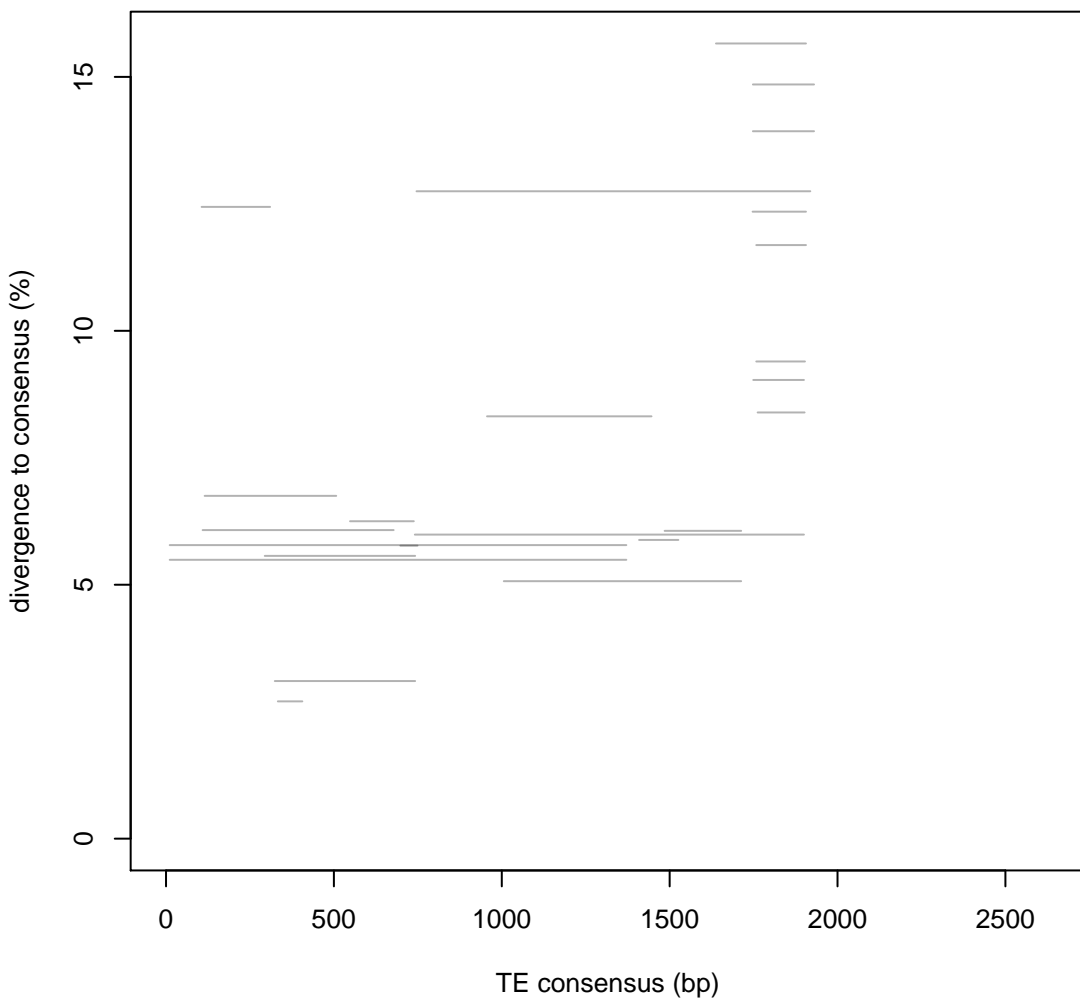
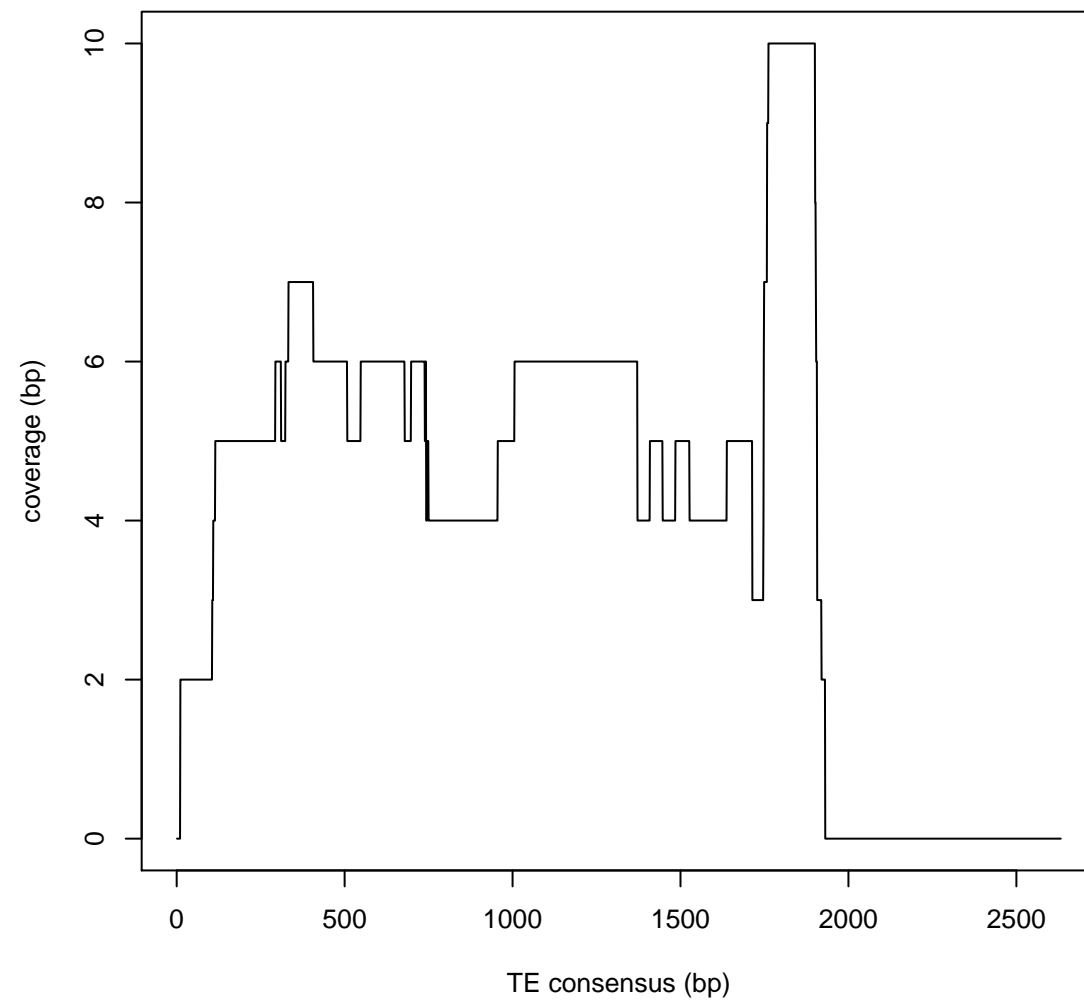


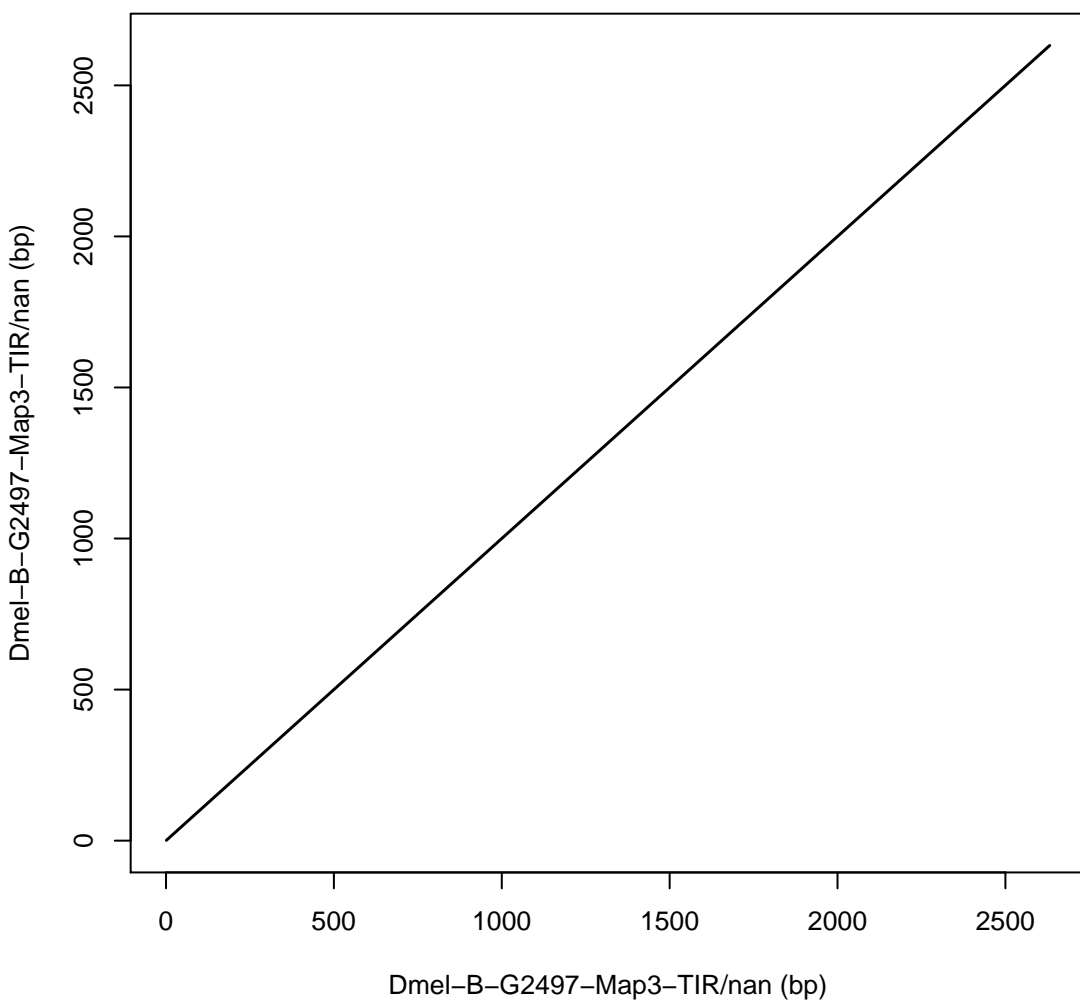
TE: Dmel-B-G2497-Map3-TIR/nan
consensus size: 2632bp; fragments: 24; full length: 0 (>=2368.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

