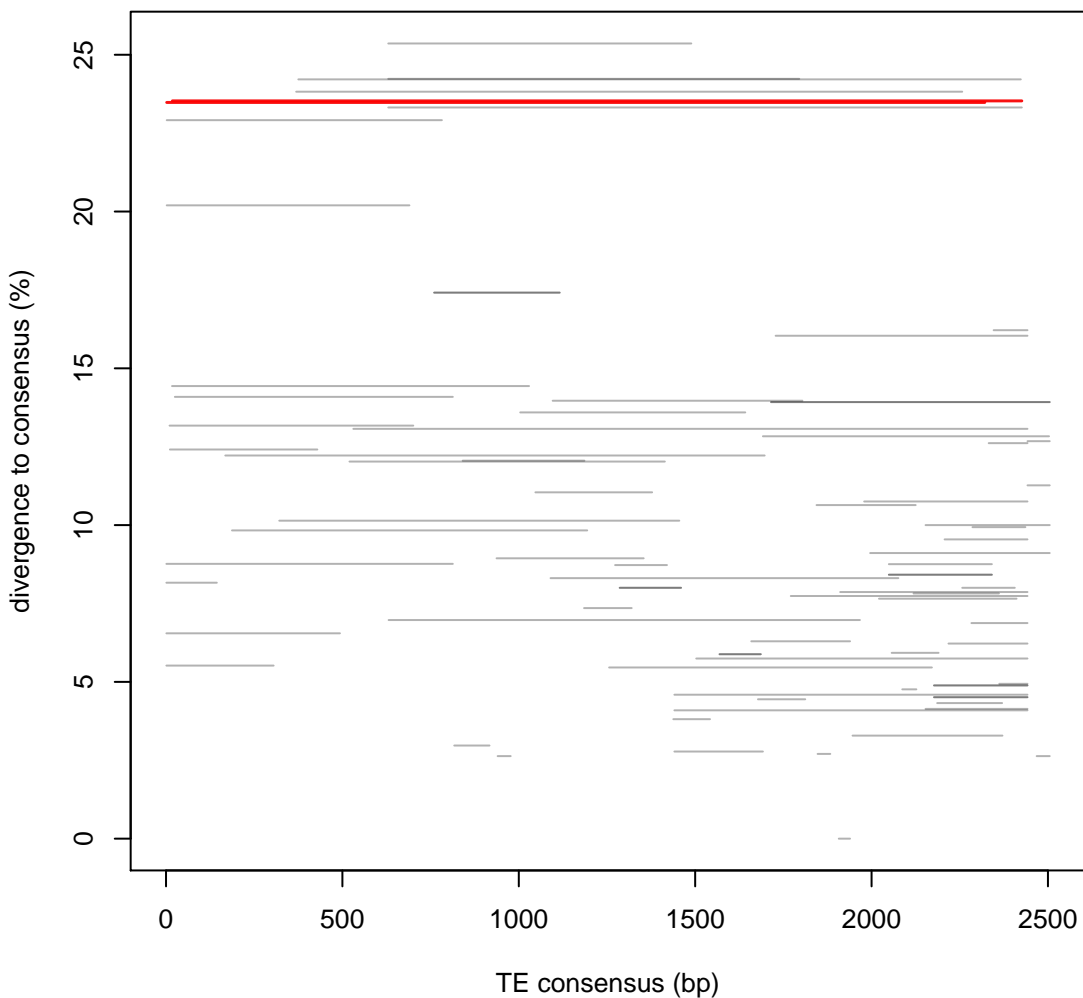
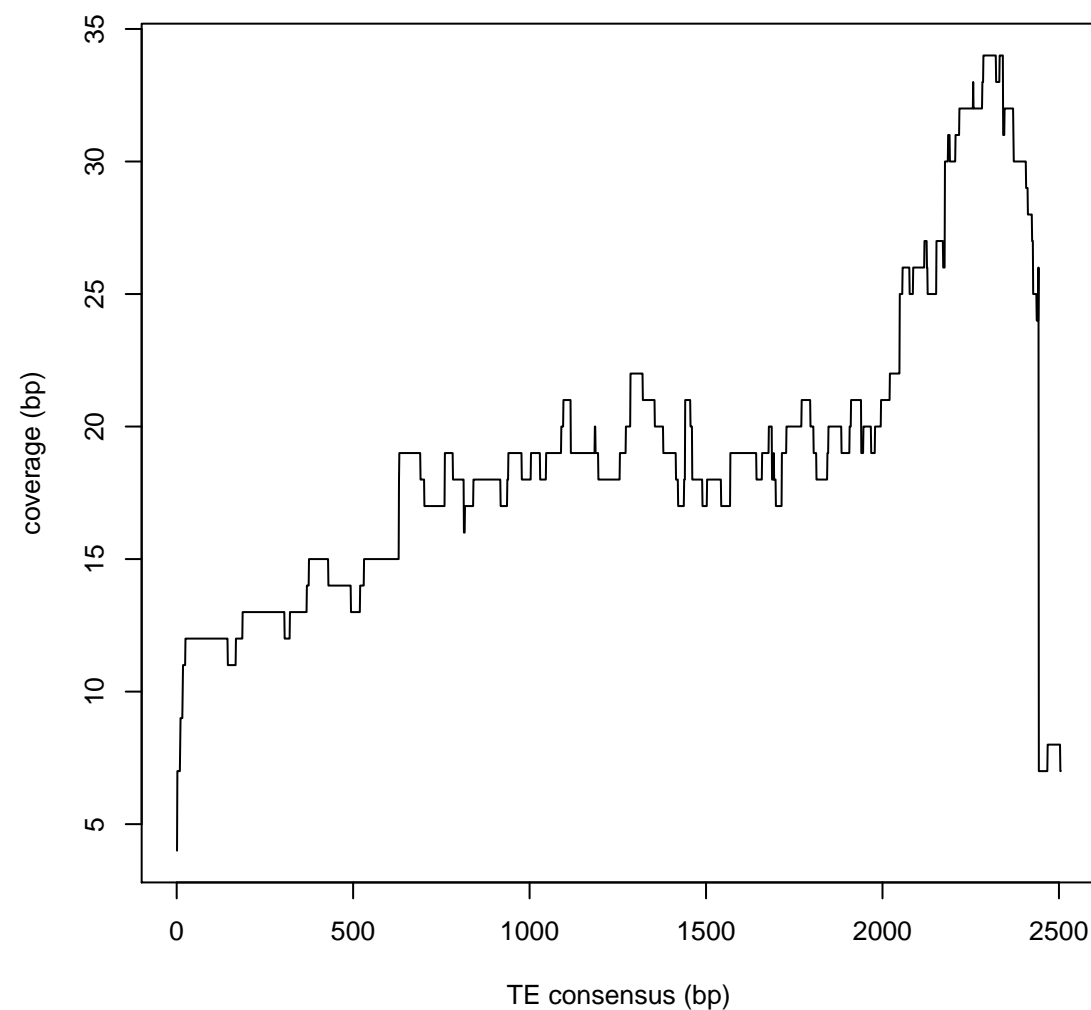


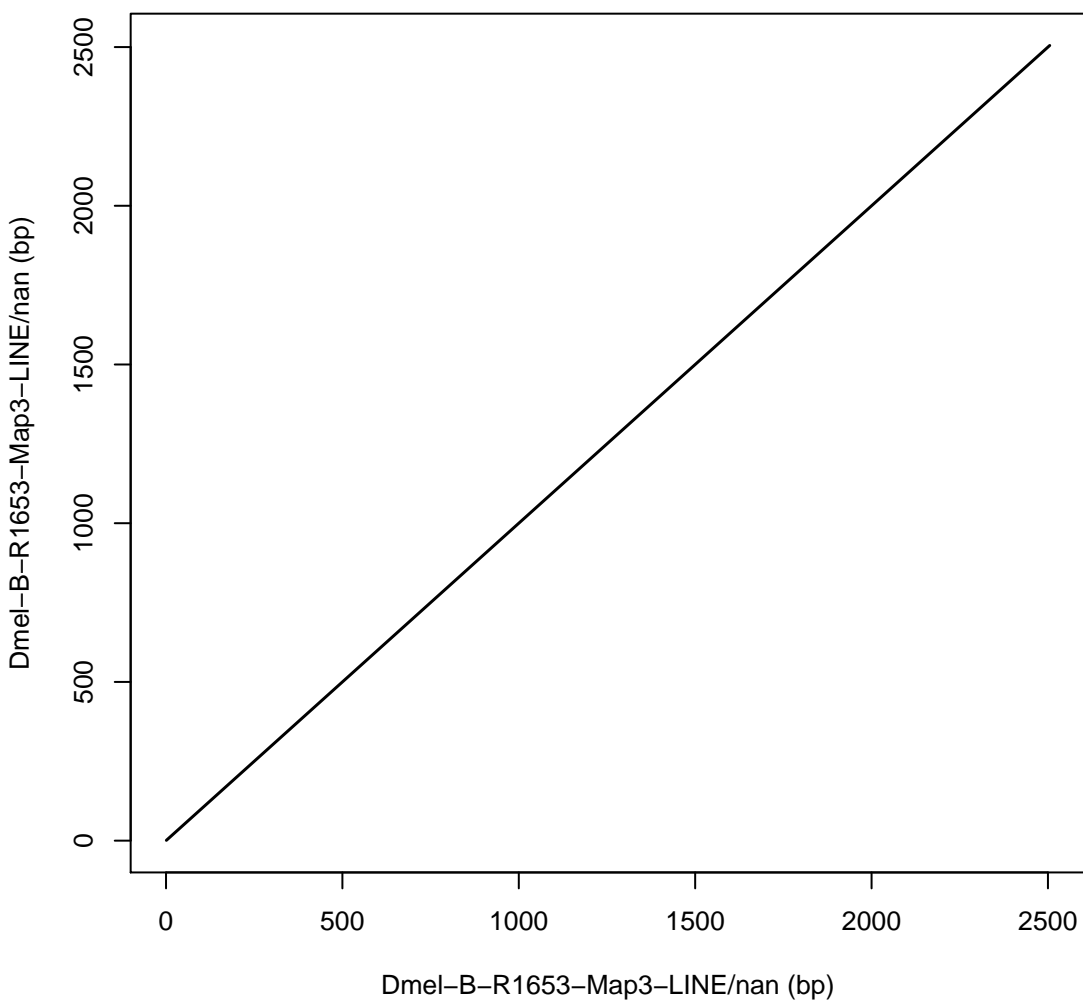
TE: Dmel-B-R1653-Map3-LINE/nan
consensus size: 2505bp; fragments: 84; full length: 2 (≥ 2254.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

