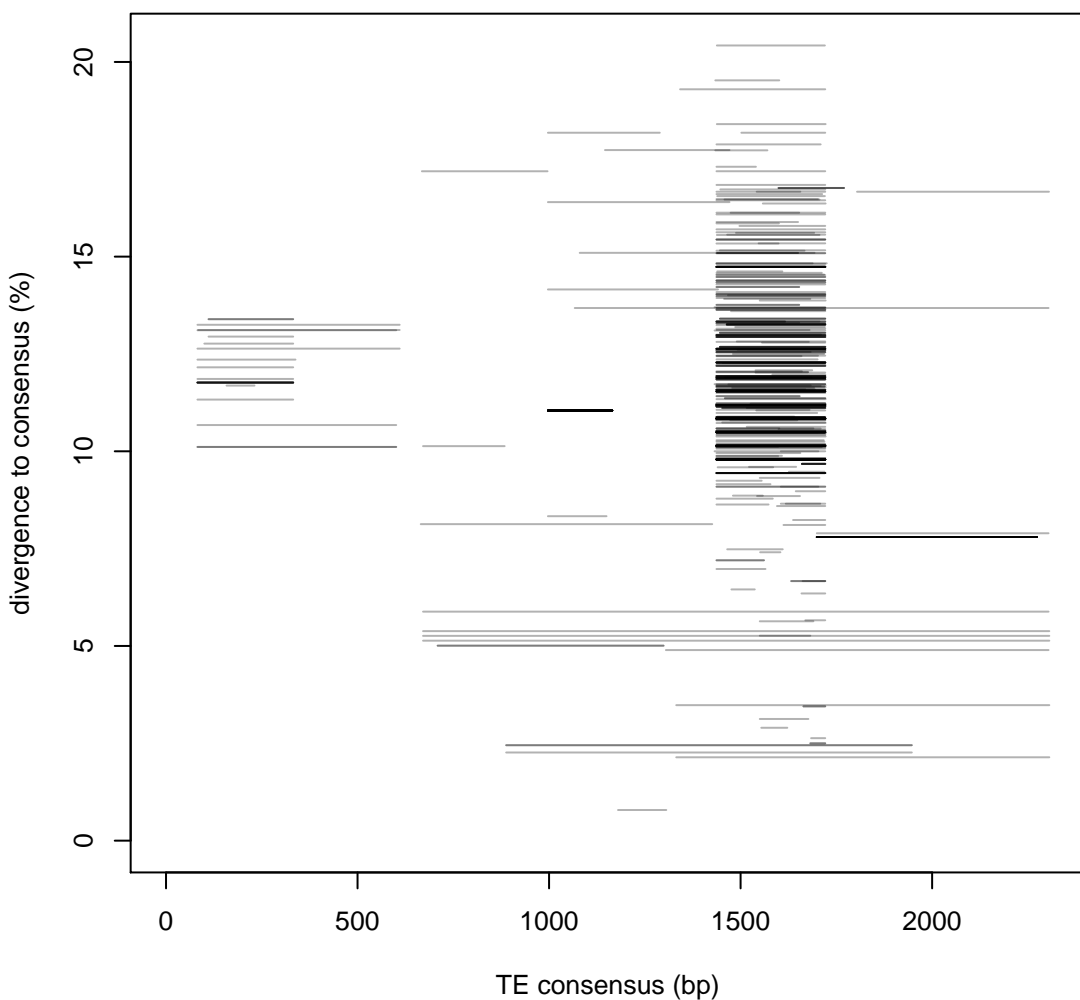
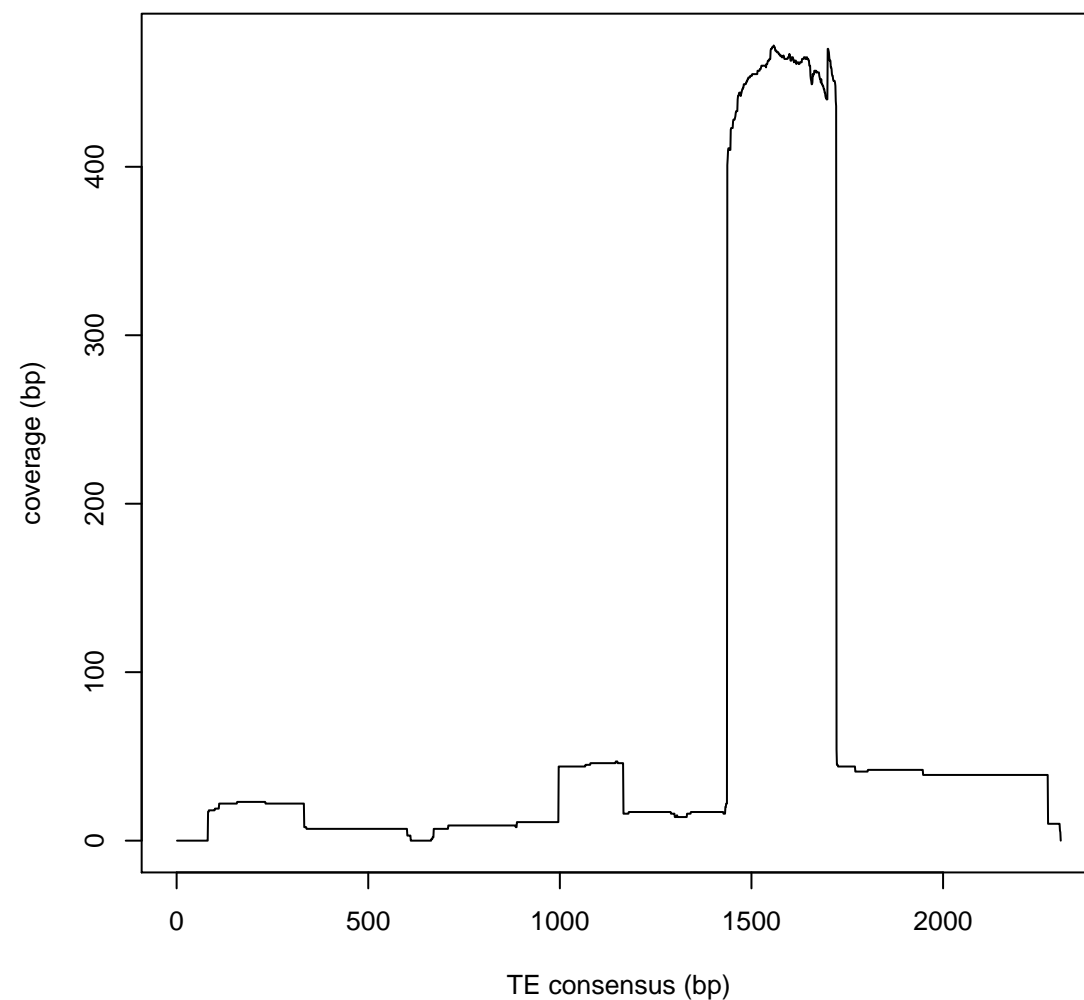


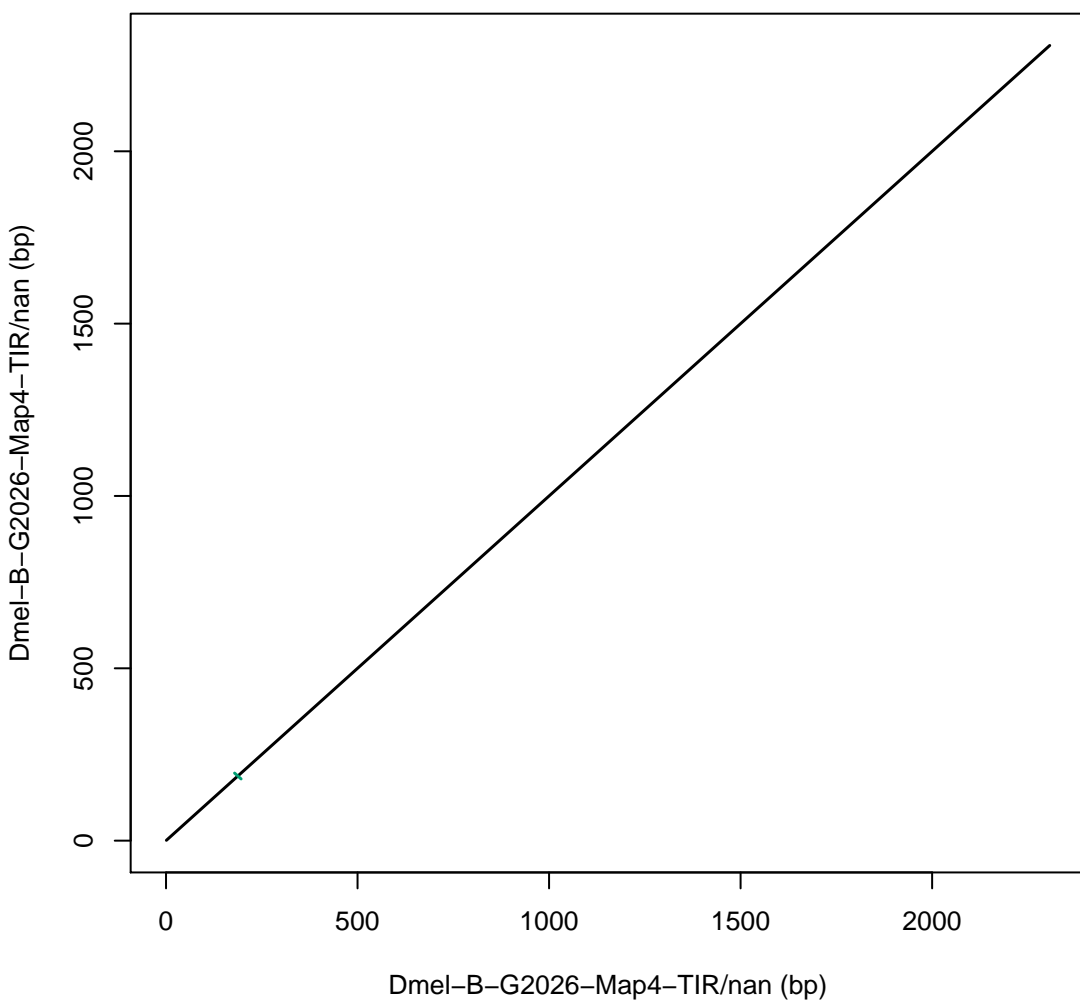
TE: Dmel-B-G2026-Map4-TIR/nan
consensus size: 2307bp; fragments: 600; full length: 0 (>=2076.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

