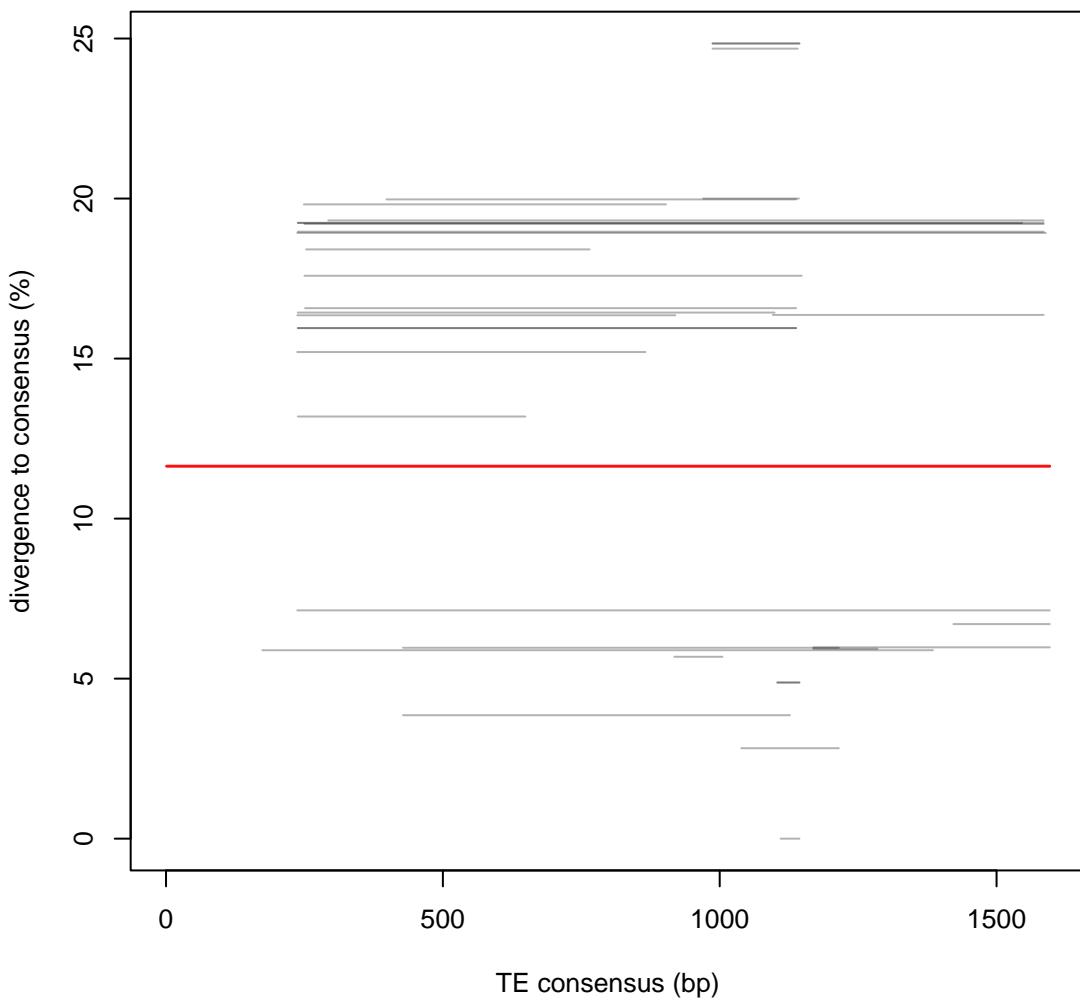
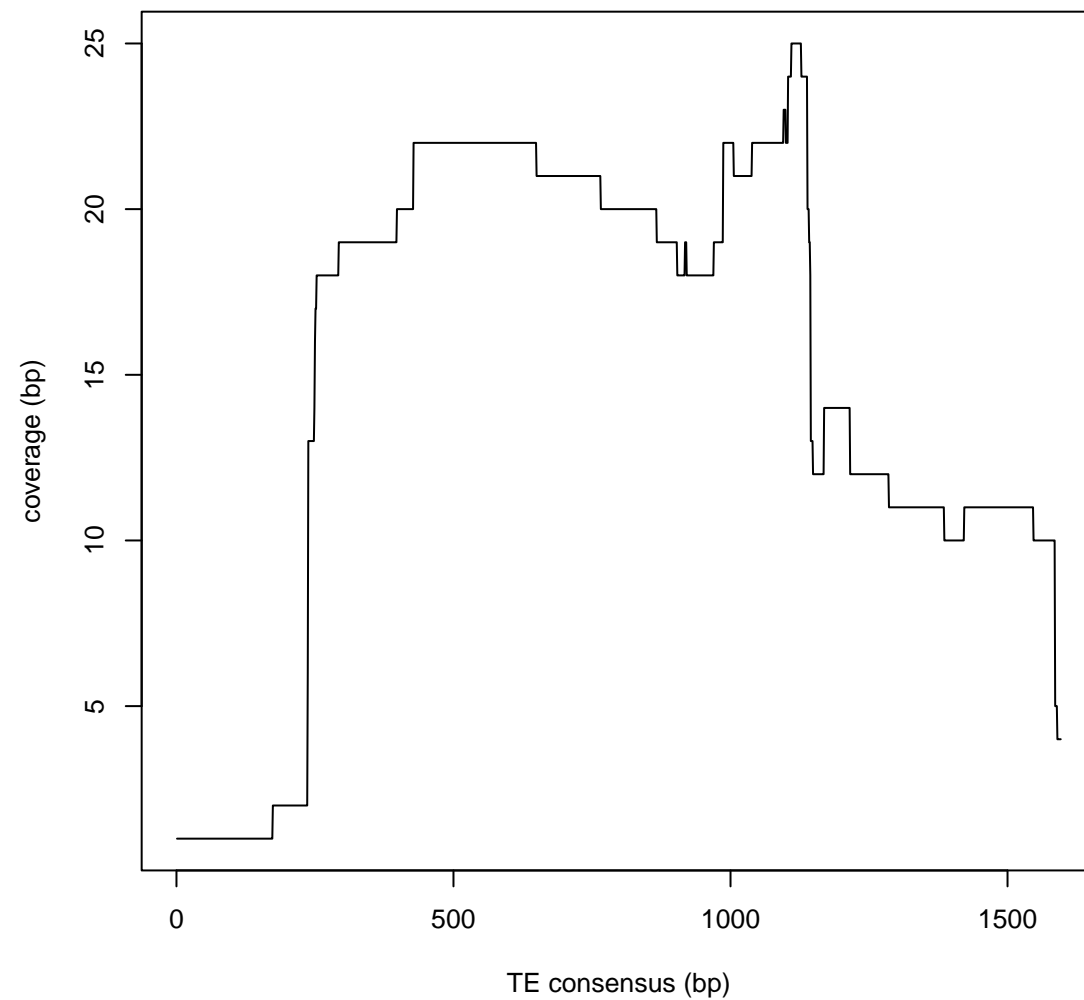


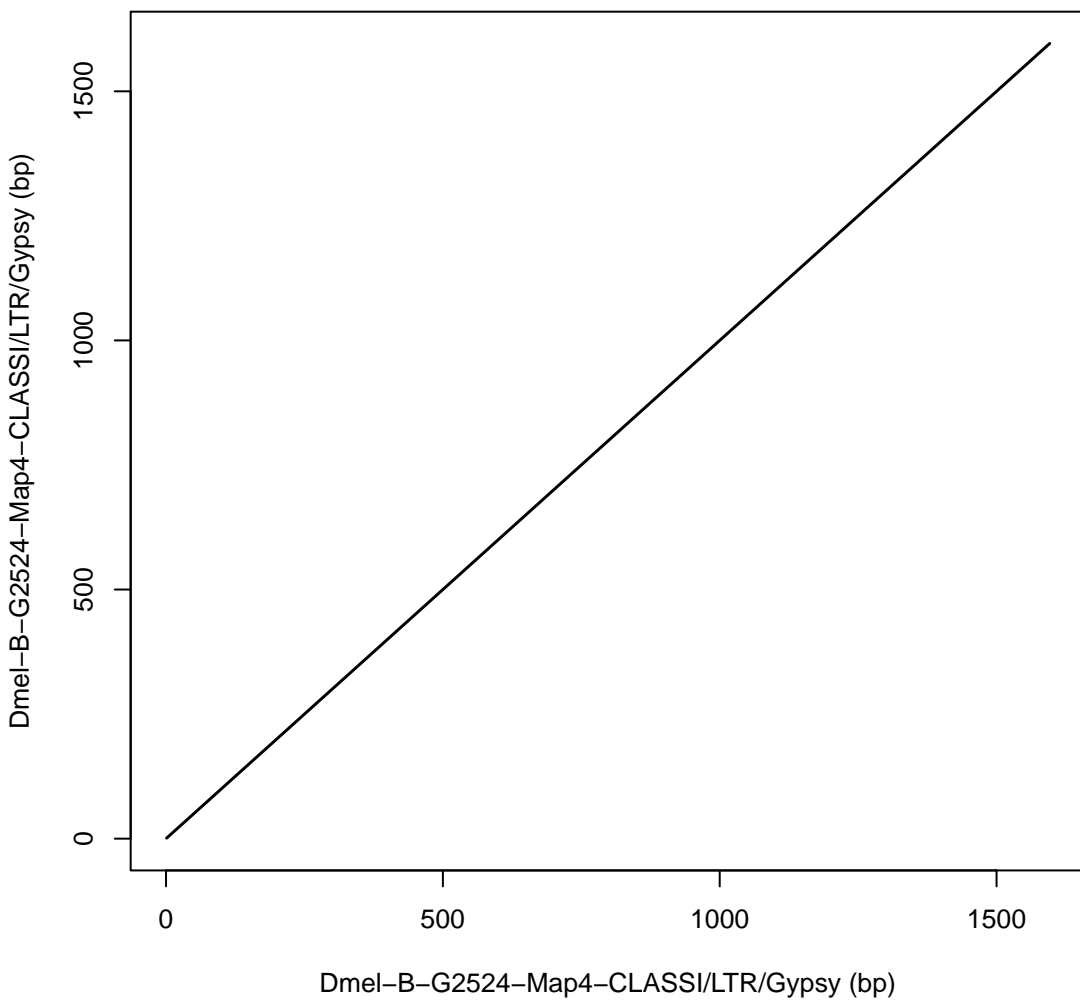
TE: Dmel-B-G2524-Map4-CLASSI/LTR/Gypsy
consensus size: 1596bp; fragments: 35; full length: 1 (>=1436.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

