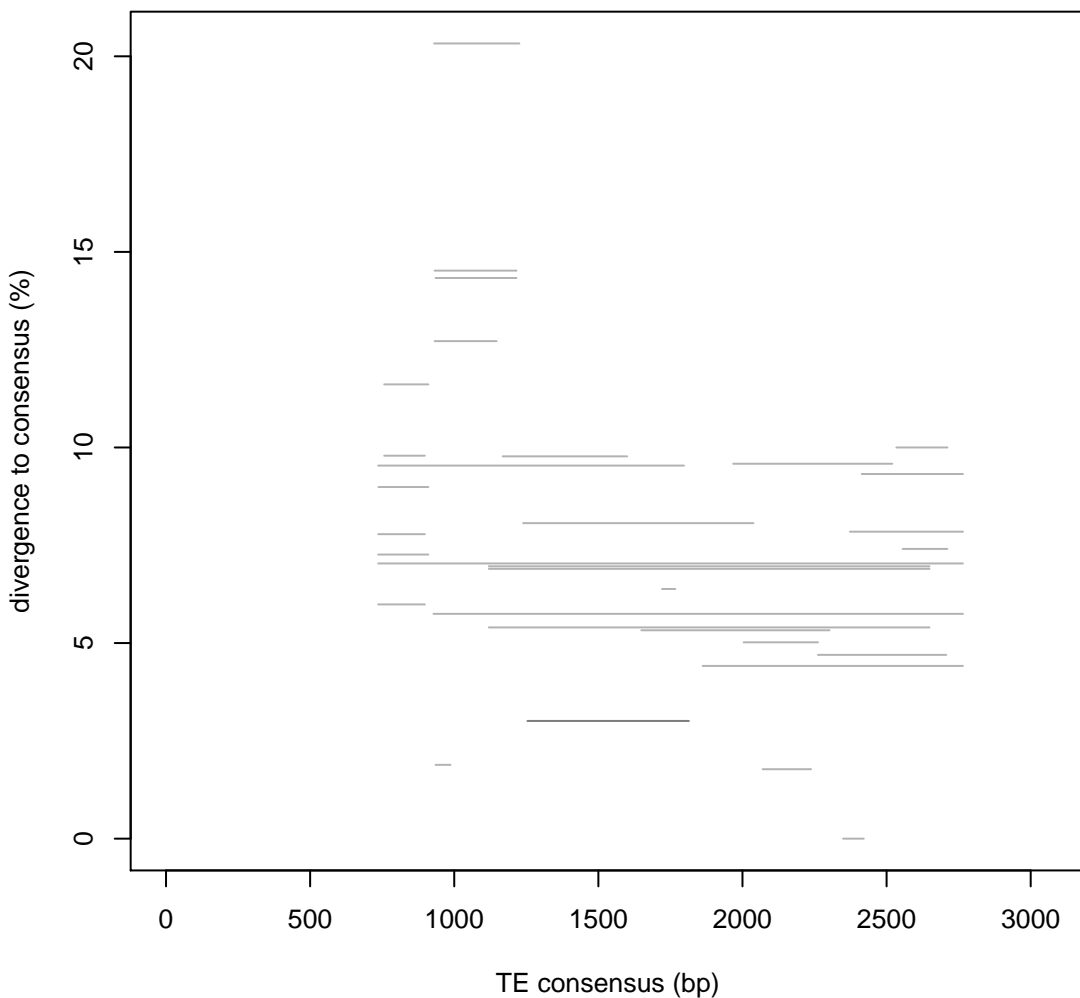
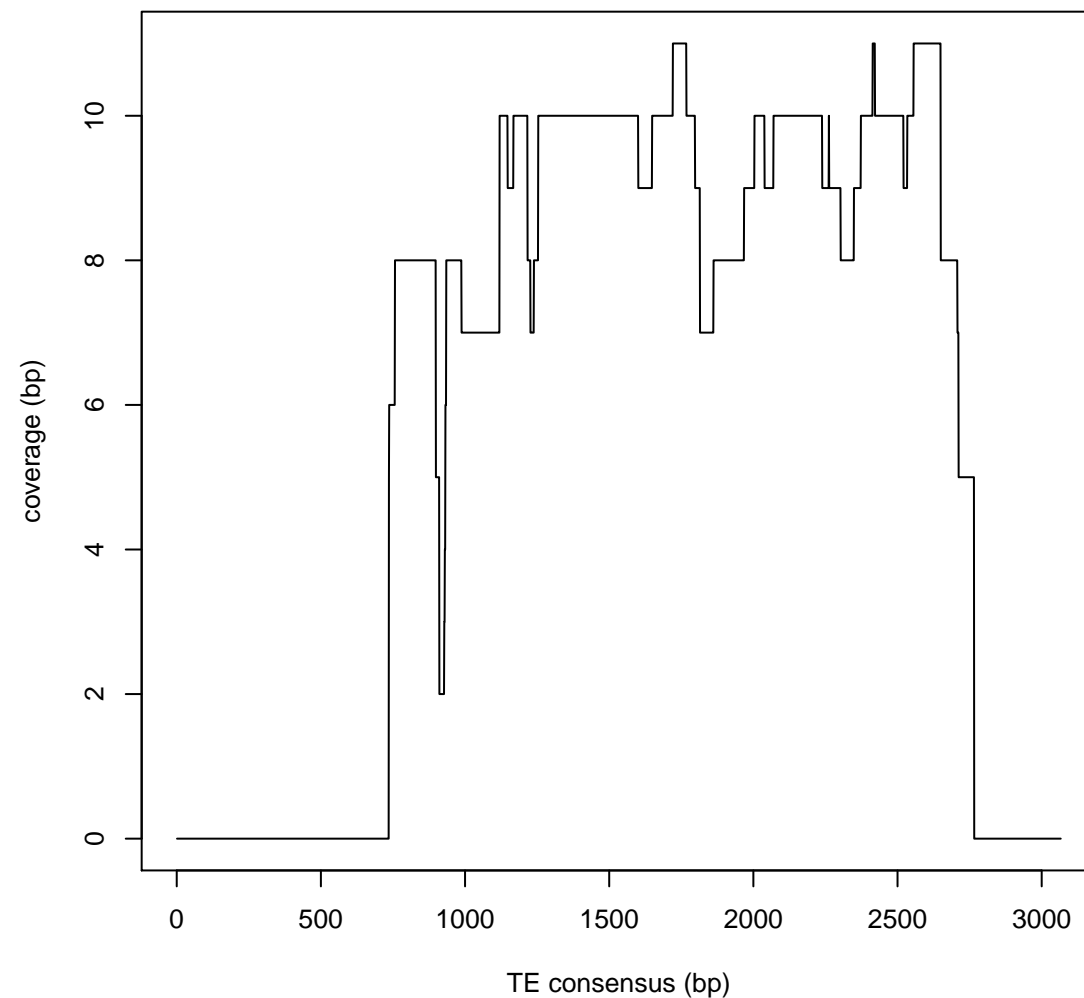


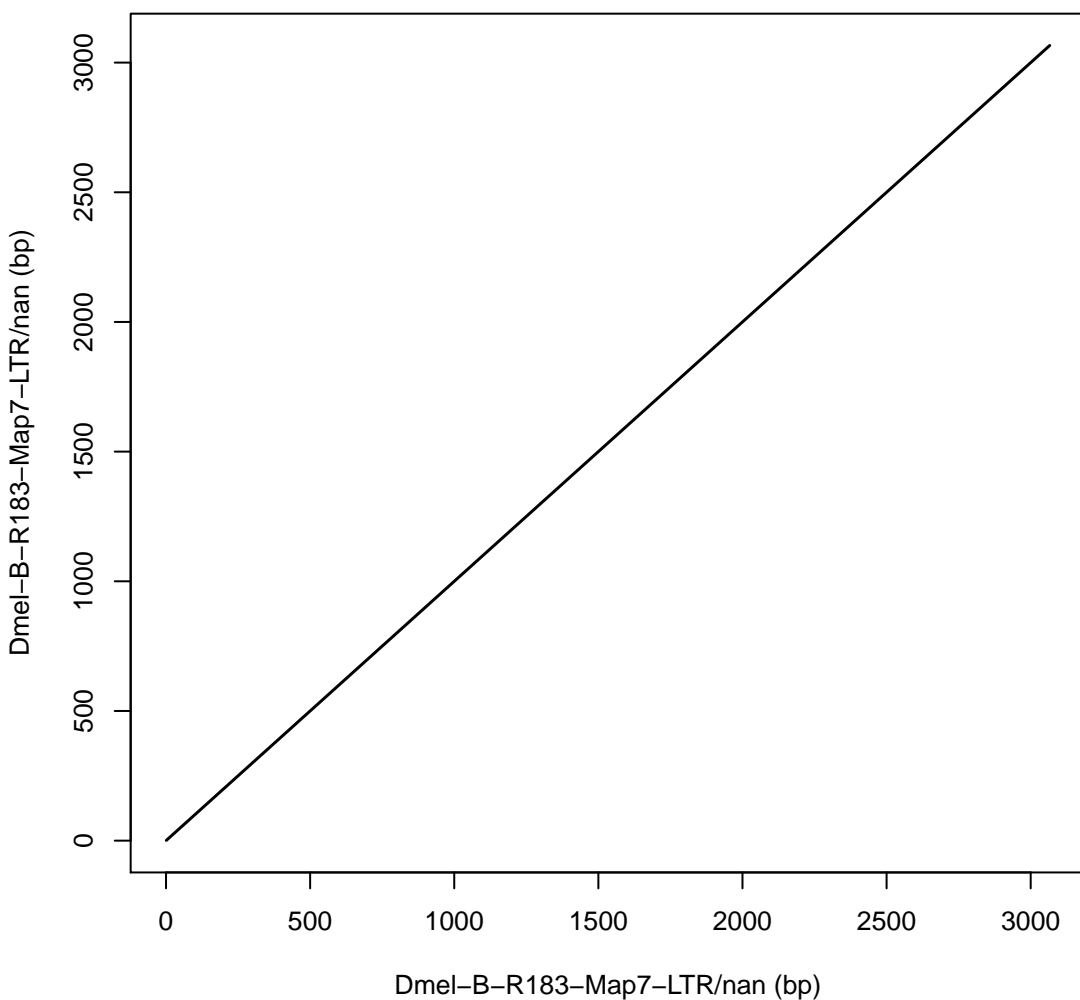
TE: Dmel-B-R183-Map7-LTR/nan
consensus size: 3066bp; fragments: 33; full length: 0 (>=2759.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

