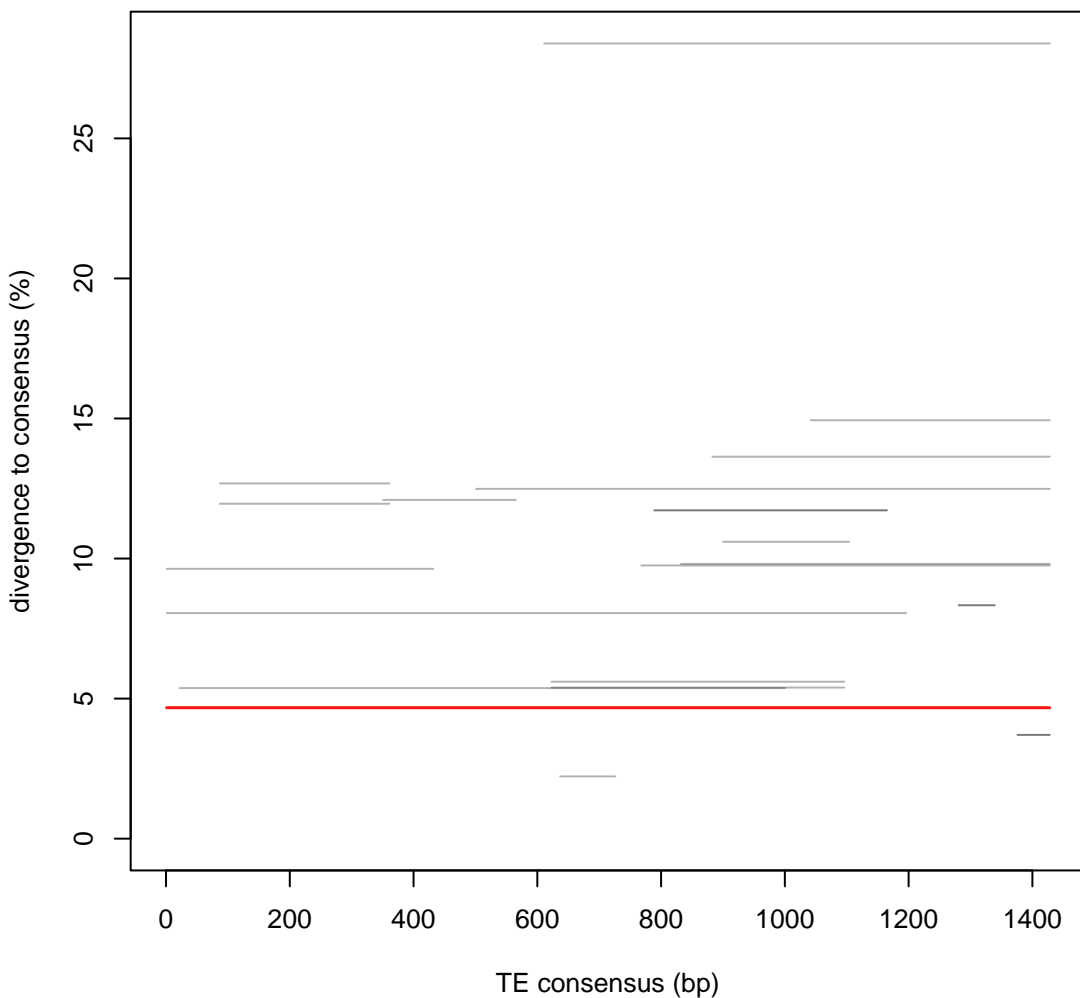
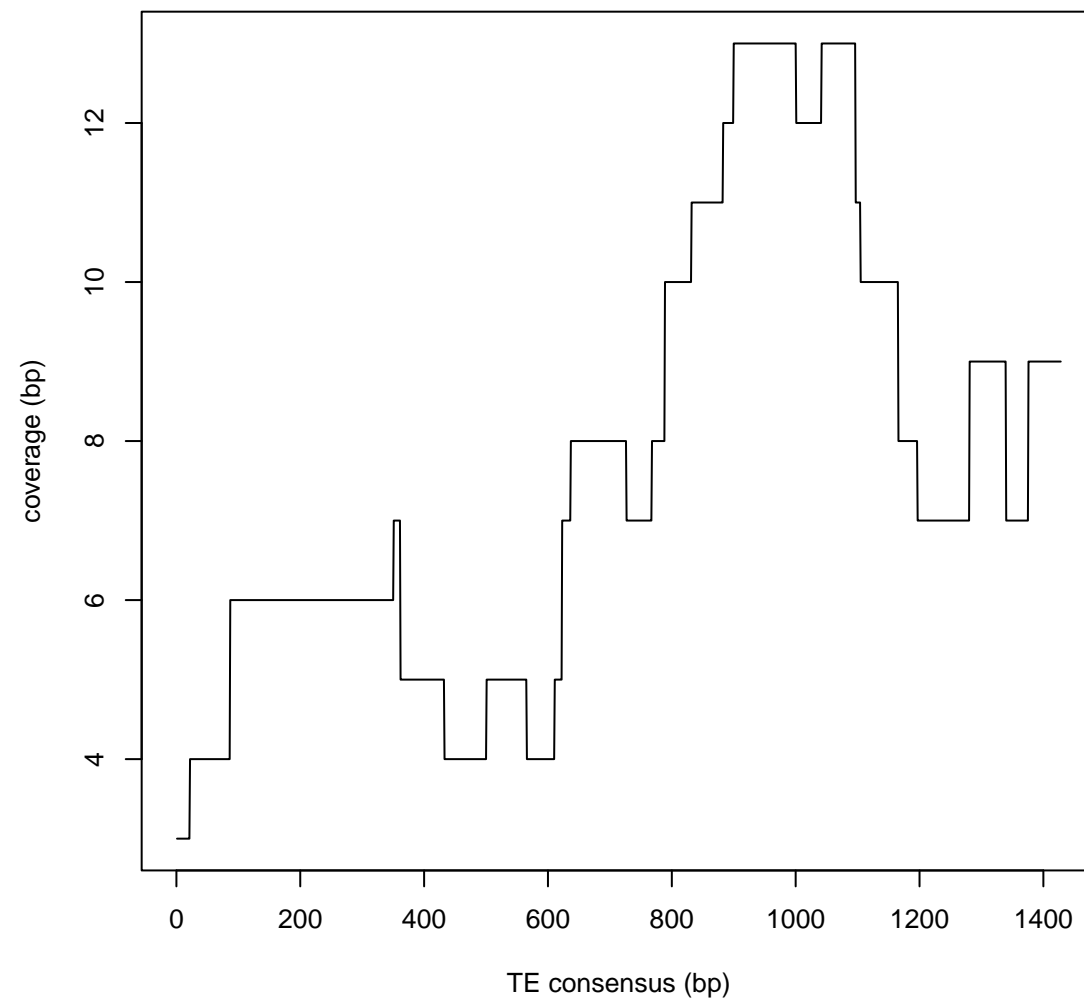


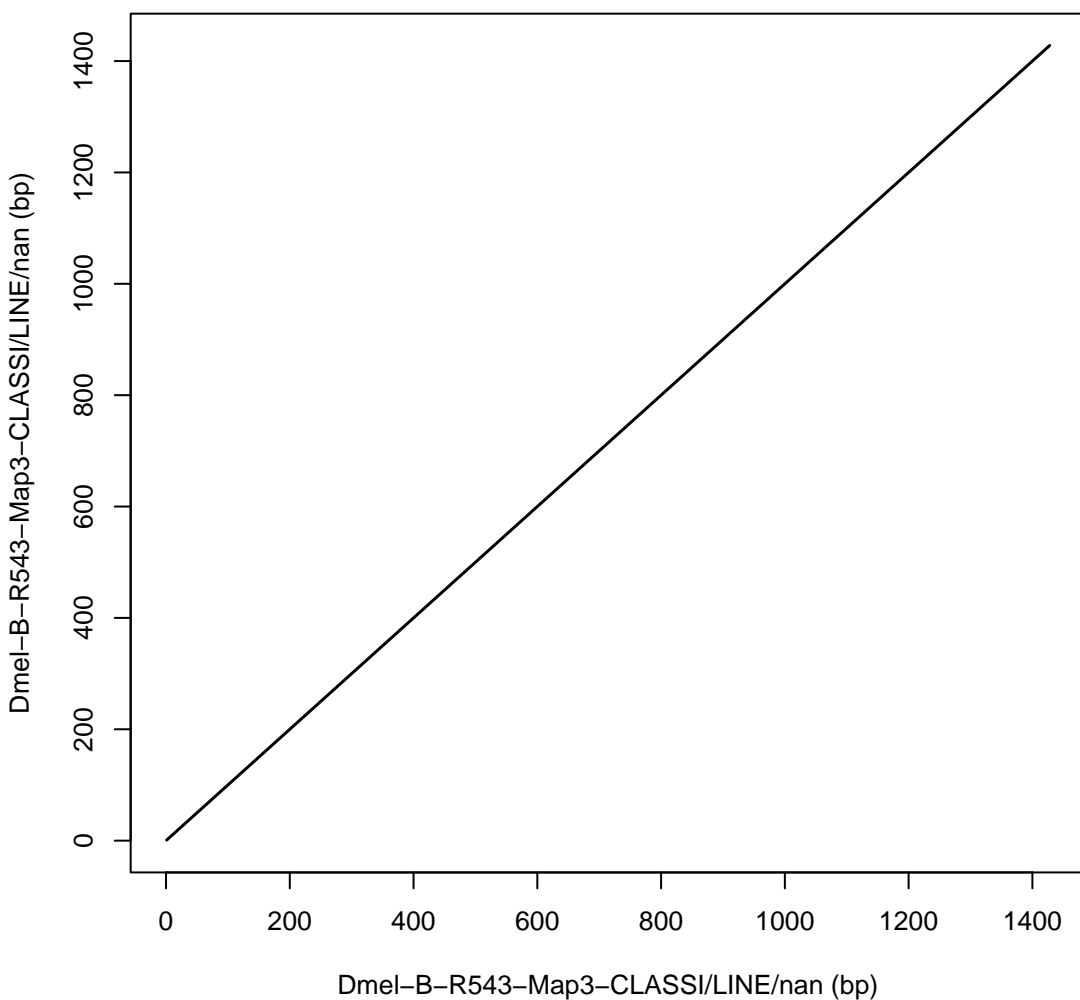
TE: Dmel-B-R543-Map3-CLASSI/LINE/nan
consensus size: 1428bp; fragments: 23; full length: 1 (≥ 1285.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

