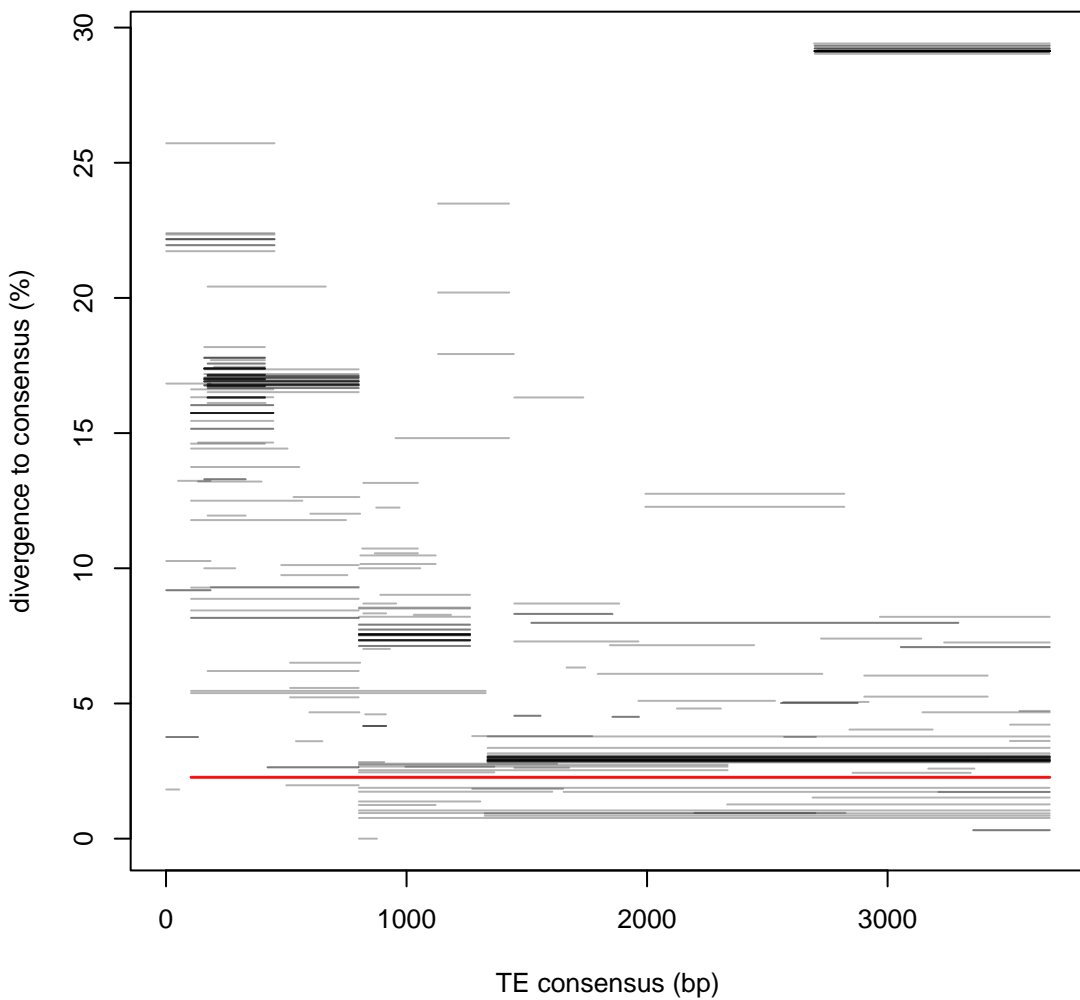
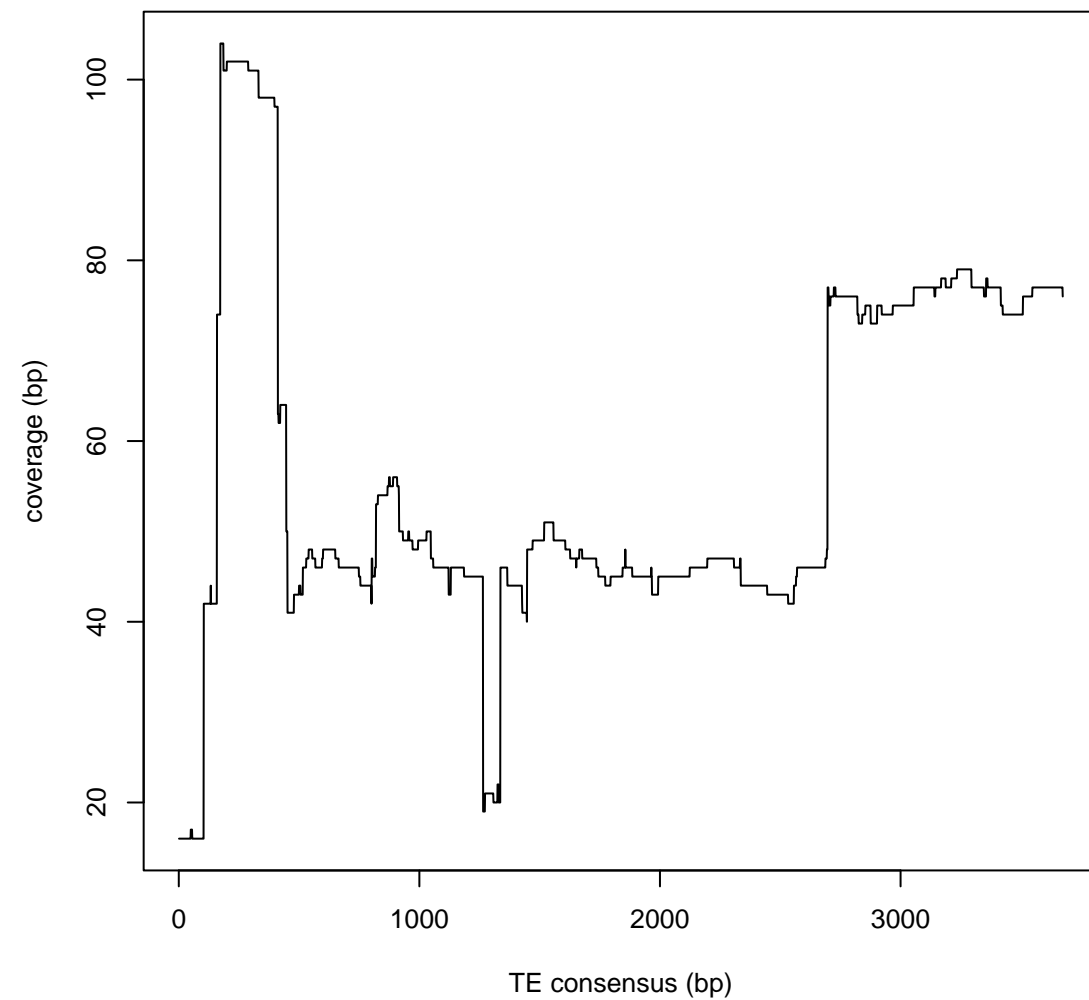


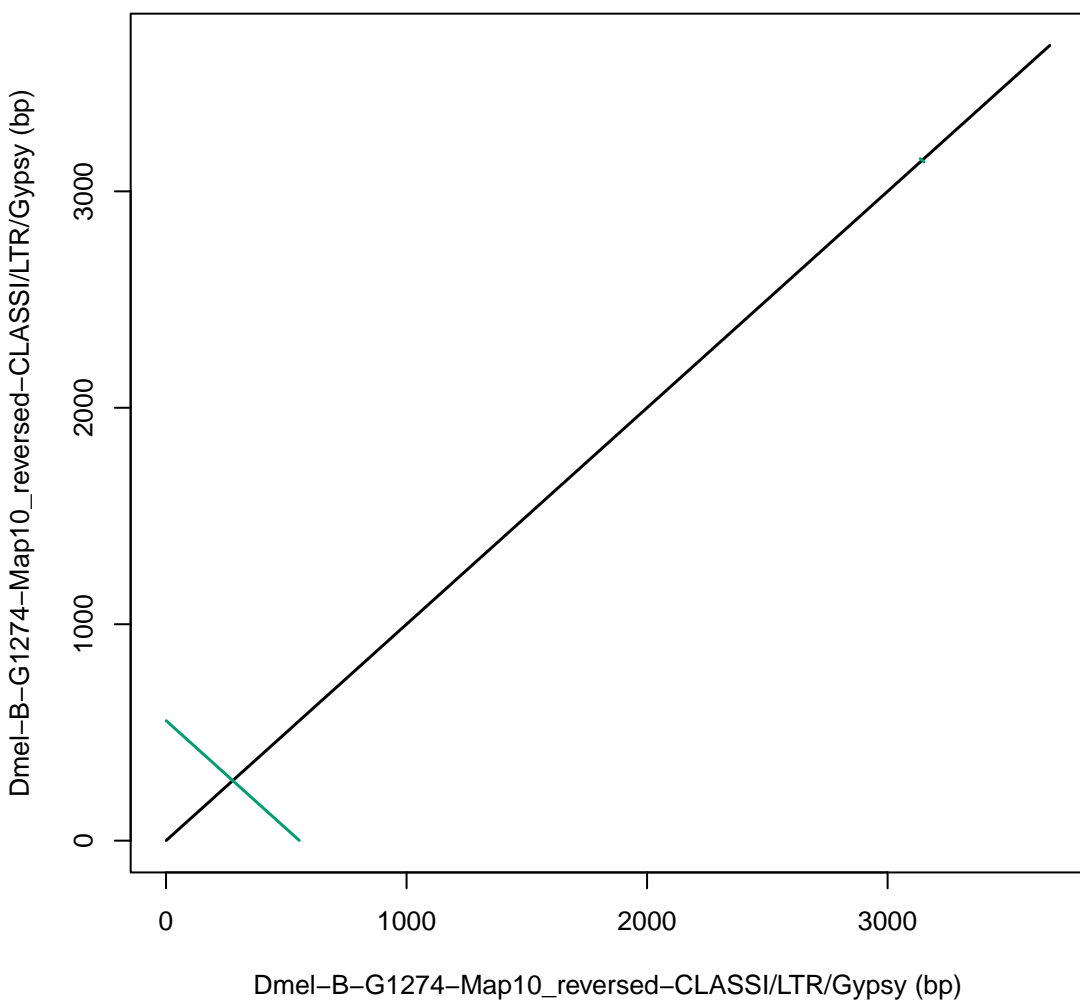
TE: Dmel-B-G1274-Map10_reversed-CLASSI/LTR/Gypsy
consensus size: 3674bp; fragments: 288; full length: 1 (>=3306.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

