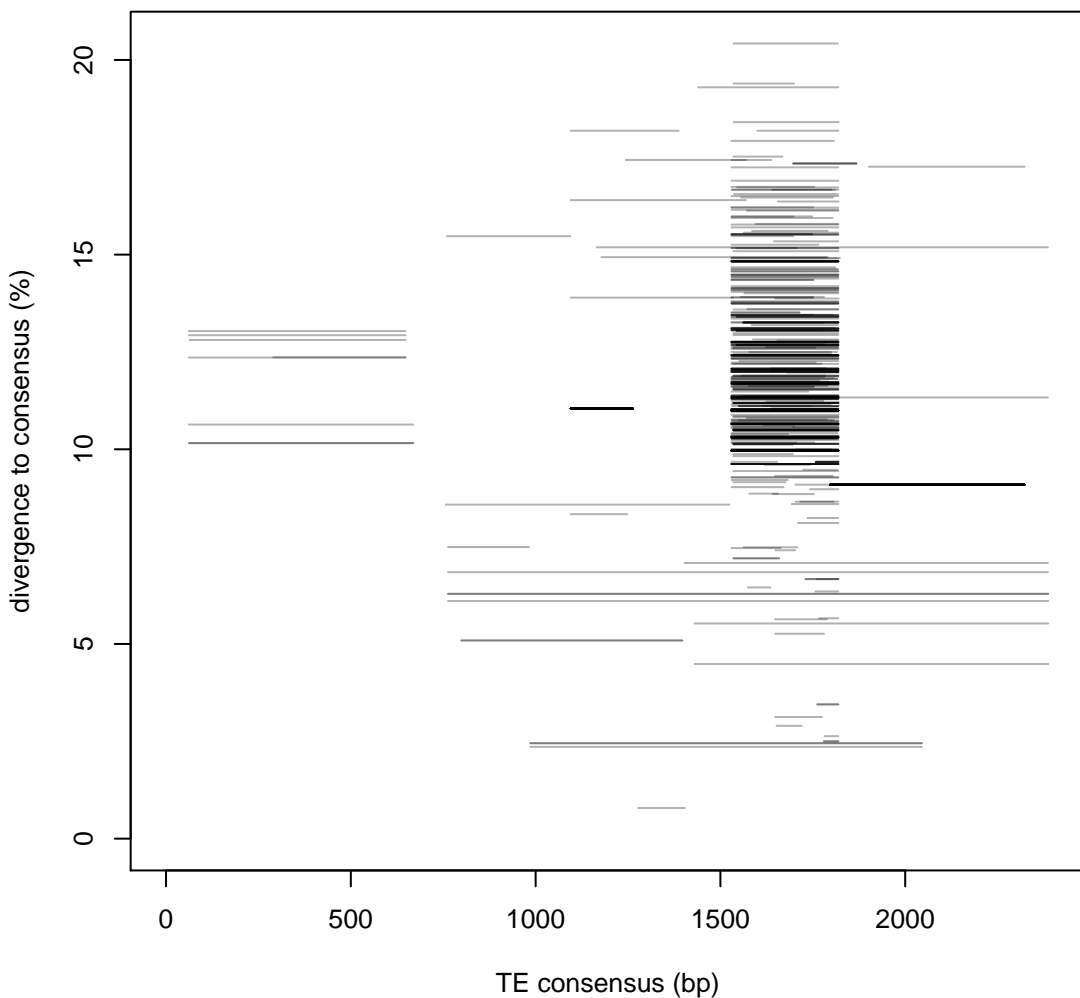
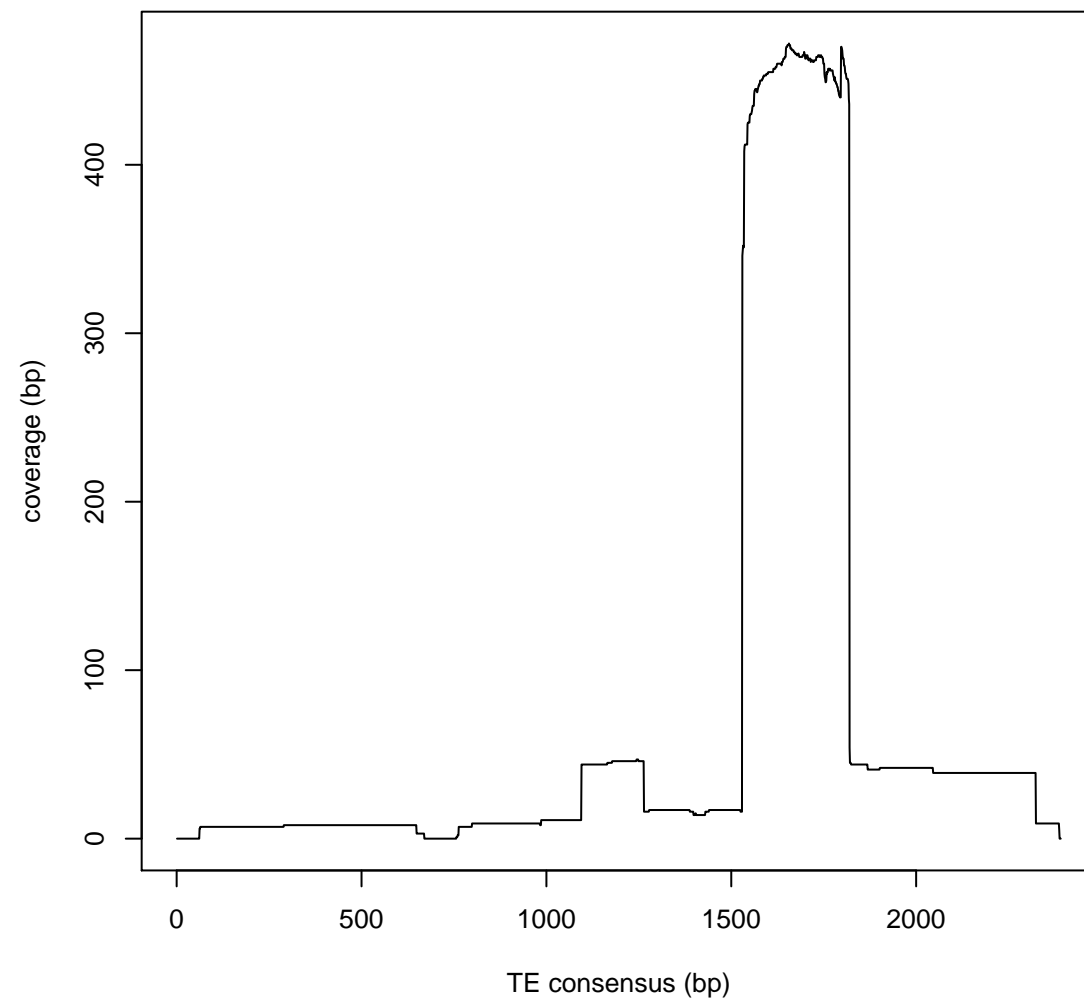


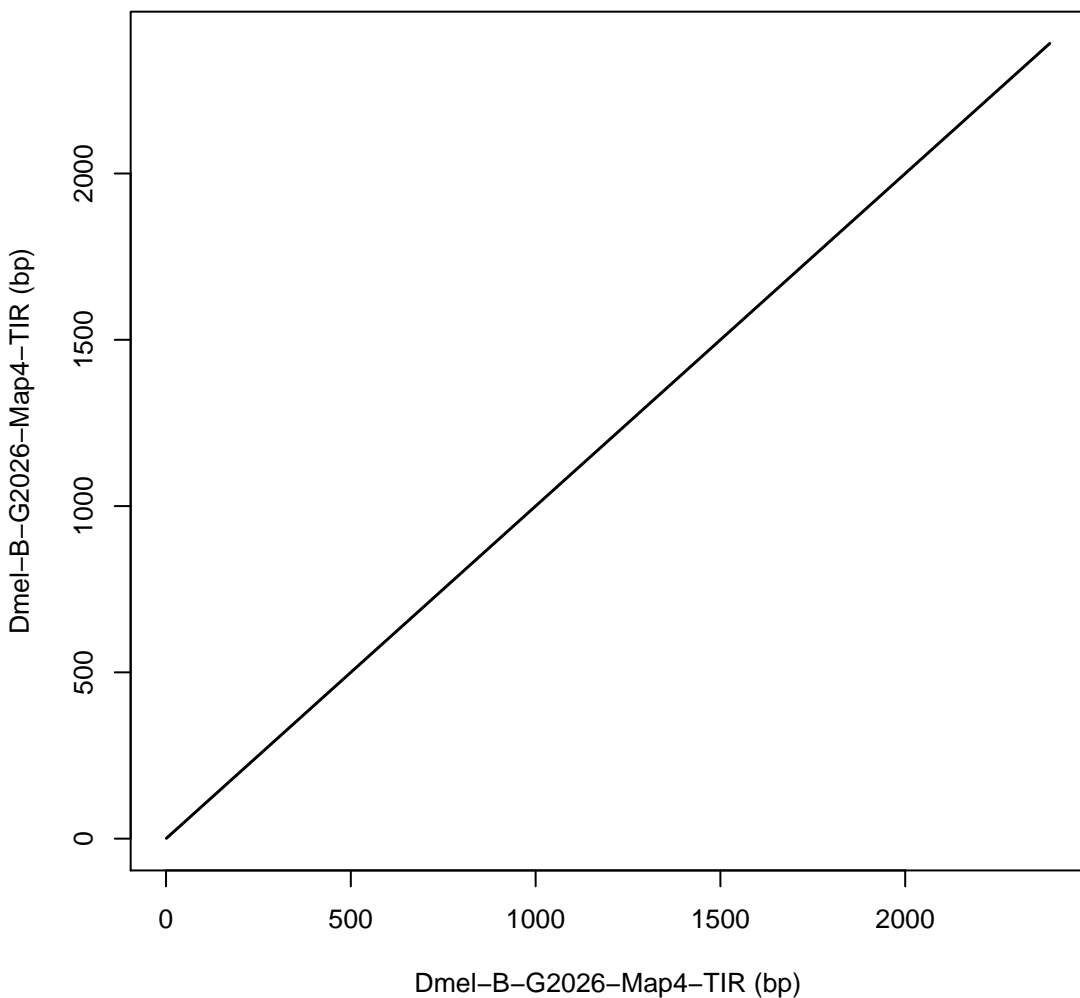
TE: Dmel-B-G2026-Map4-TIR
consensus size: 2391bp; fragments: 585; full length: 0 (>=2151.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

