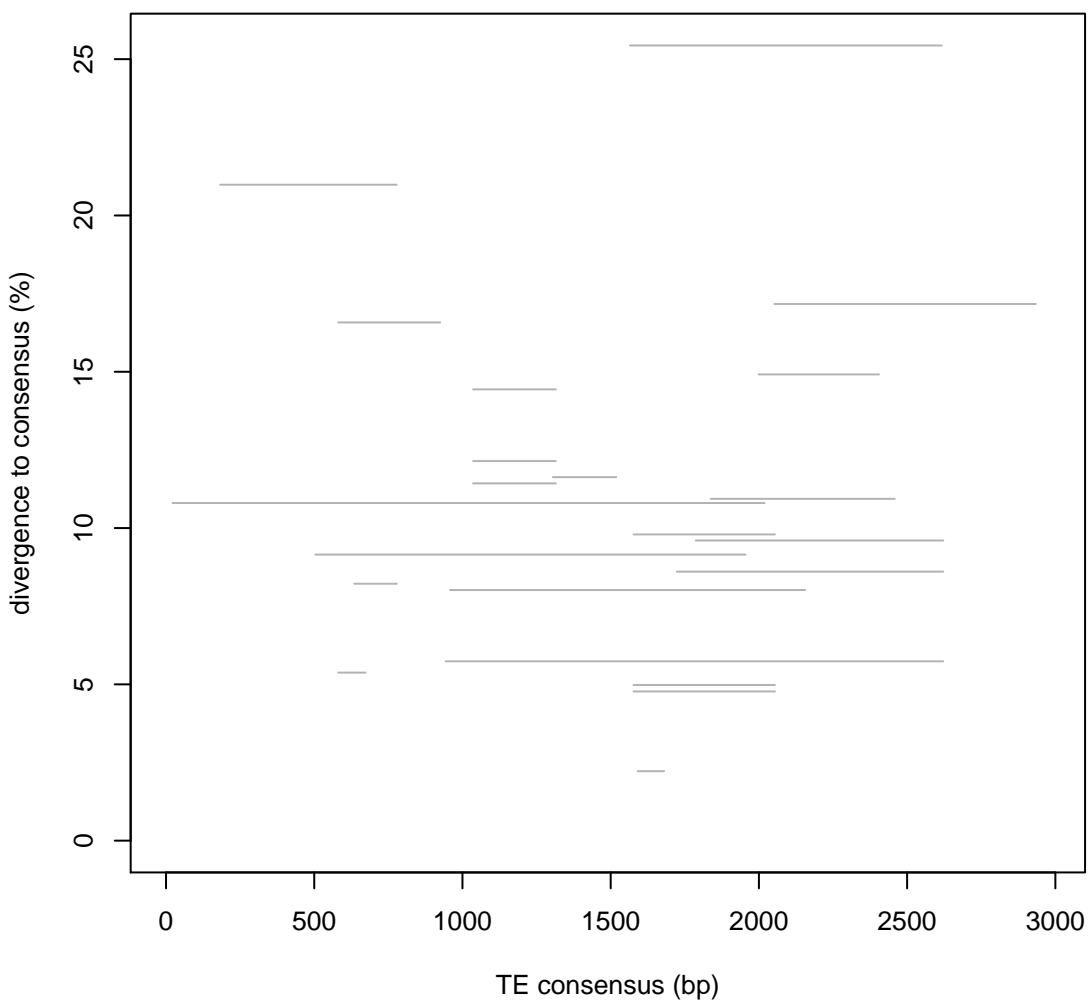
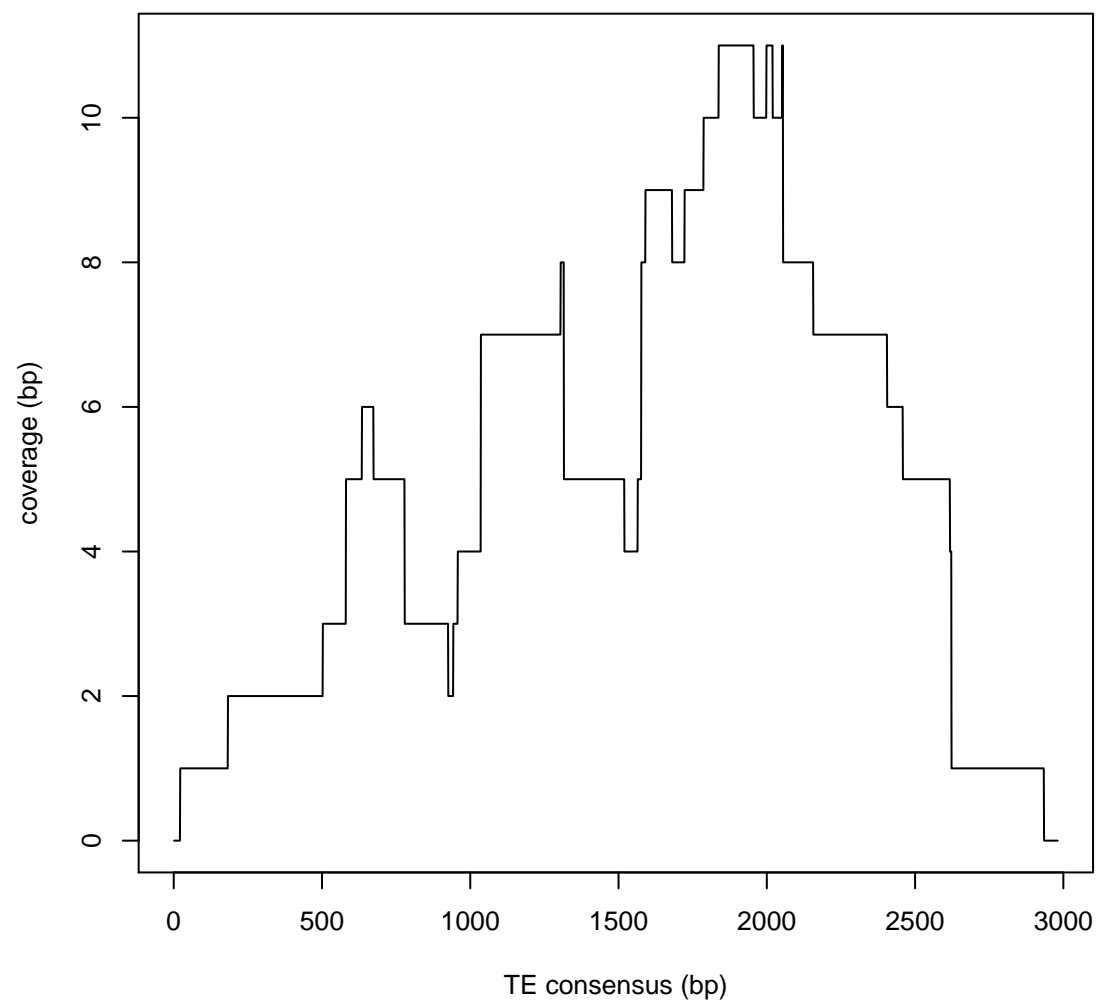


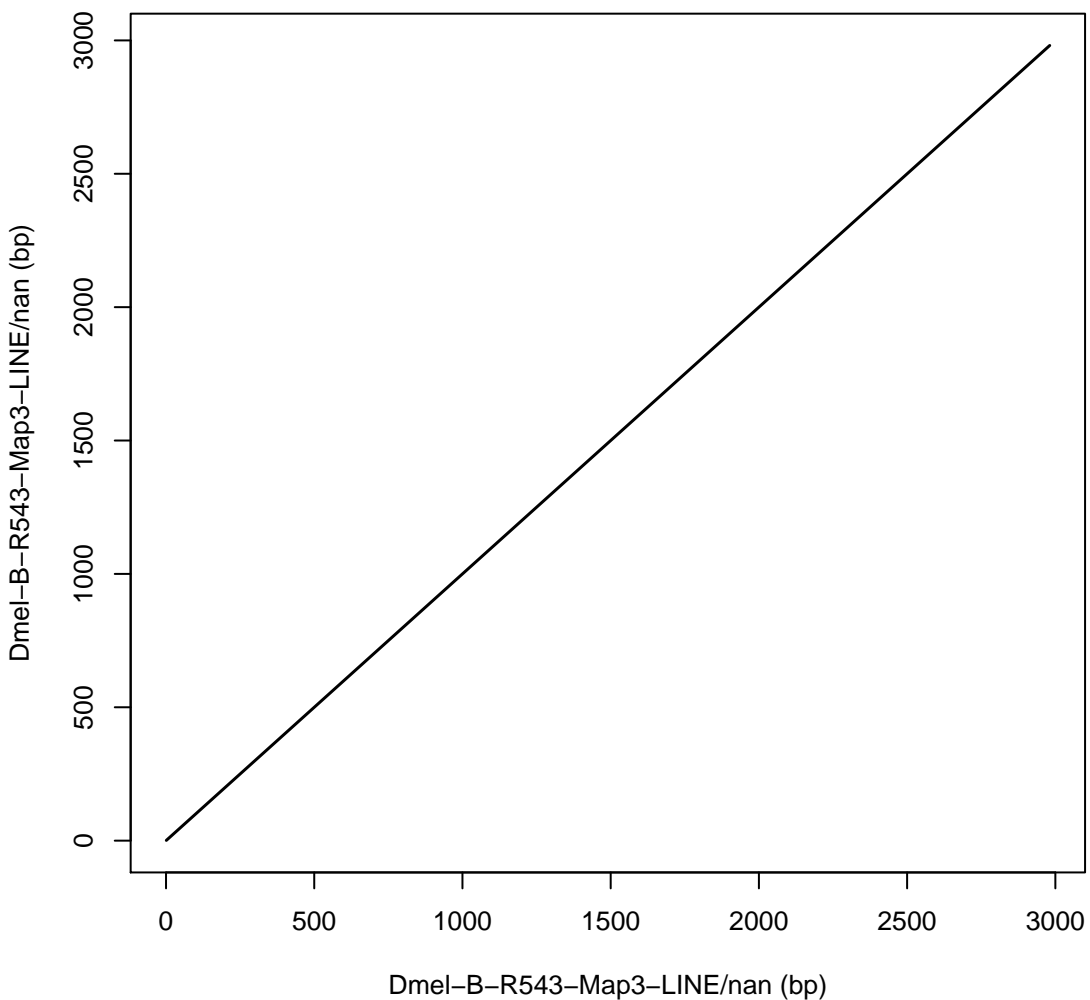
TE: Dmel-B-R543-Map3-LINE/nan
consensus size: 2981bp; fragments: 22; full length: 0 (≥ 2682.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

