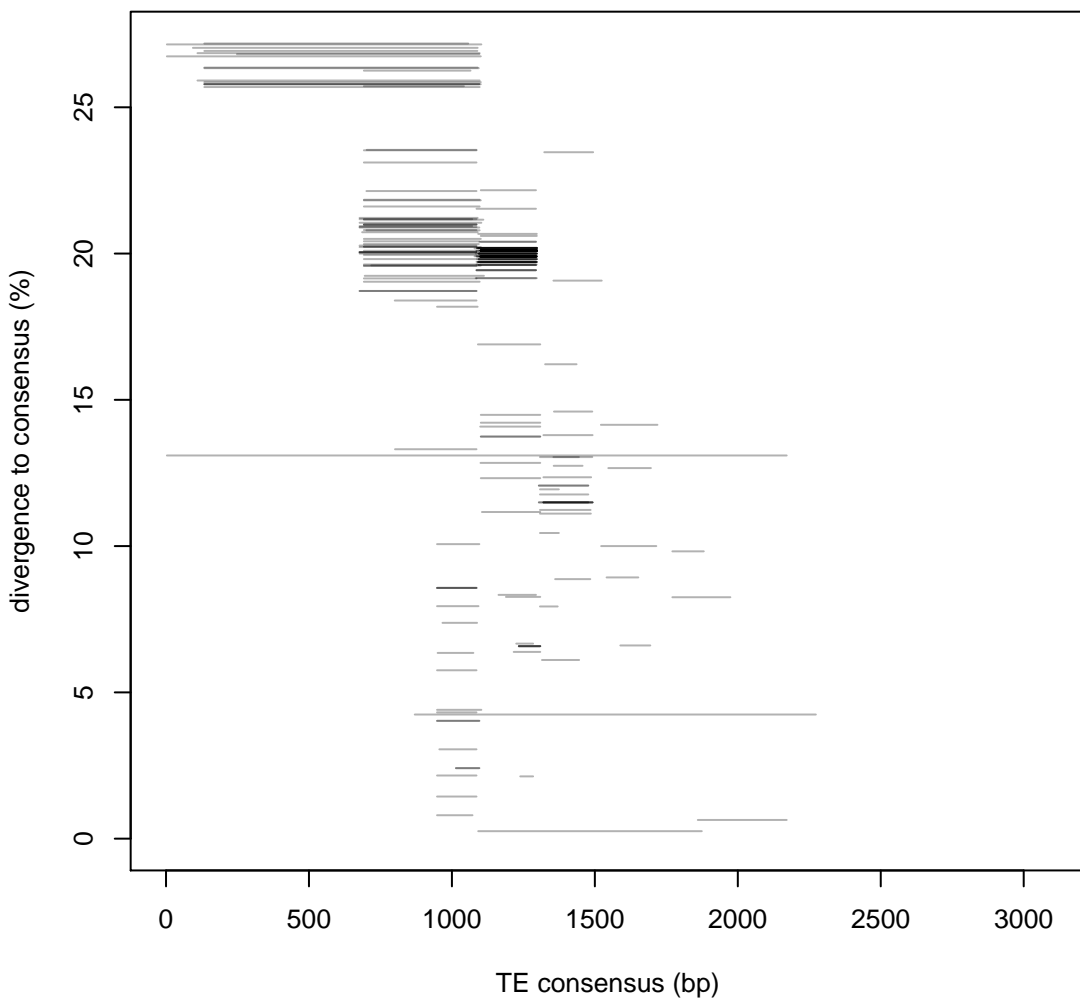
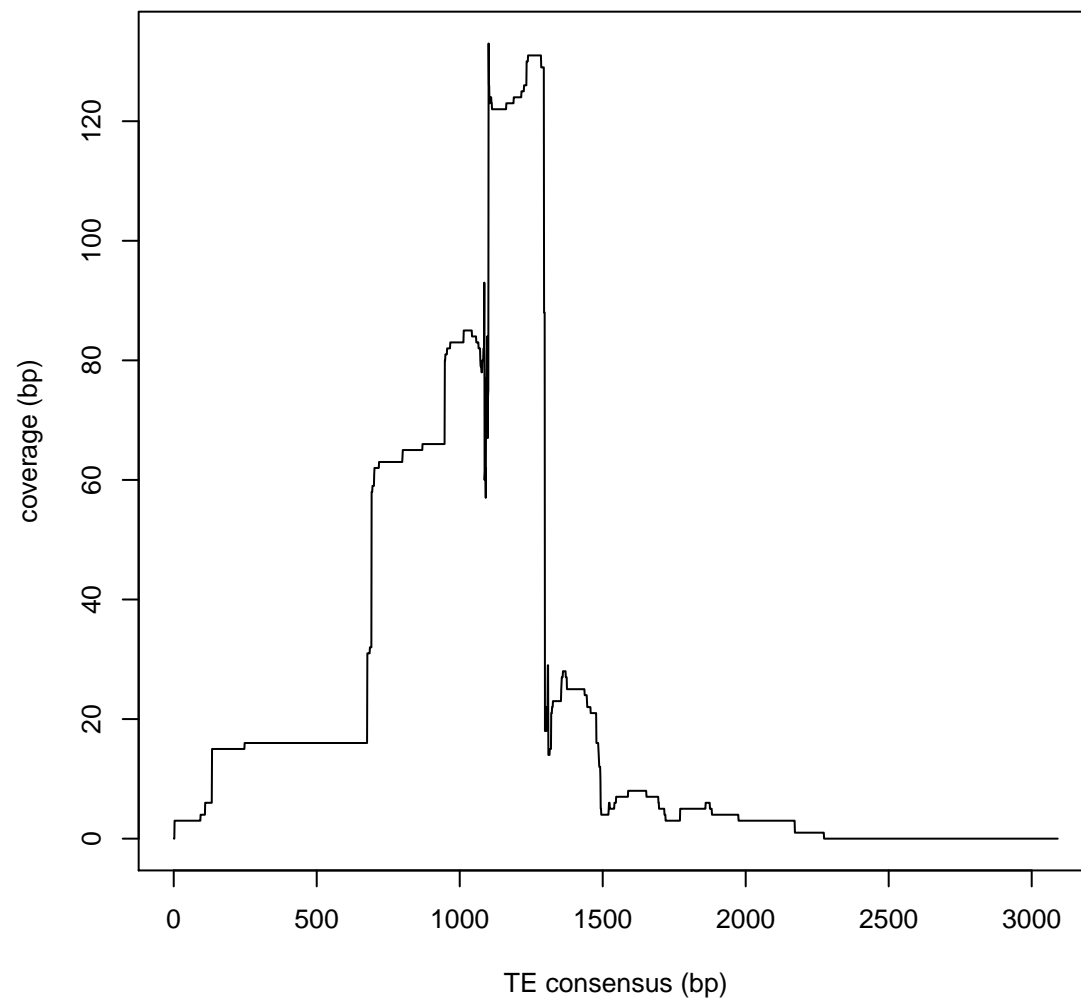


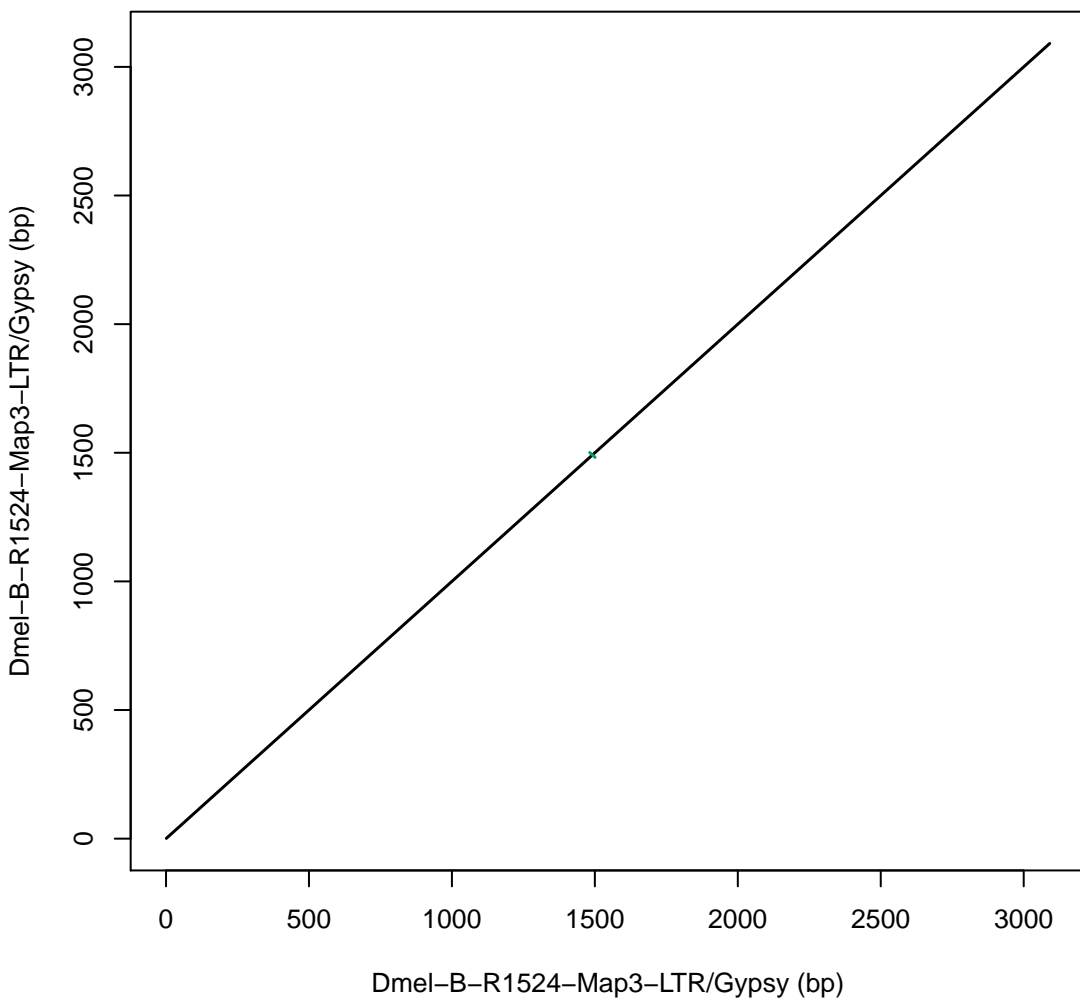
TE: Dmel-B-R1524-Map3-LTR/Gypsy
consensus size: 3091bp; fragments: 247; full length: 0 (>=2781.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

