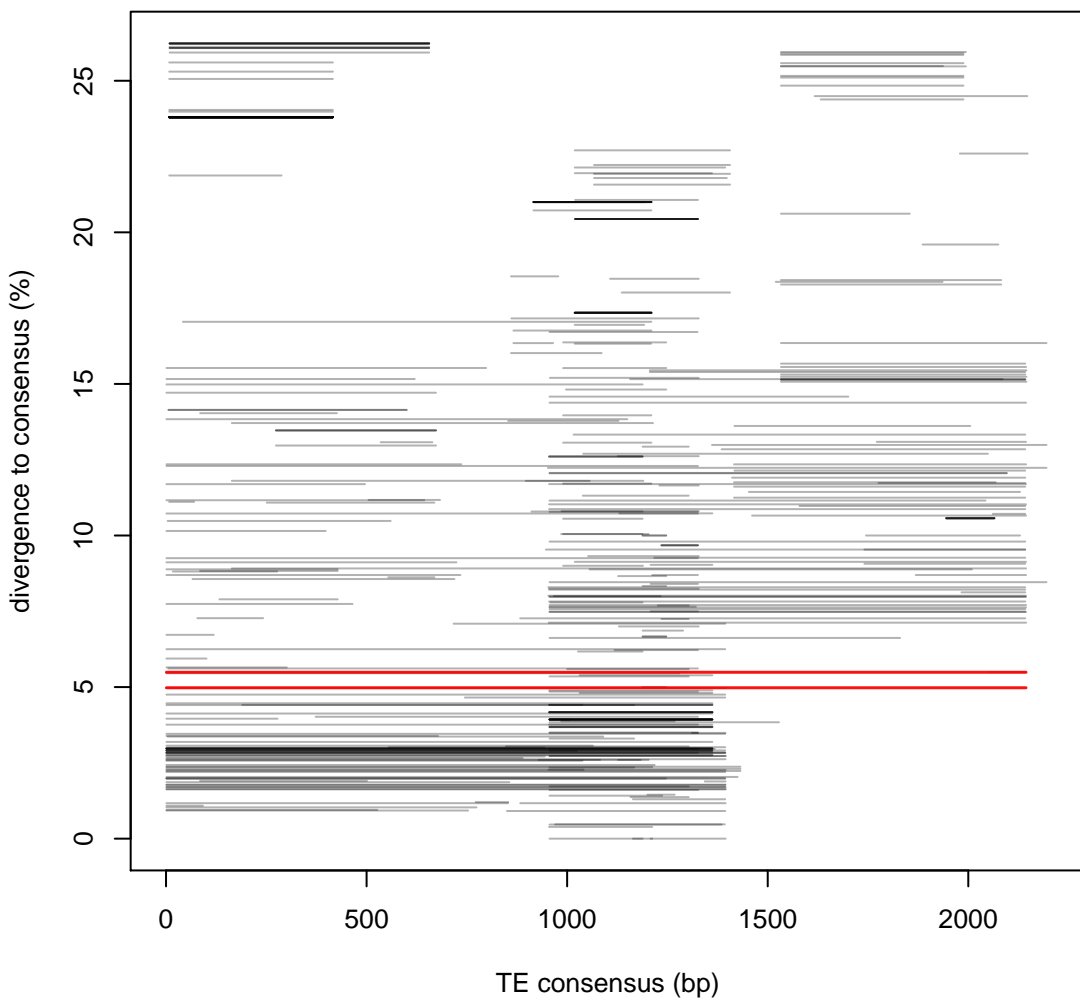
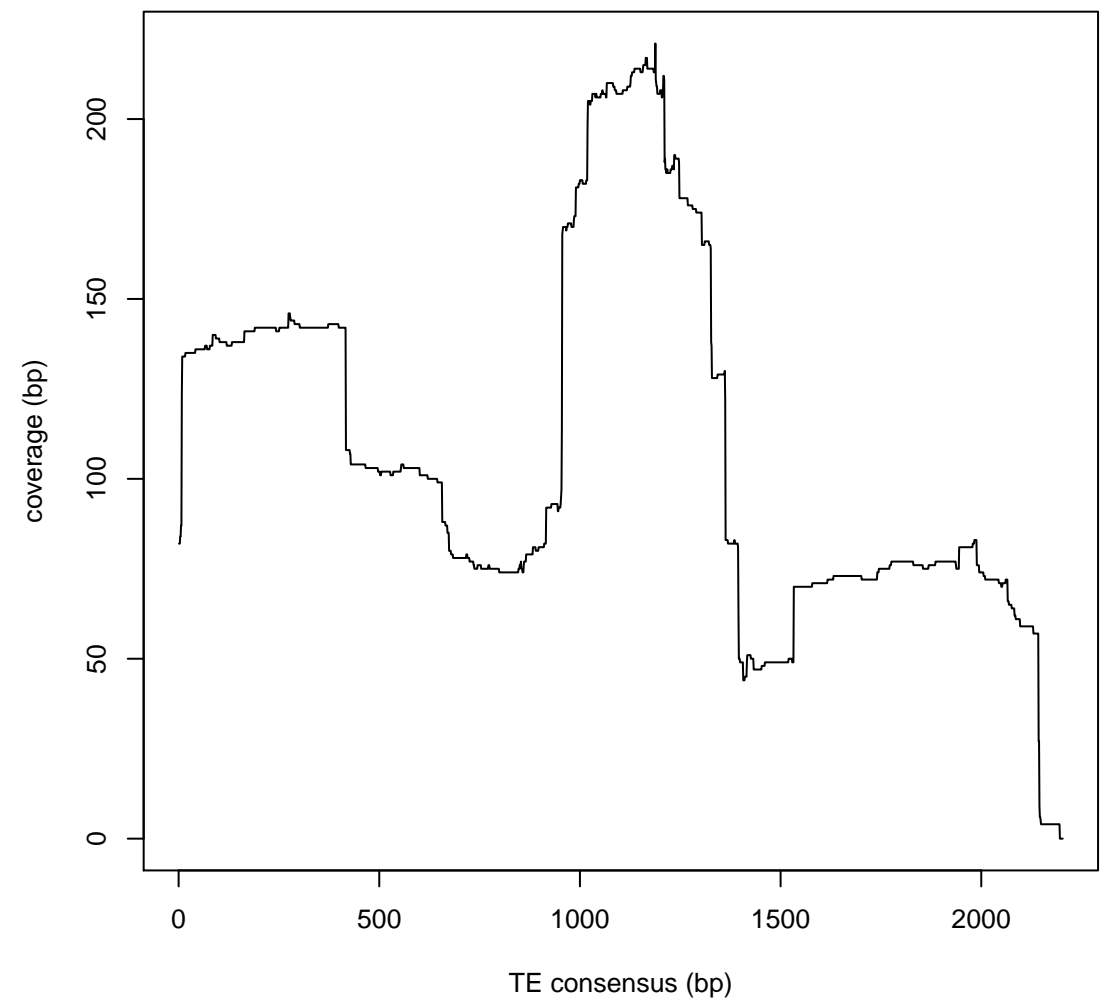


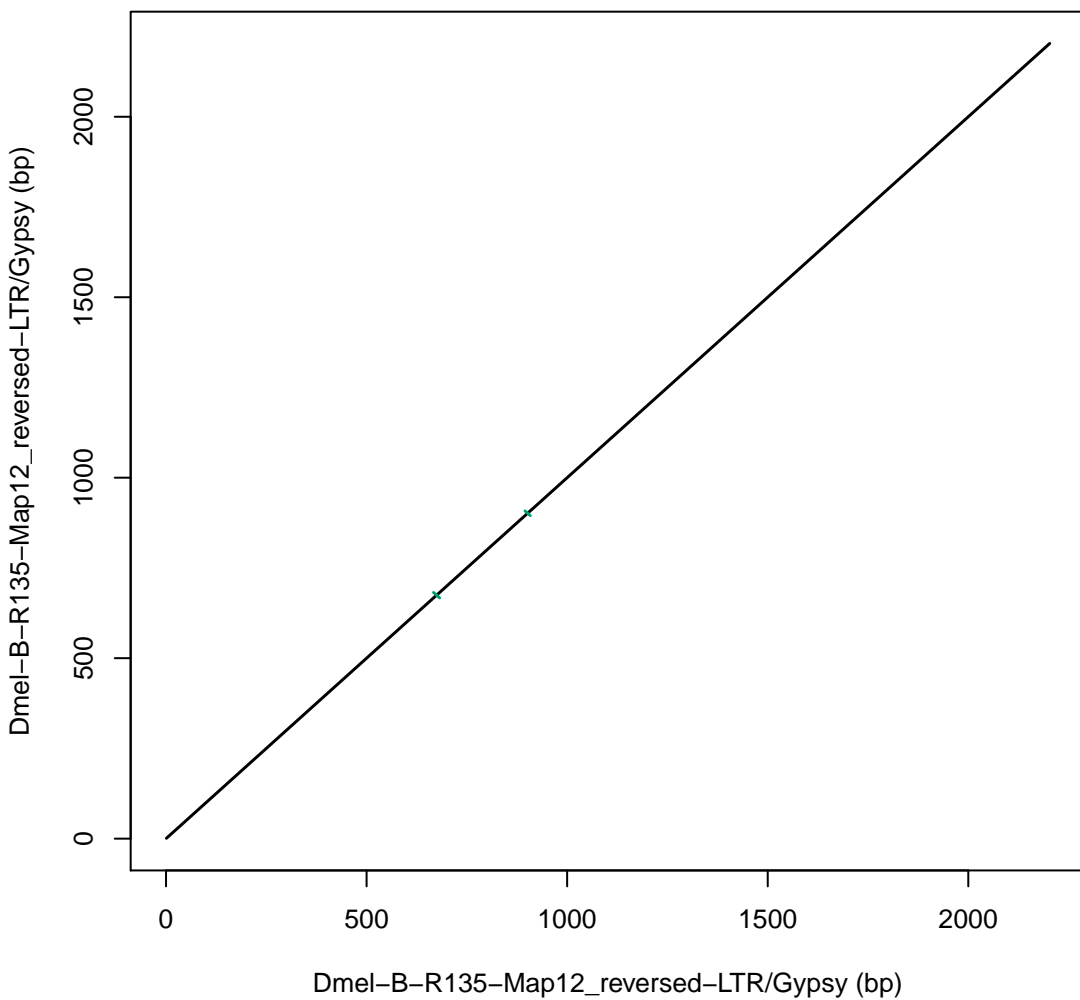
TE: Dmel-B-R135-Map12_reversed-LTR/Gypsy
consensus size: 2203bp; fragments: 400; full length: 2 (≥ 1982.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

