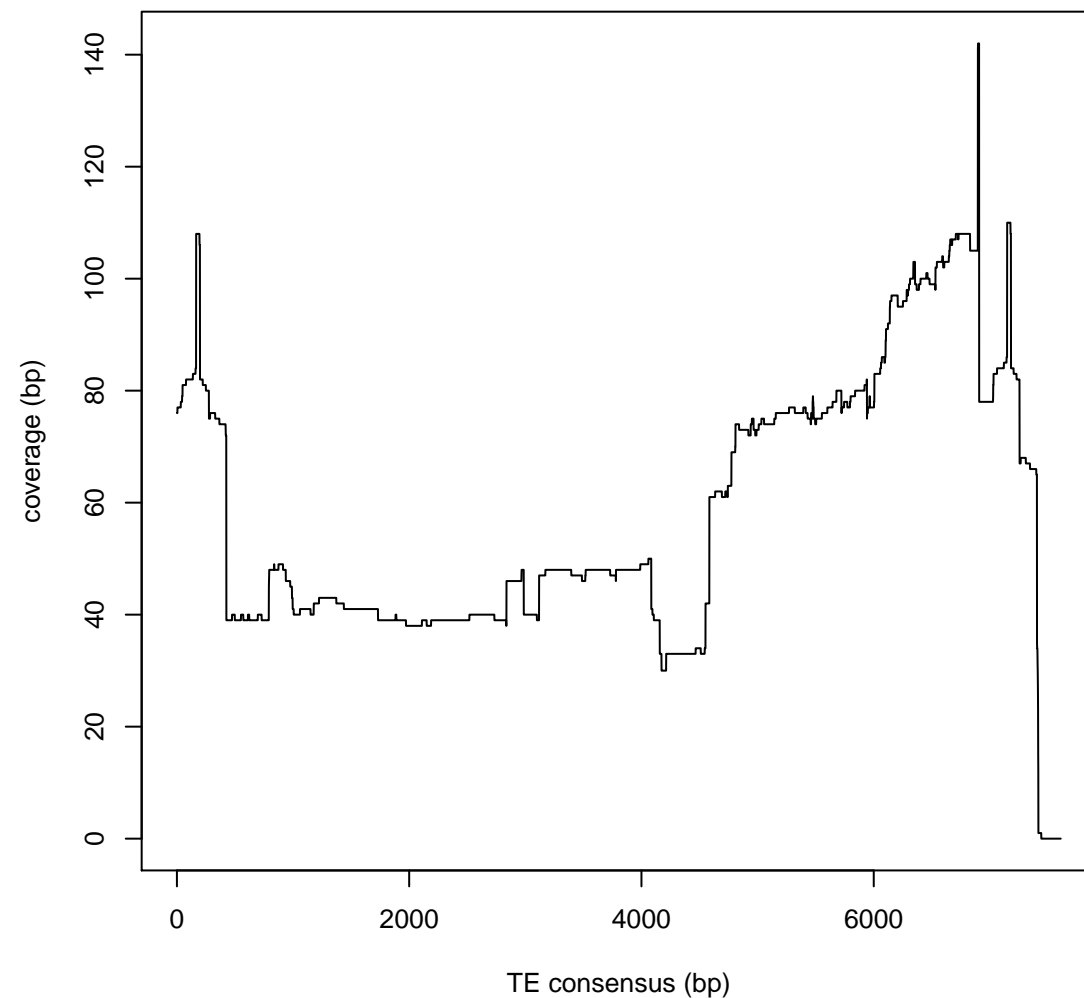


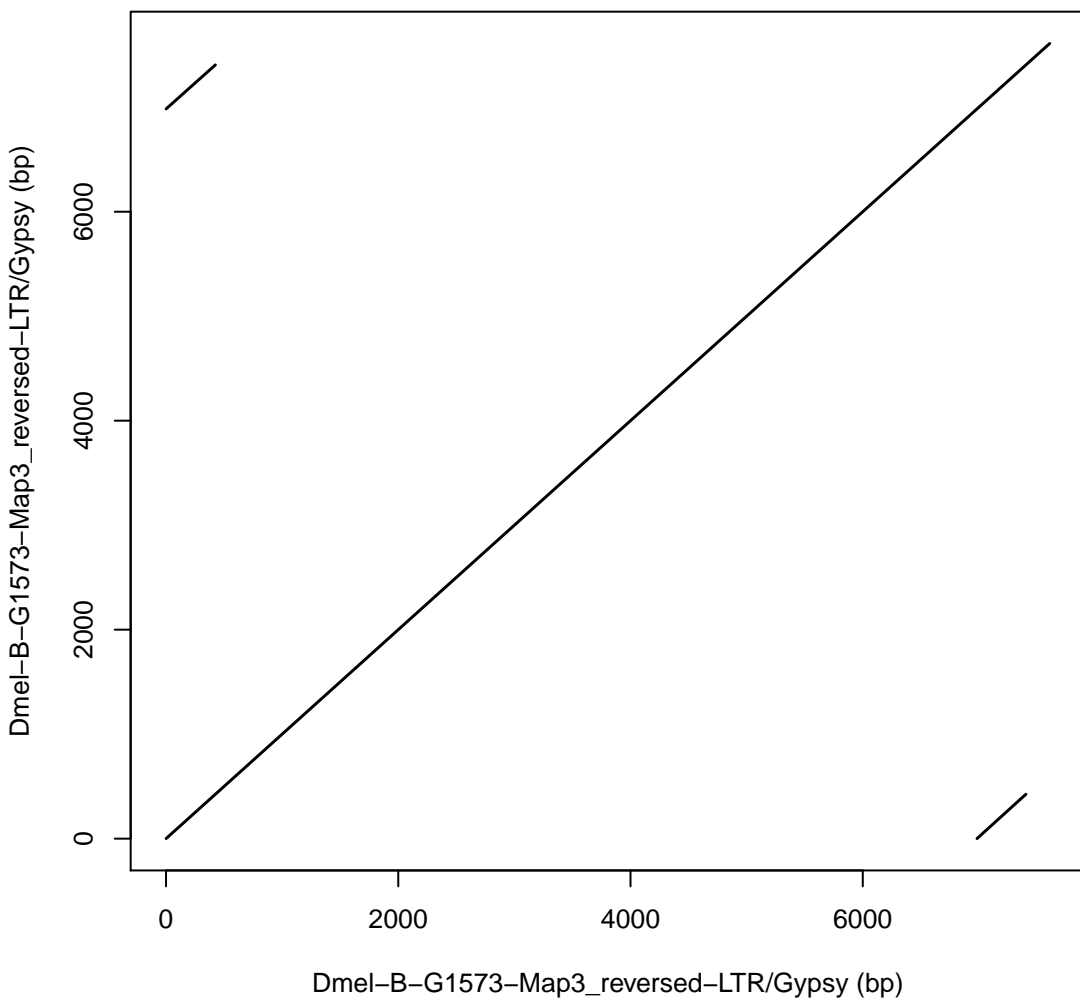
TE: Dmel-B-G1573-Map3_reversed-LTR/Gypsy
consensus size: 7610bp; fragments: 355; full length: 11 (>=6849bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

