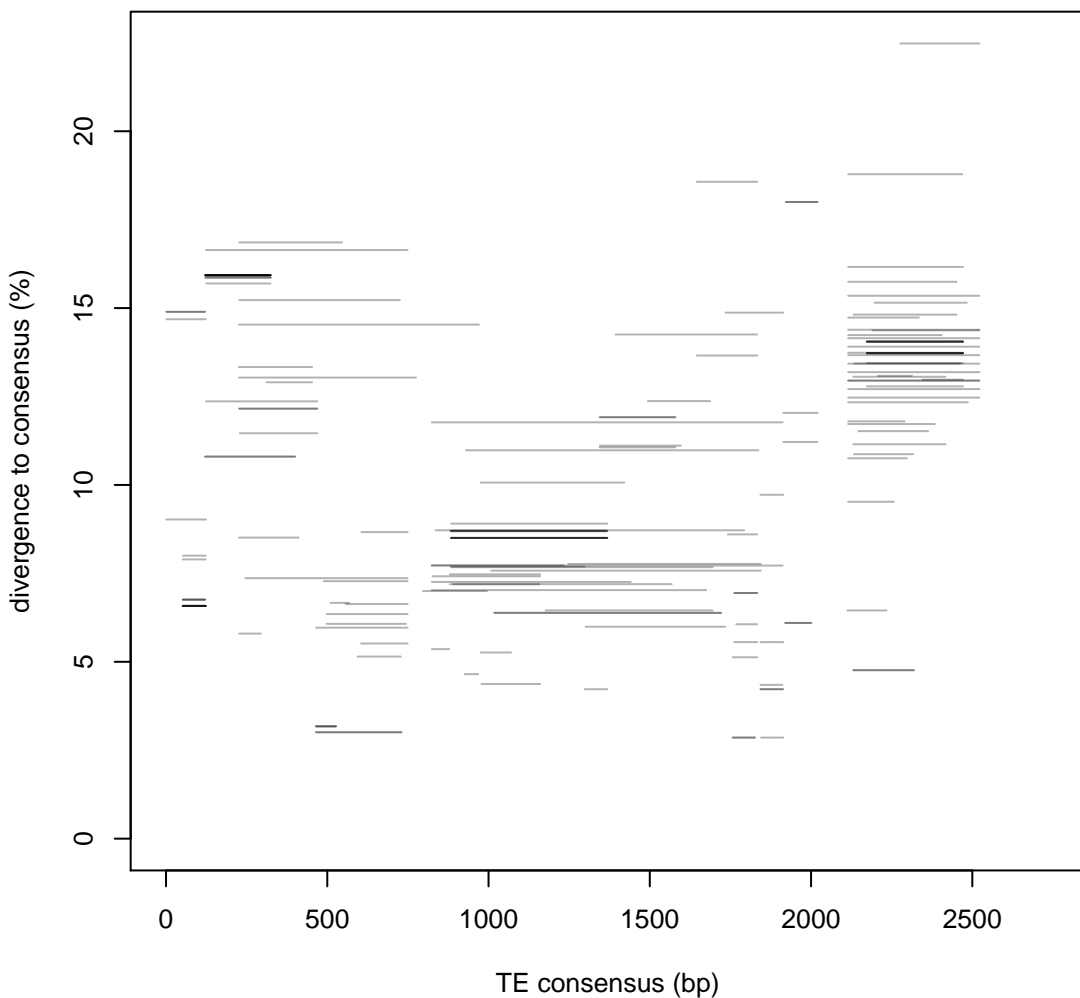
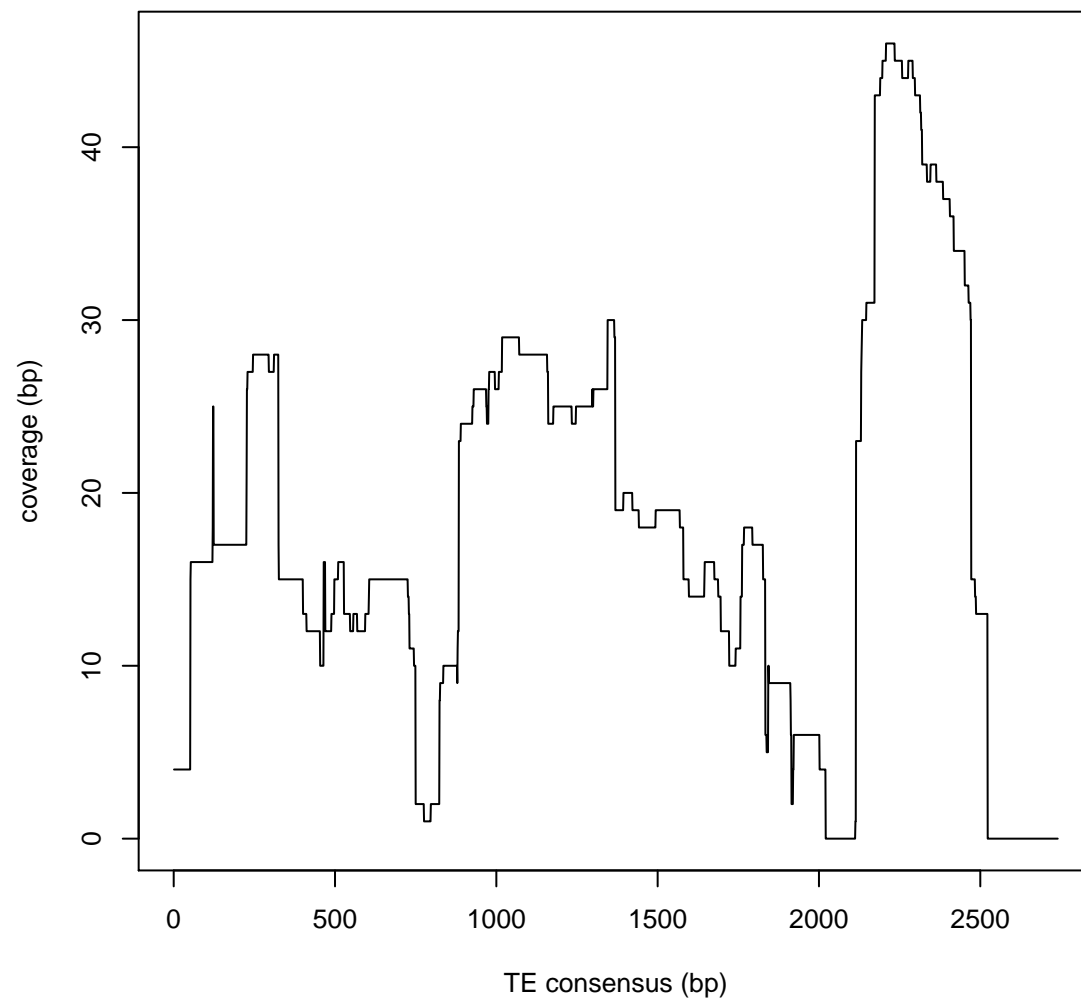


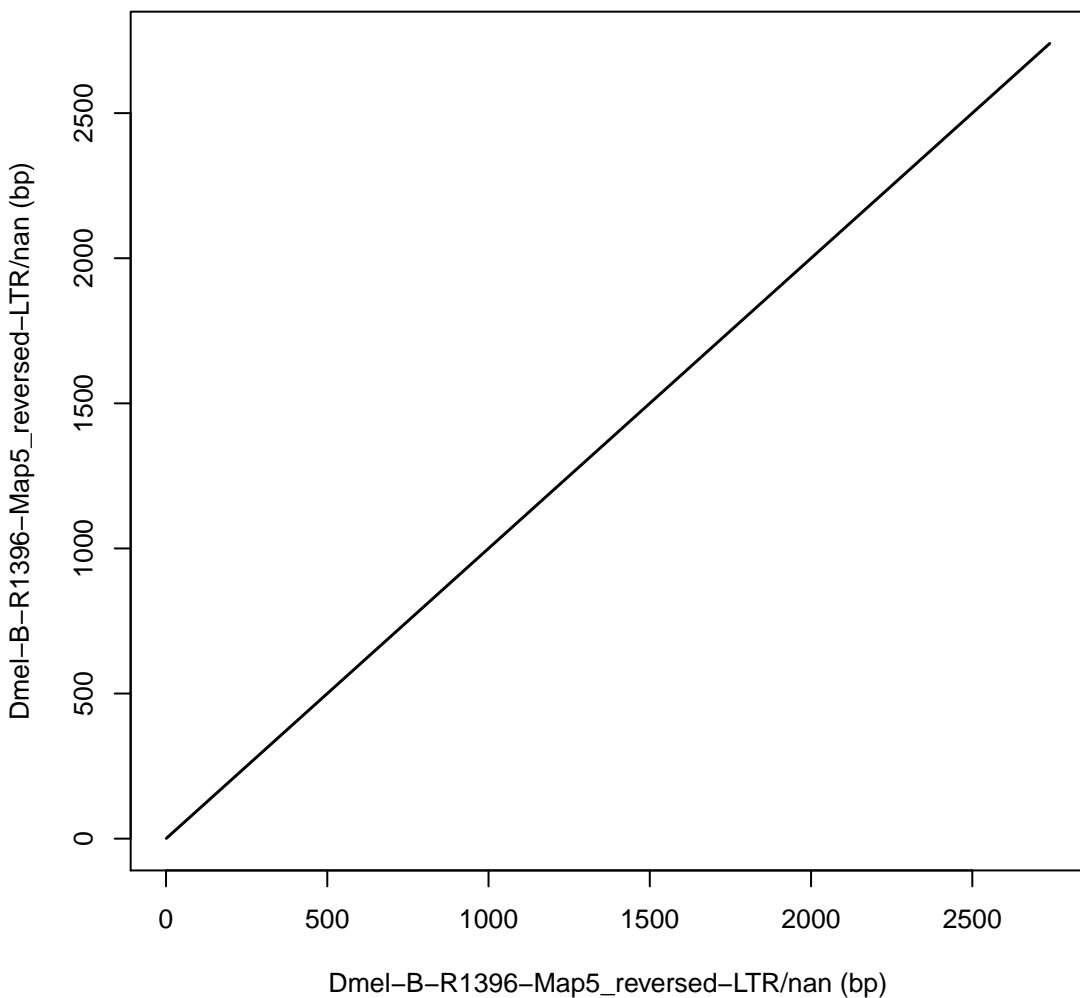
TE: Dmel-B-R1396-Map5_reversed-LTR/nan
consensus size: 2740bp; fragments: 172; full length: 0 (>=2466bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

