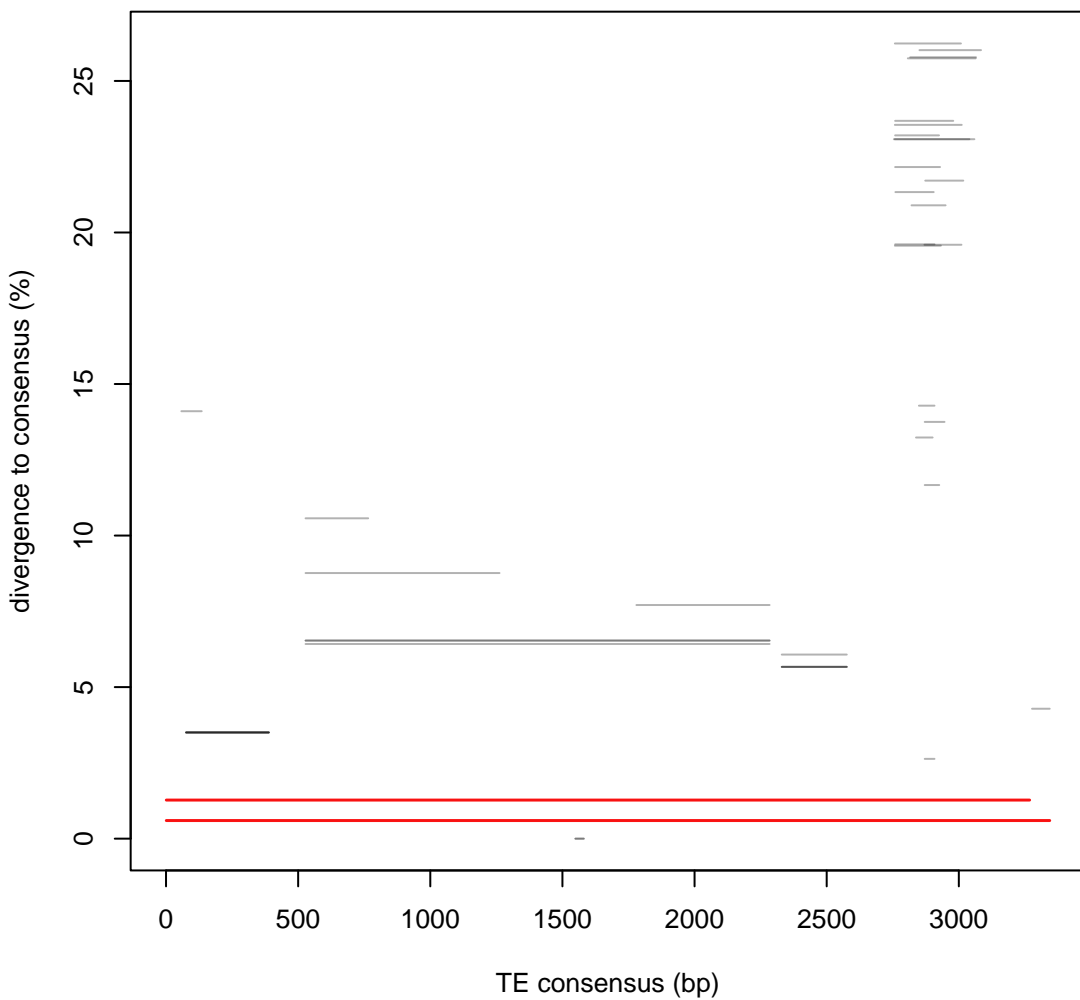
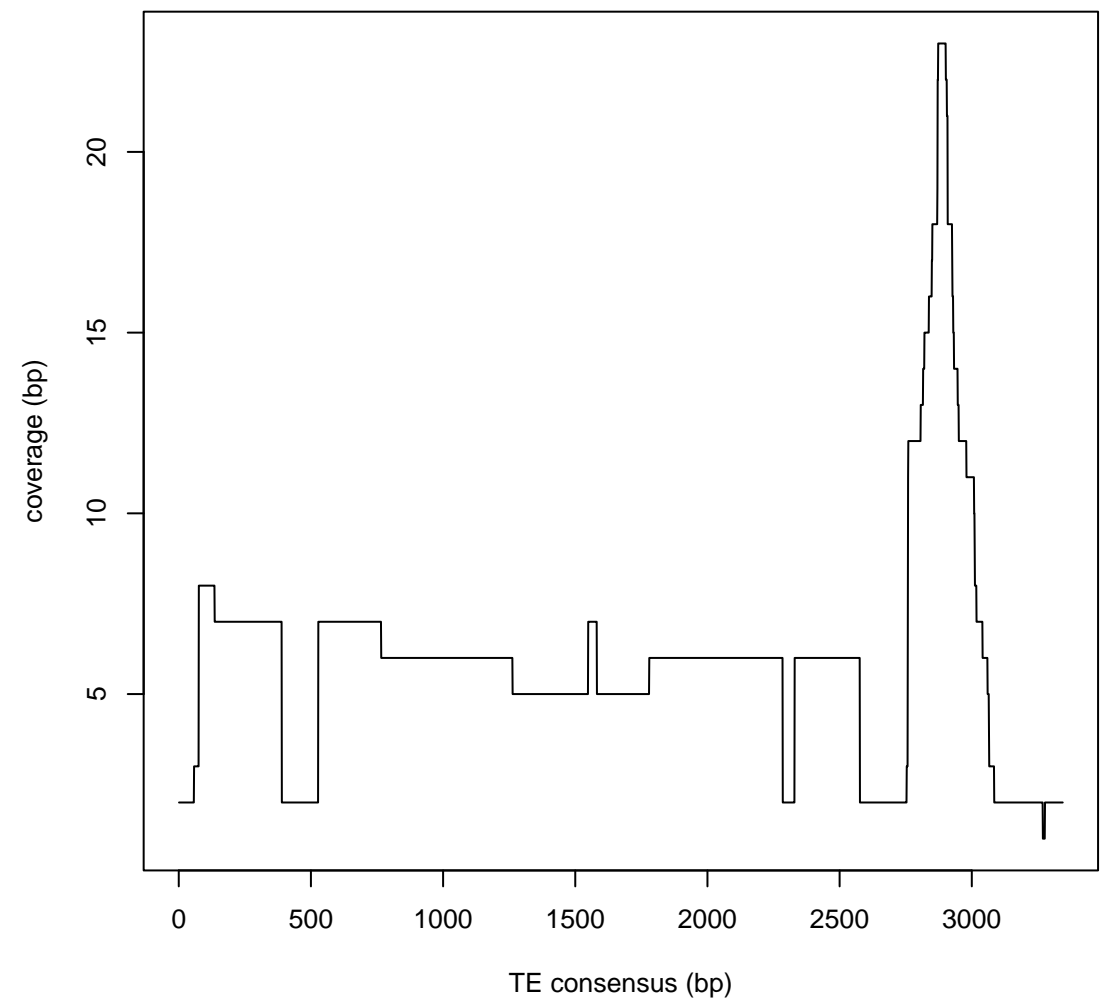


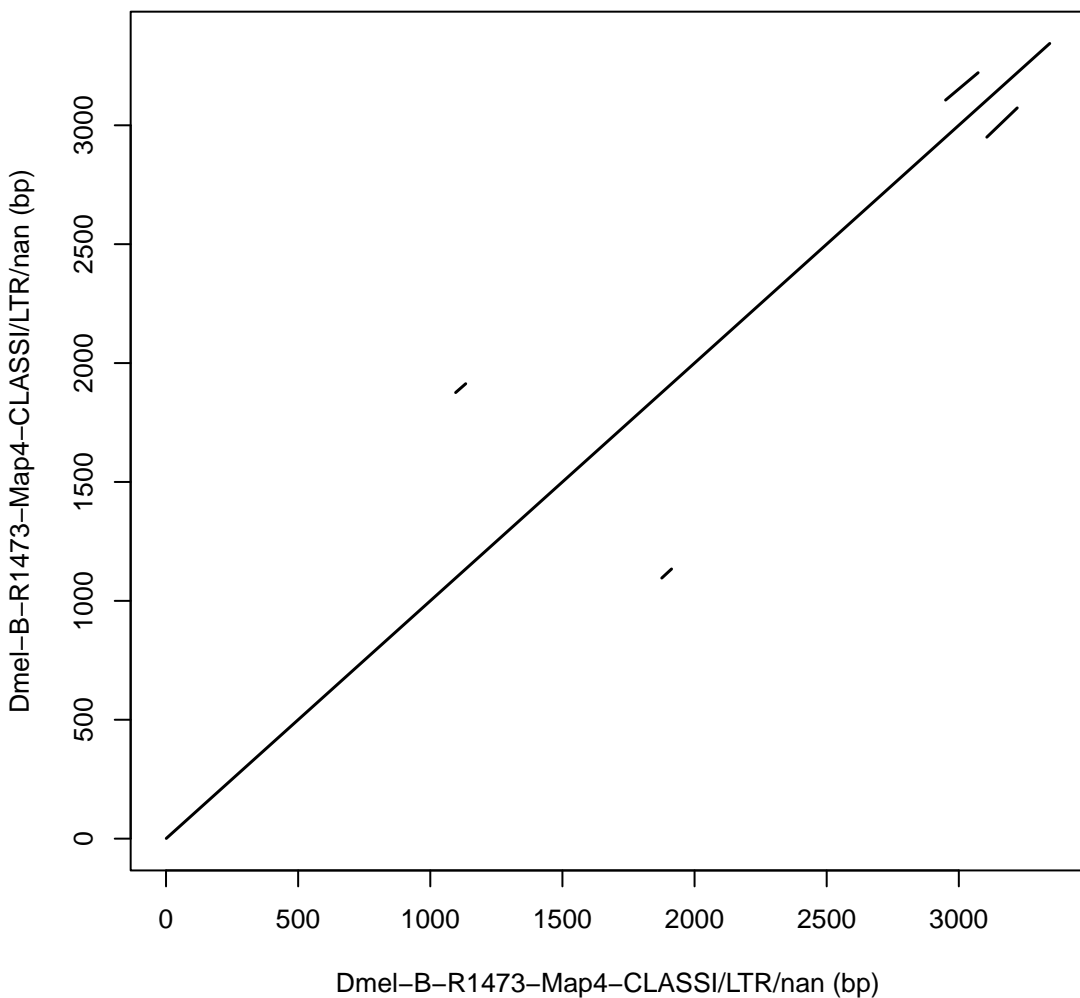
TE: Dmel-B-R1473-Map4-CLASSI/LTR/nan
consensus size: 3344bp; fragments: 42; full length: 2 (>=3009.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

