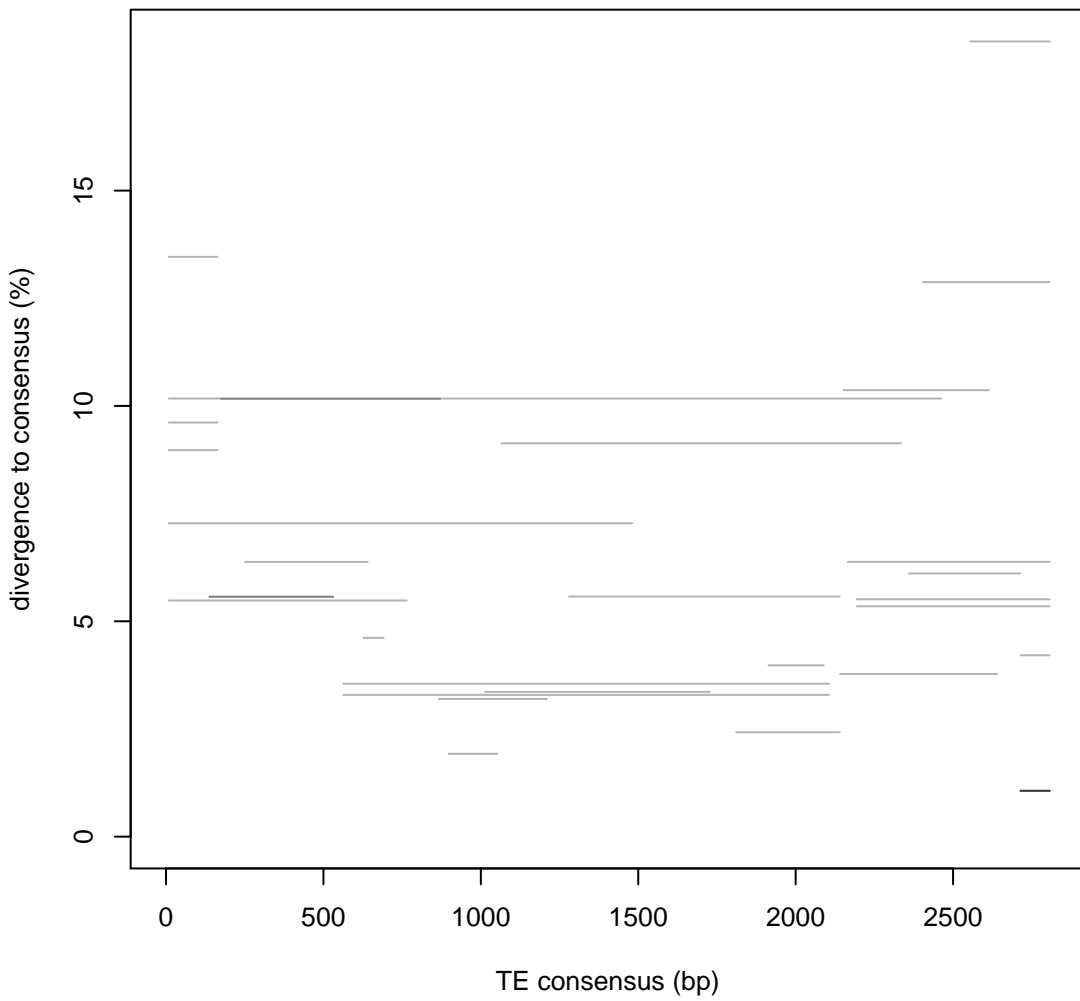
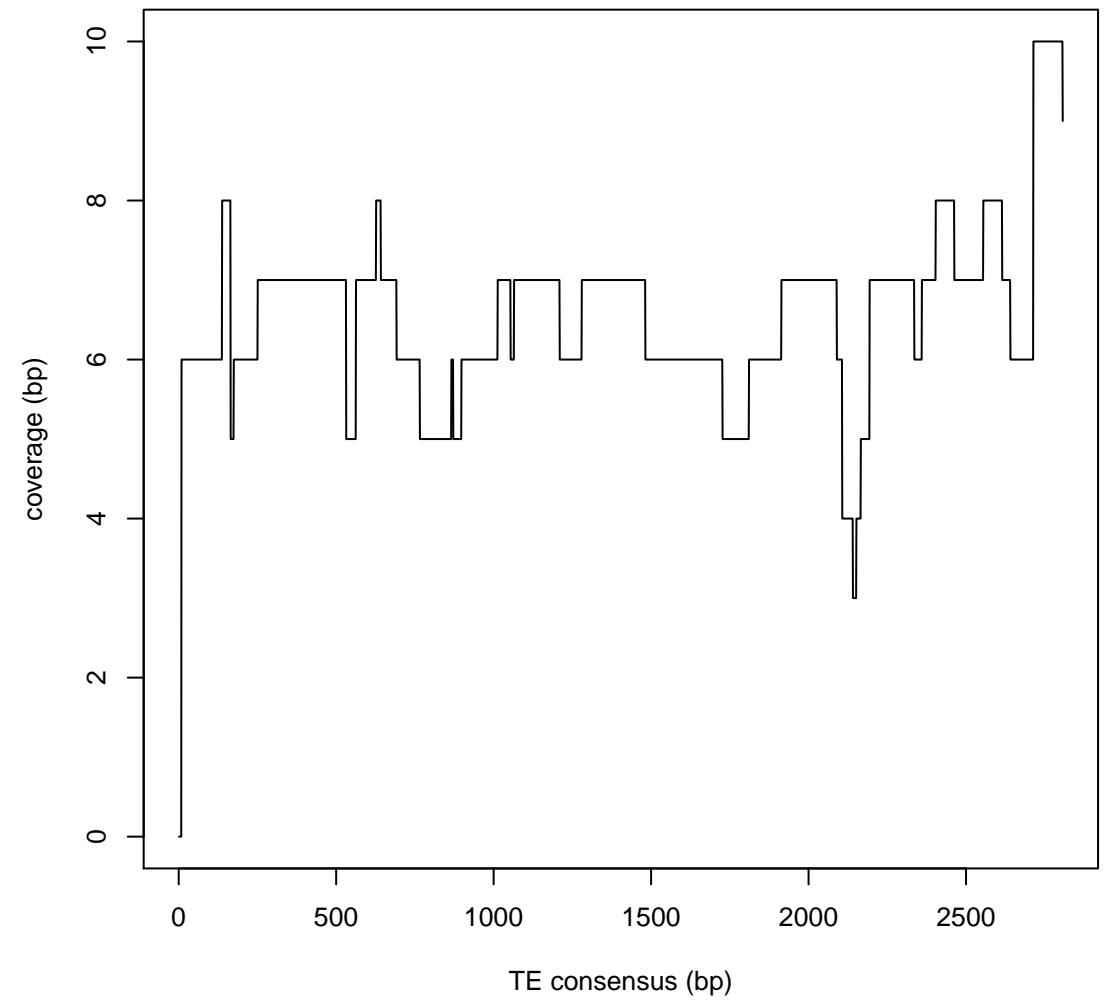


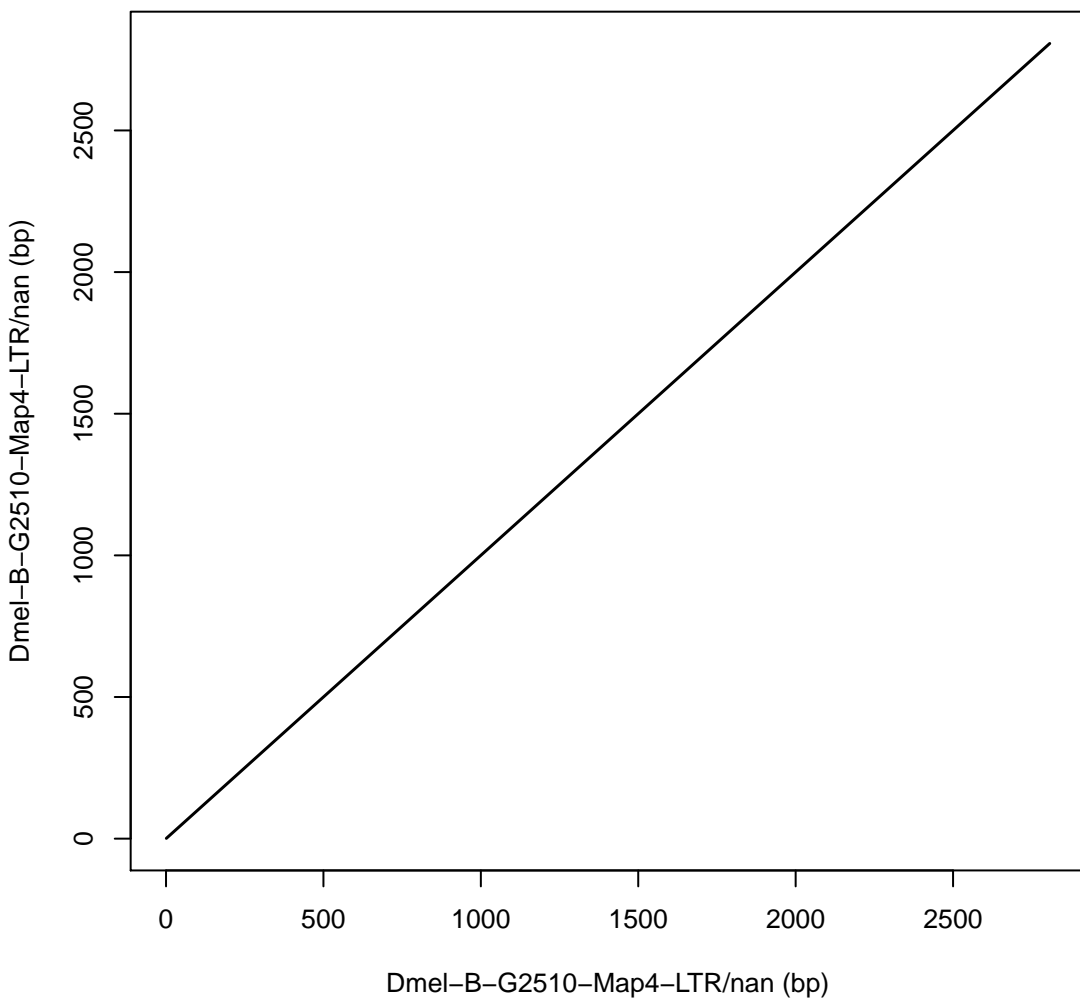
TE: Dmel-B-G2510-Map4-LTR/nan
consensus size: 2807bp; fragments: 33; full length: 0 (>=2526.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

