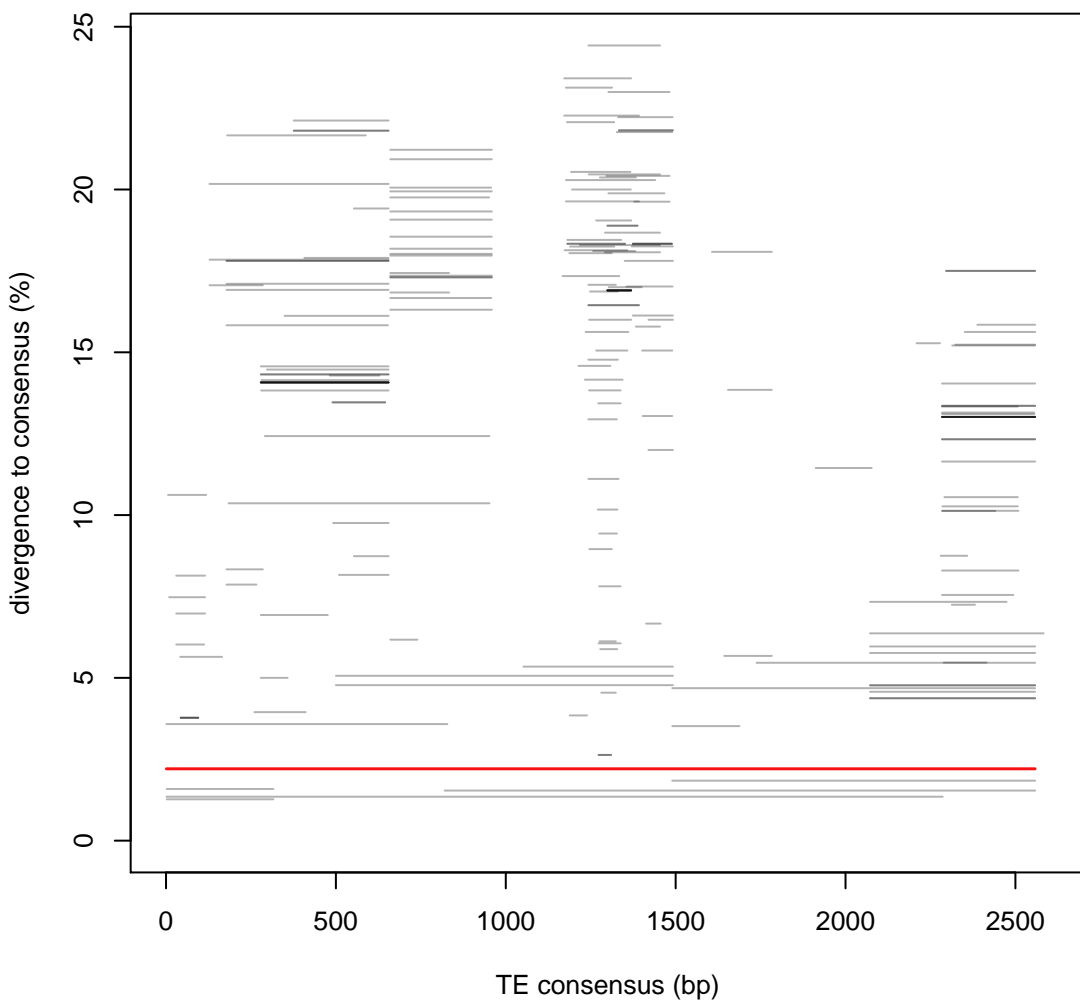
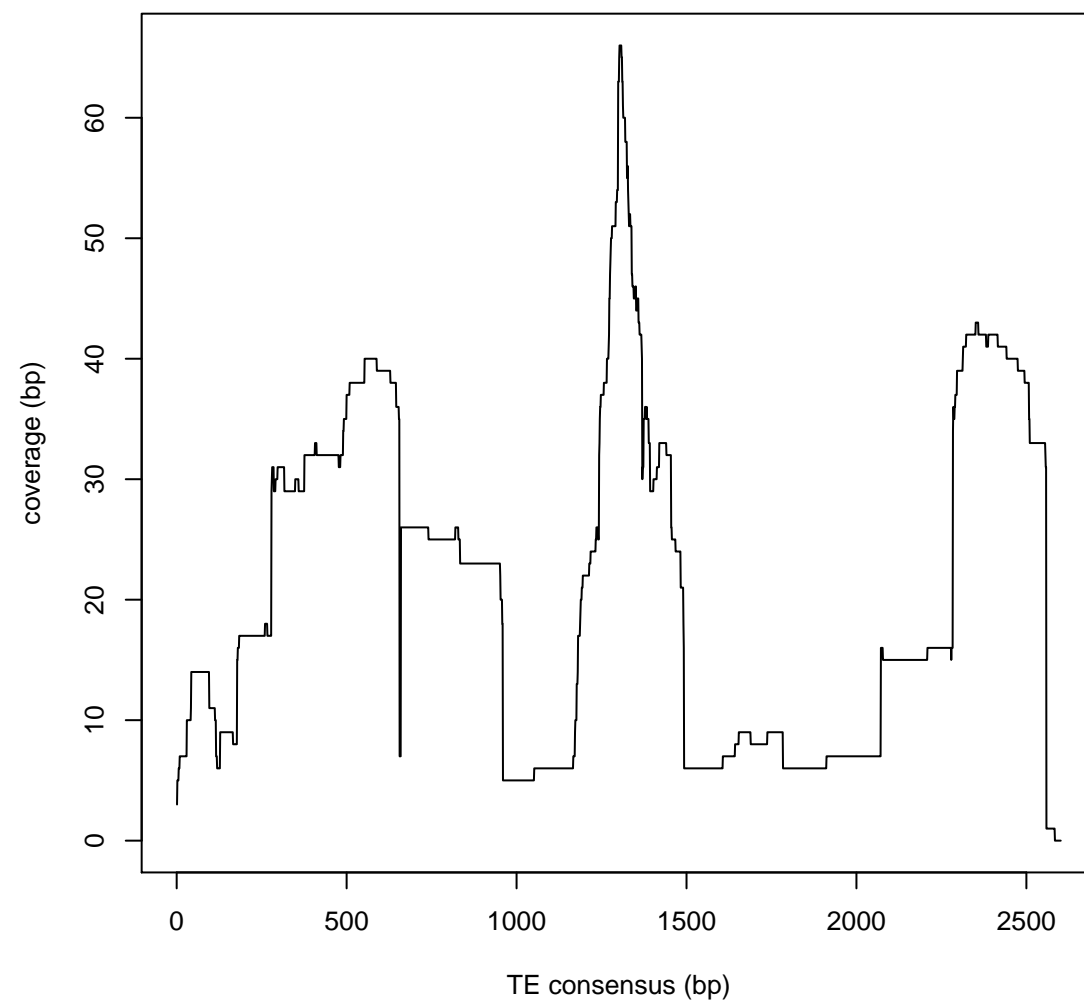


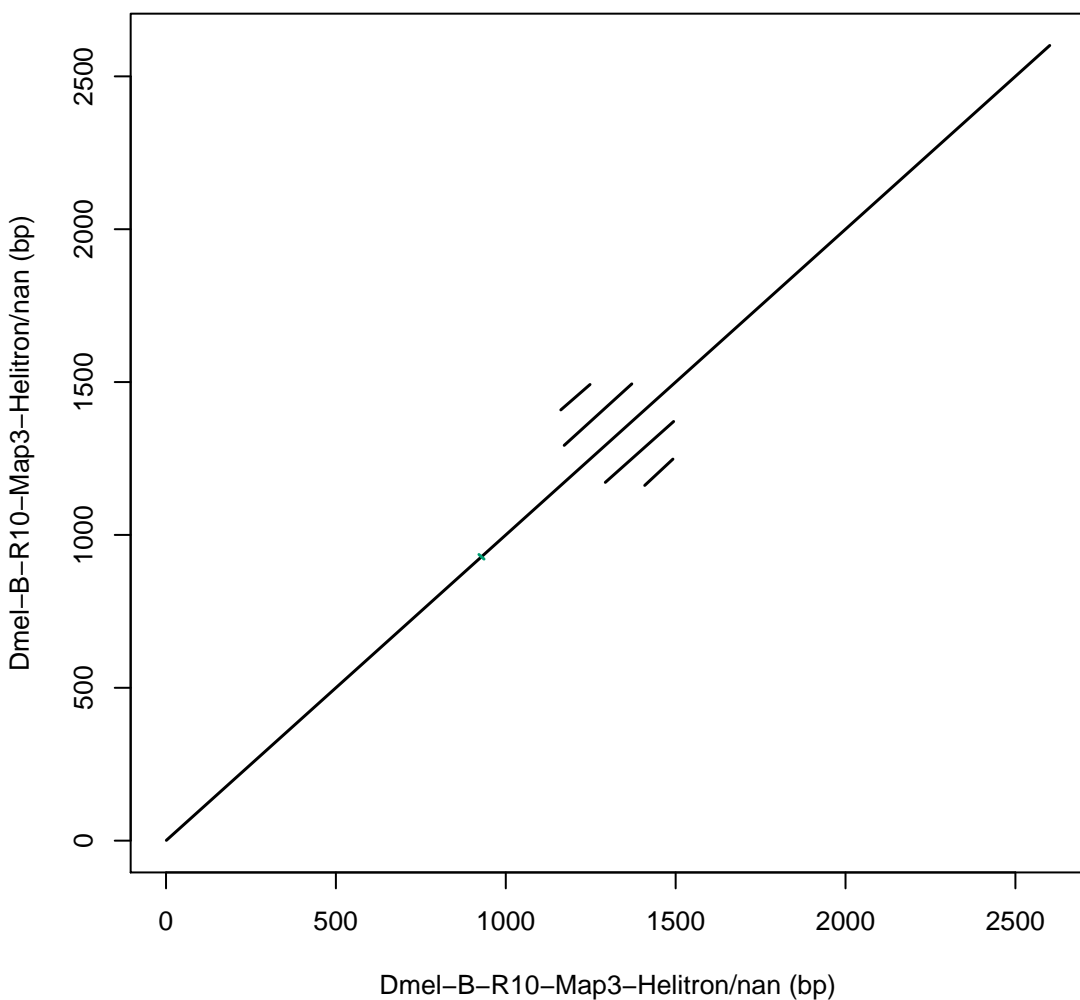
TE: Dmel-B-R10-Map3-Helitron/nan
consensus size: 2601bp; fragments: 205; full length: 1 (≥ 2340.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

