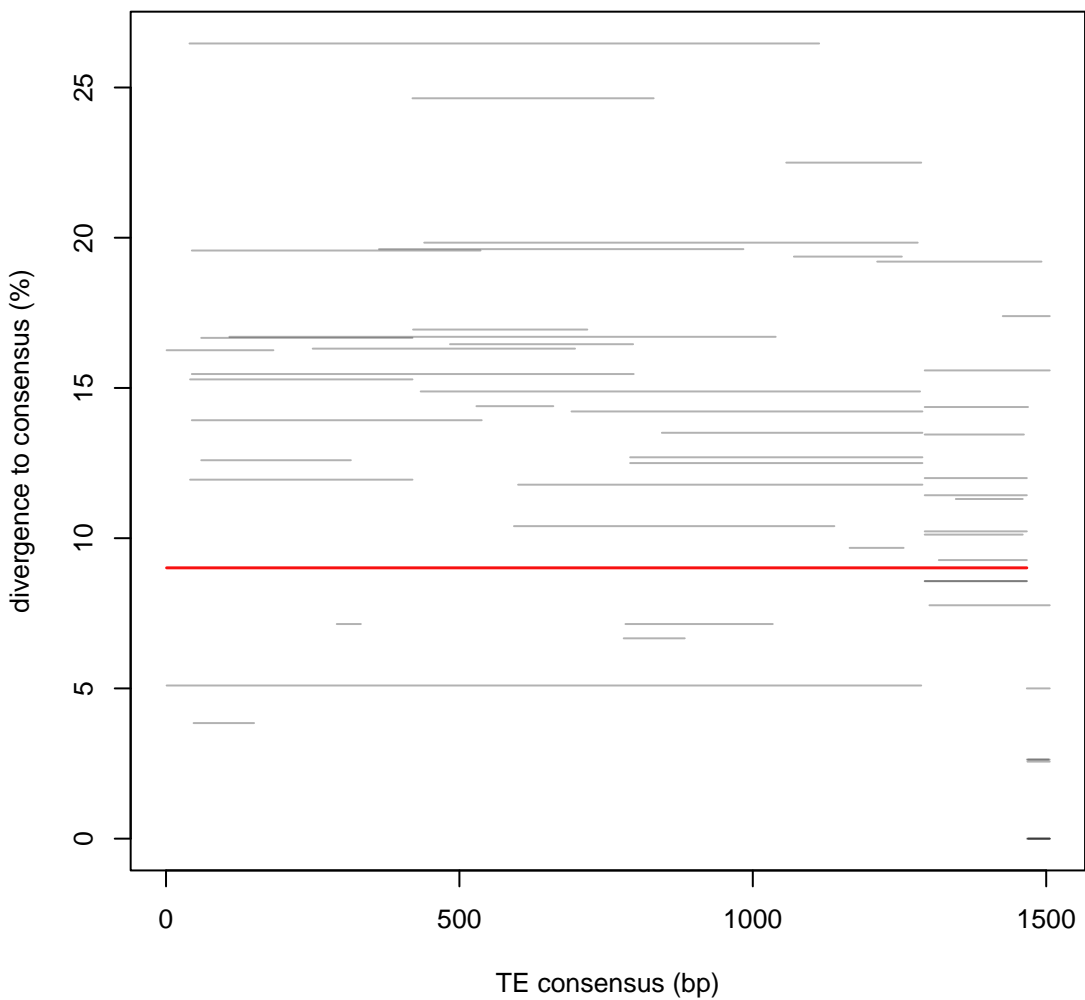
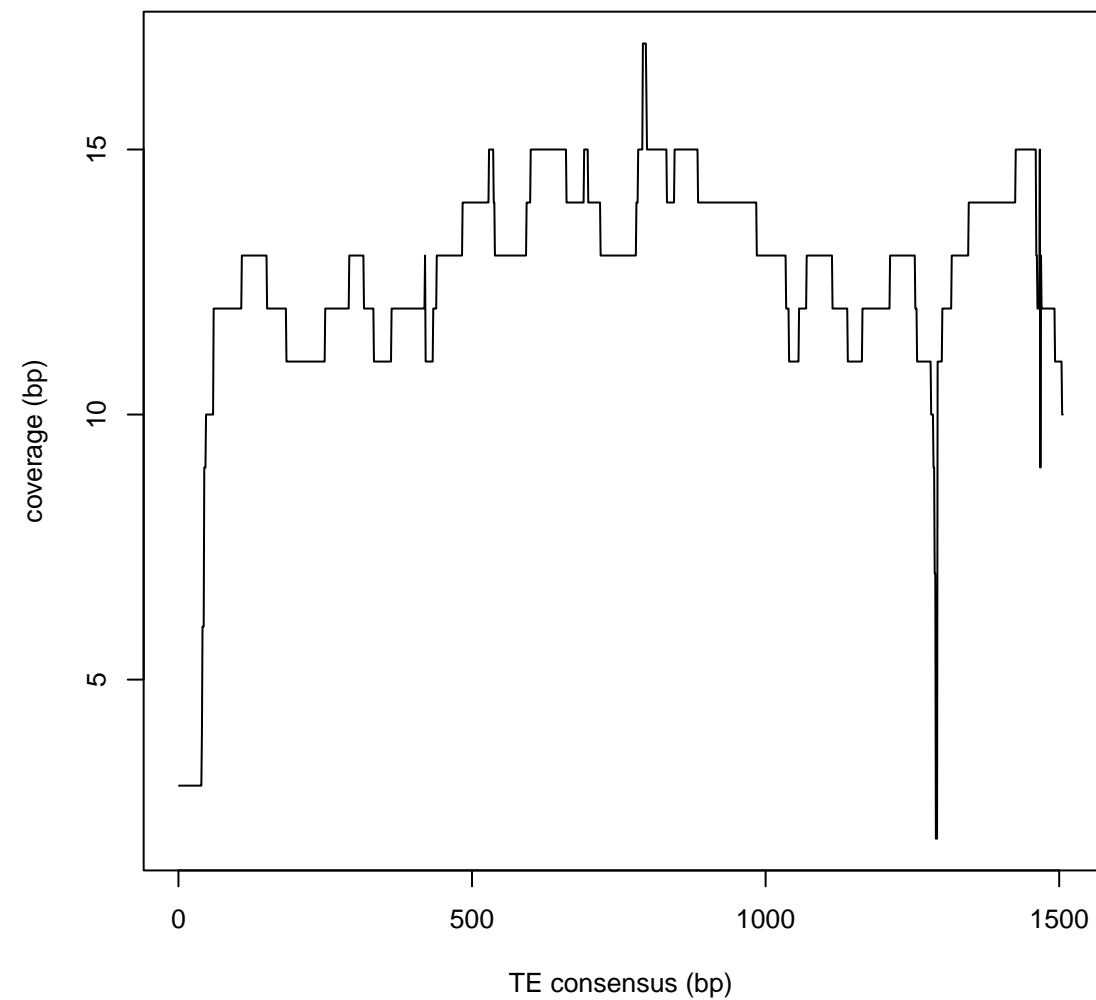


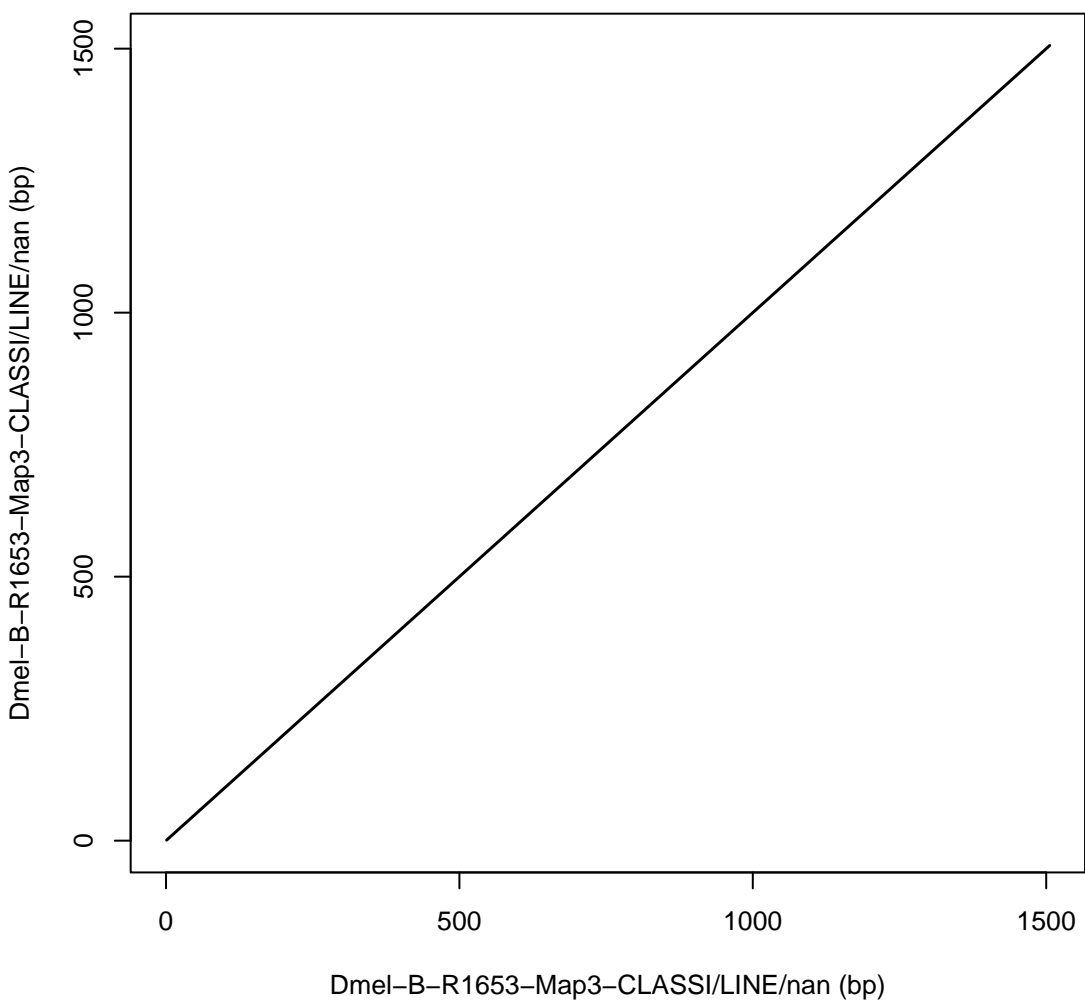
TE: Dmel-B-R1653-Map3-CLASSI/LINE/nan
consensus size: 1506bp; fragments: 55; full length: 1 (≥ 1355.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

