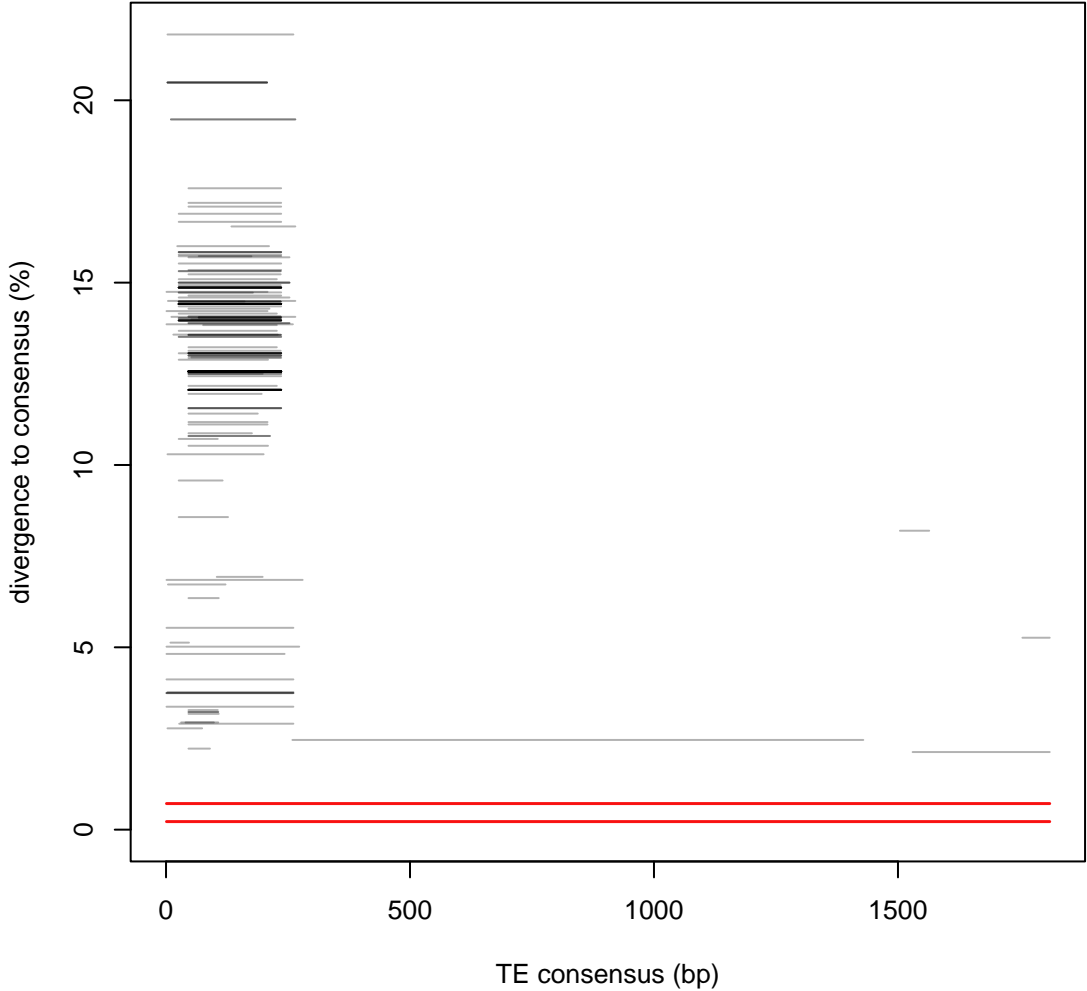
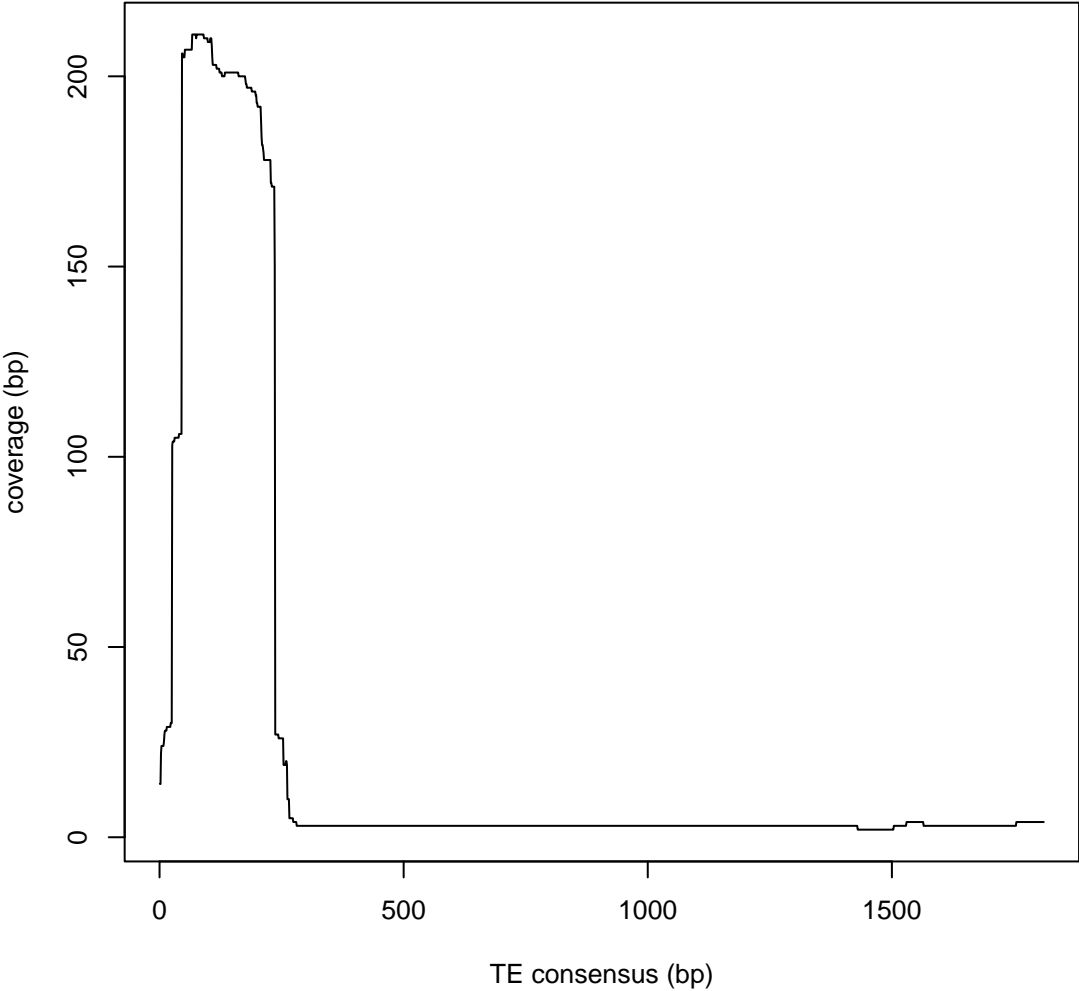


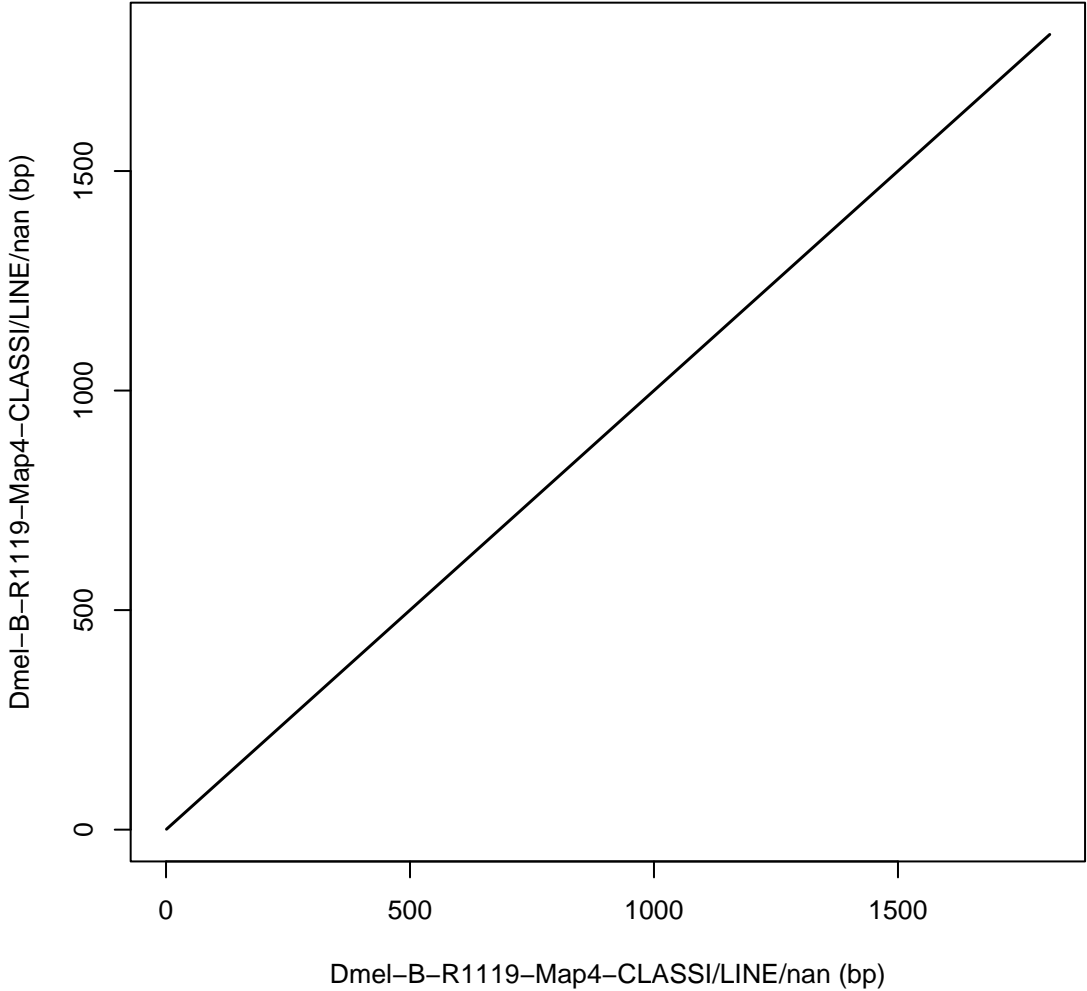
TE: Dmel-B-R1119-Map4-CLASSI/LINE/nan
consensus size: 1811bp; fragments: 219; full length: 2 (≥ 1629.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

