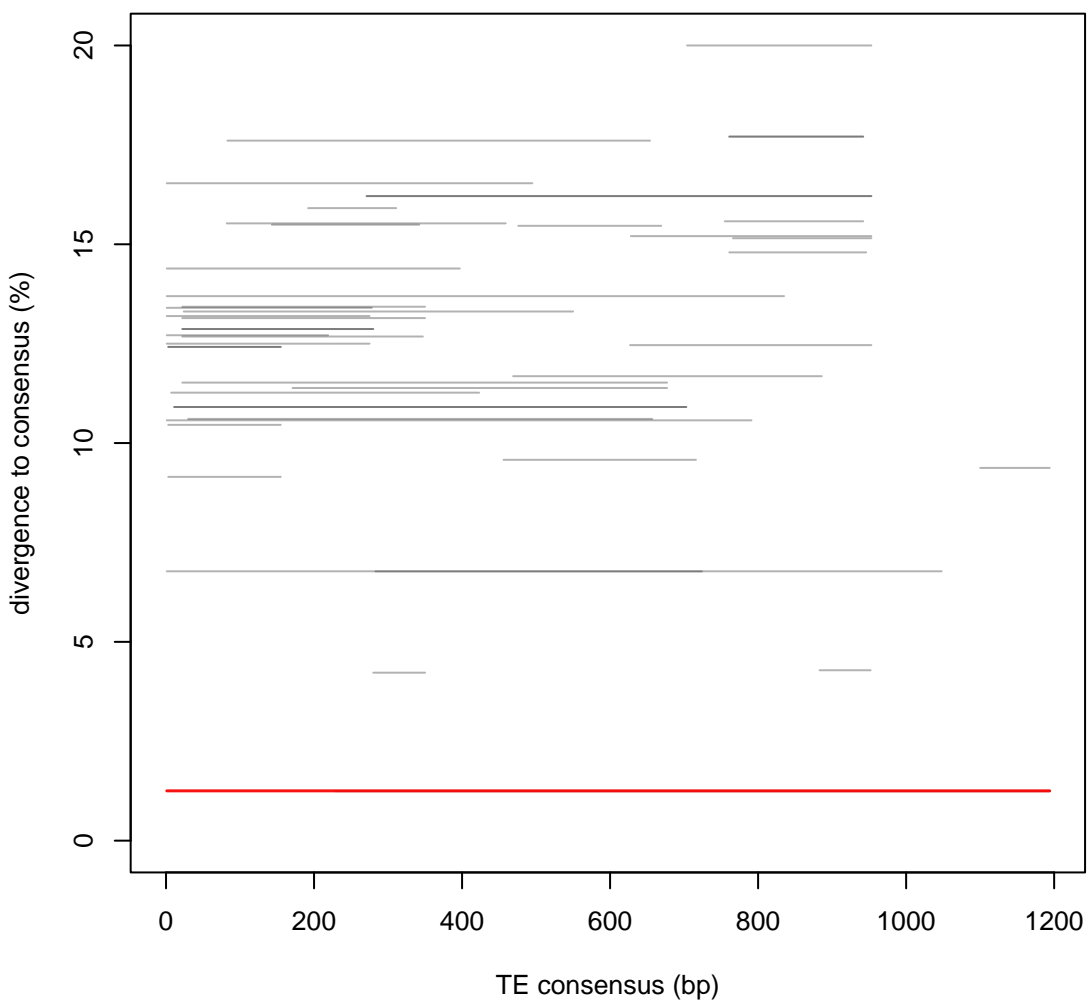
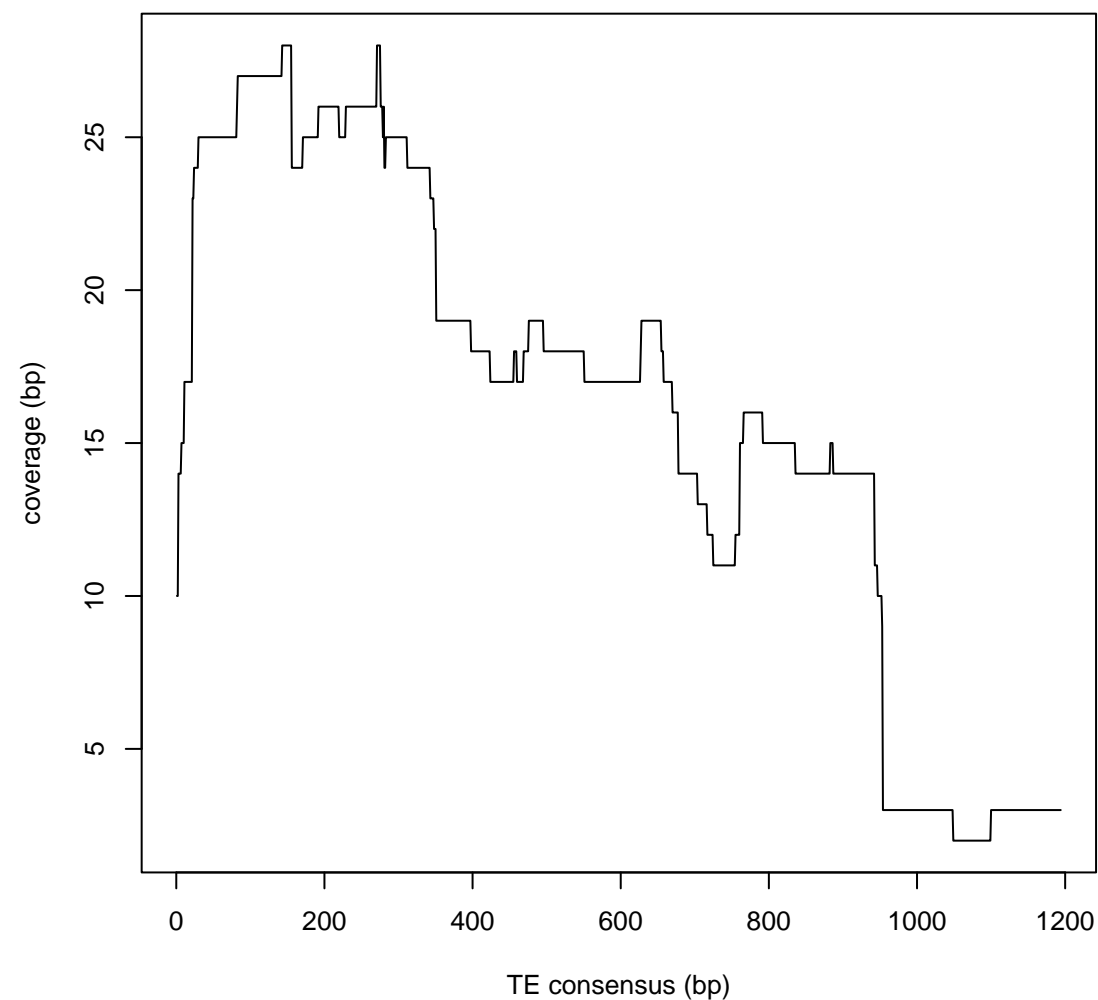


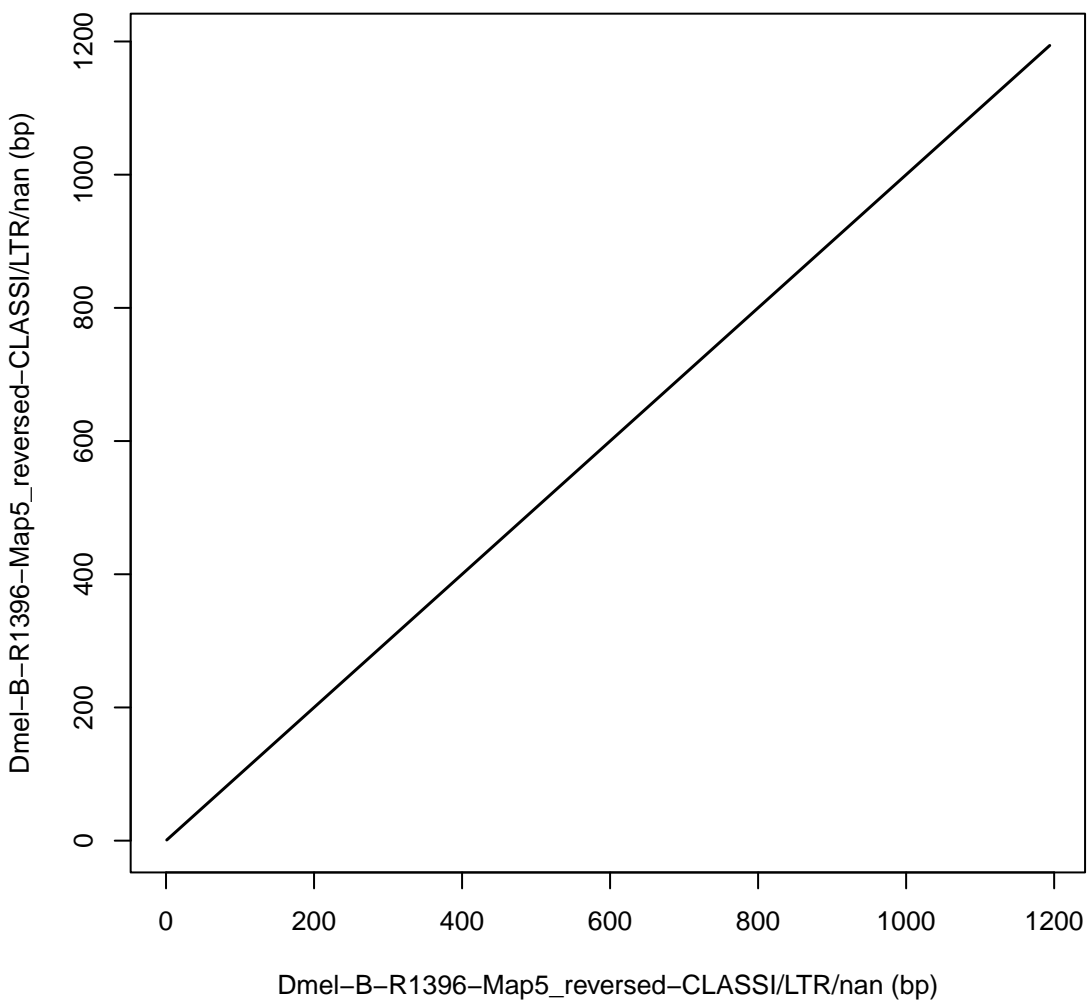
TE: Dmel-B-R1396-Map5_reversed-CLASSI/LTR/nan
consensus size: 1194bp; fragments: 48; full length: 1 (>=1074.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

