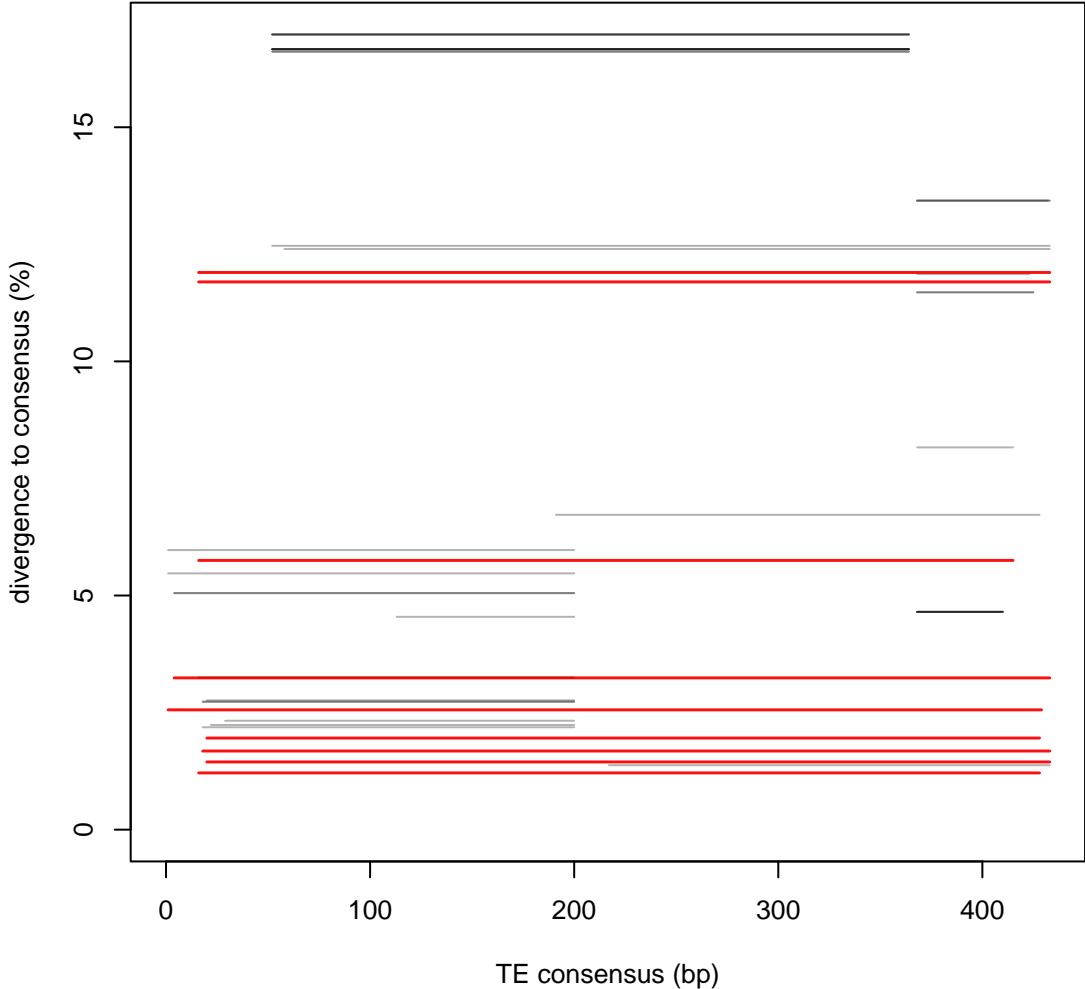
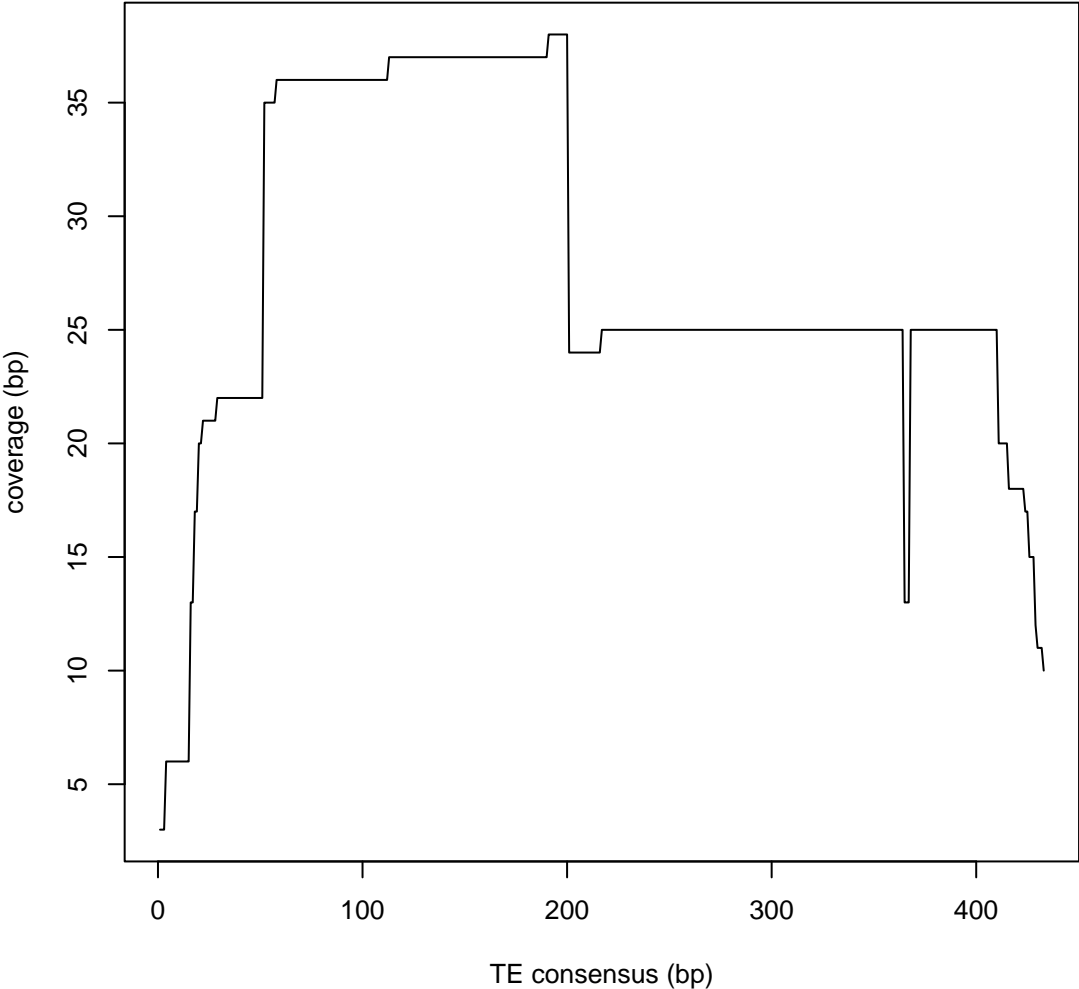


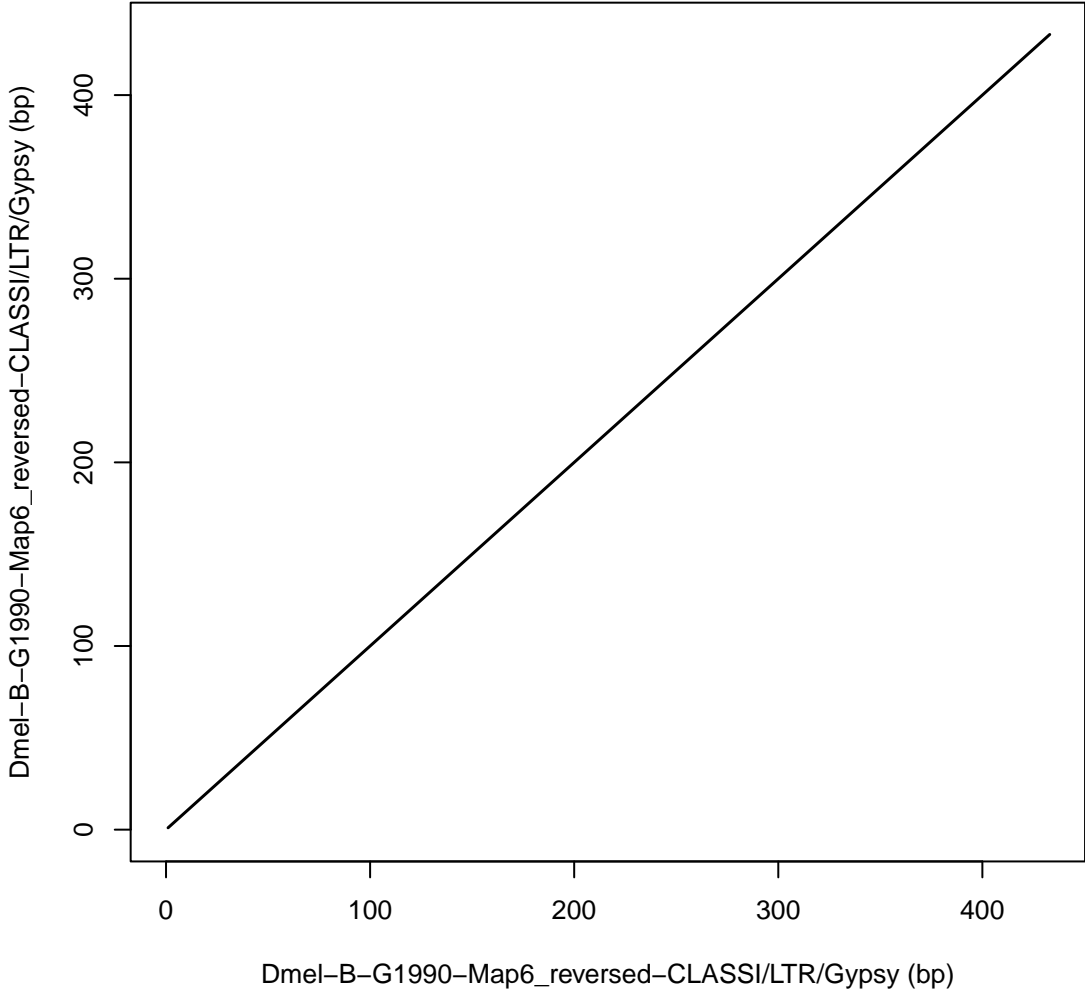
TE: Dmel-B-G1990-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 433bp; fragments: 51; full length: 9 (>=389.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

