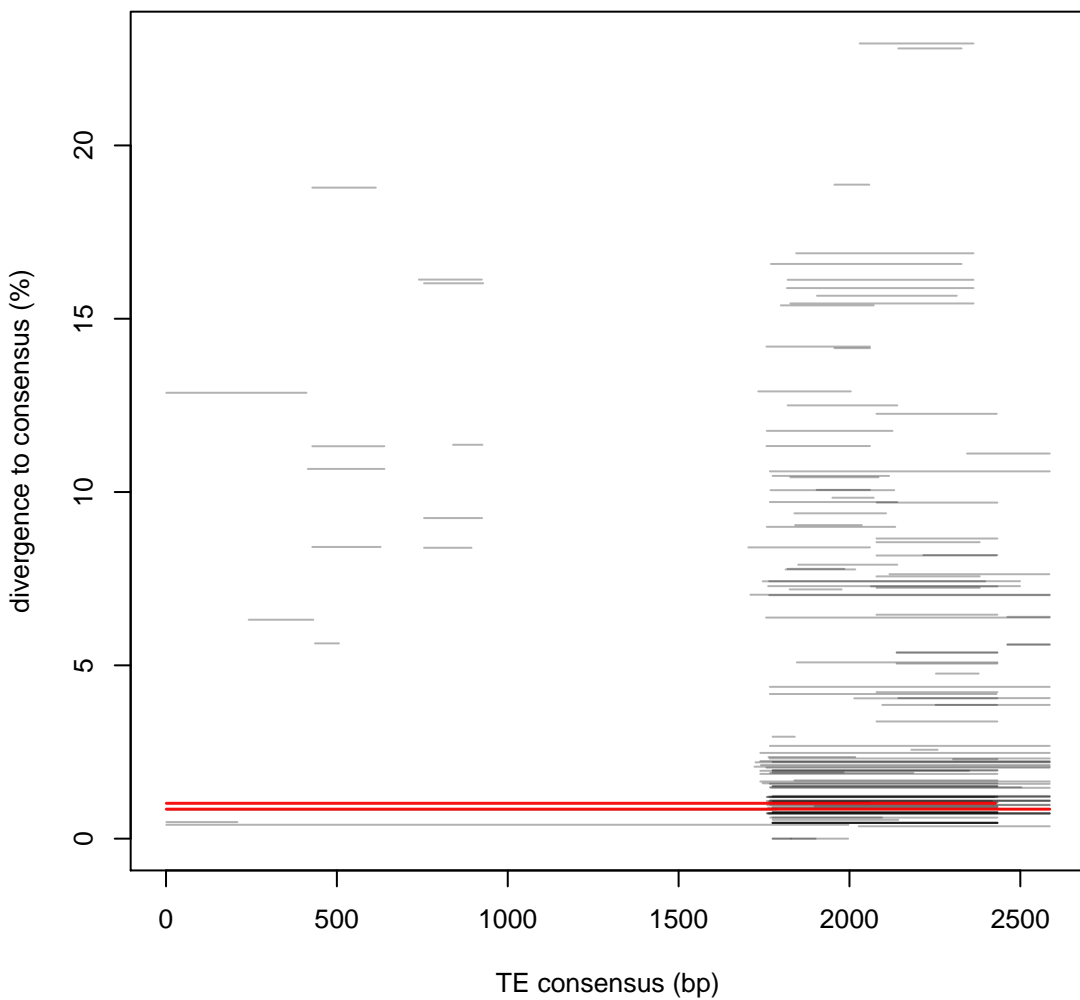
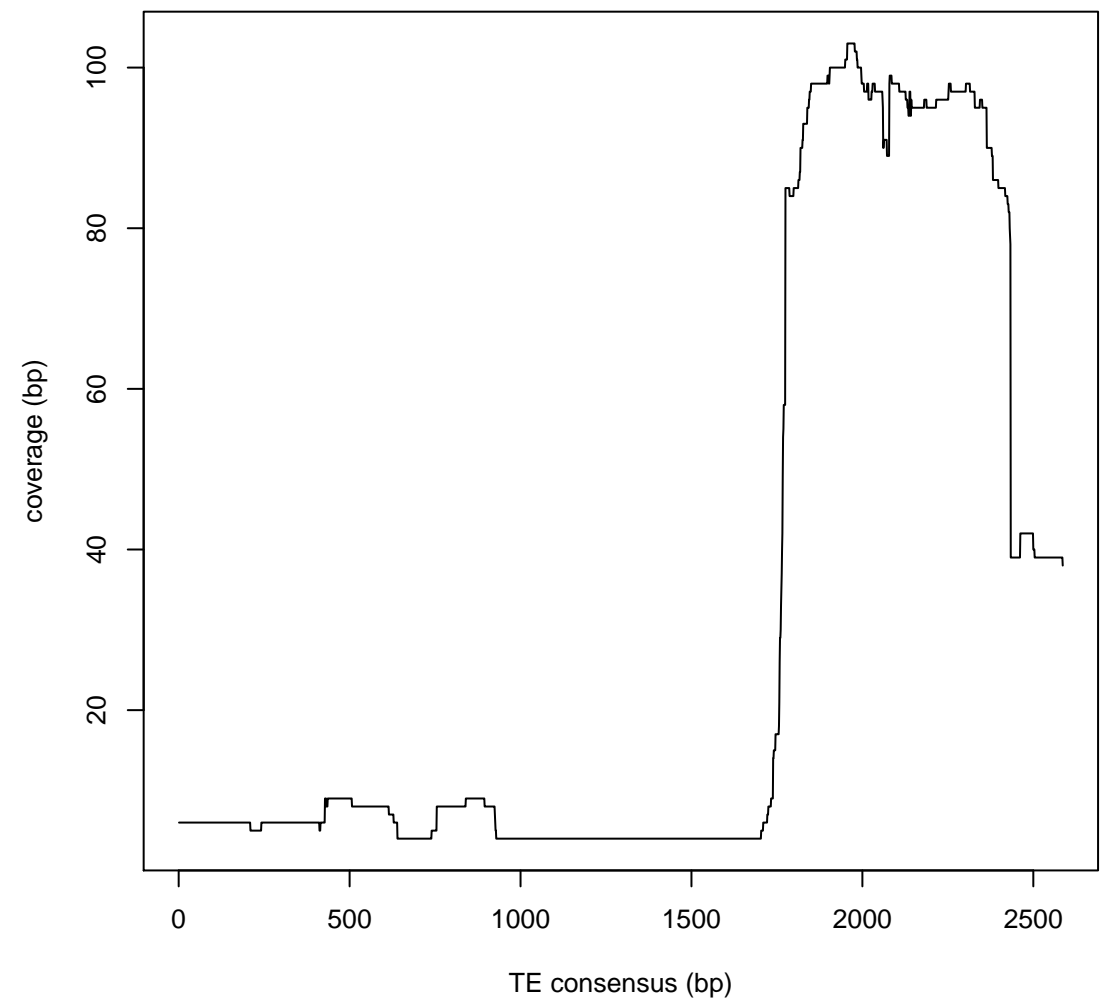


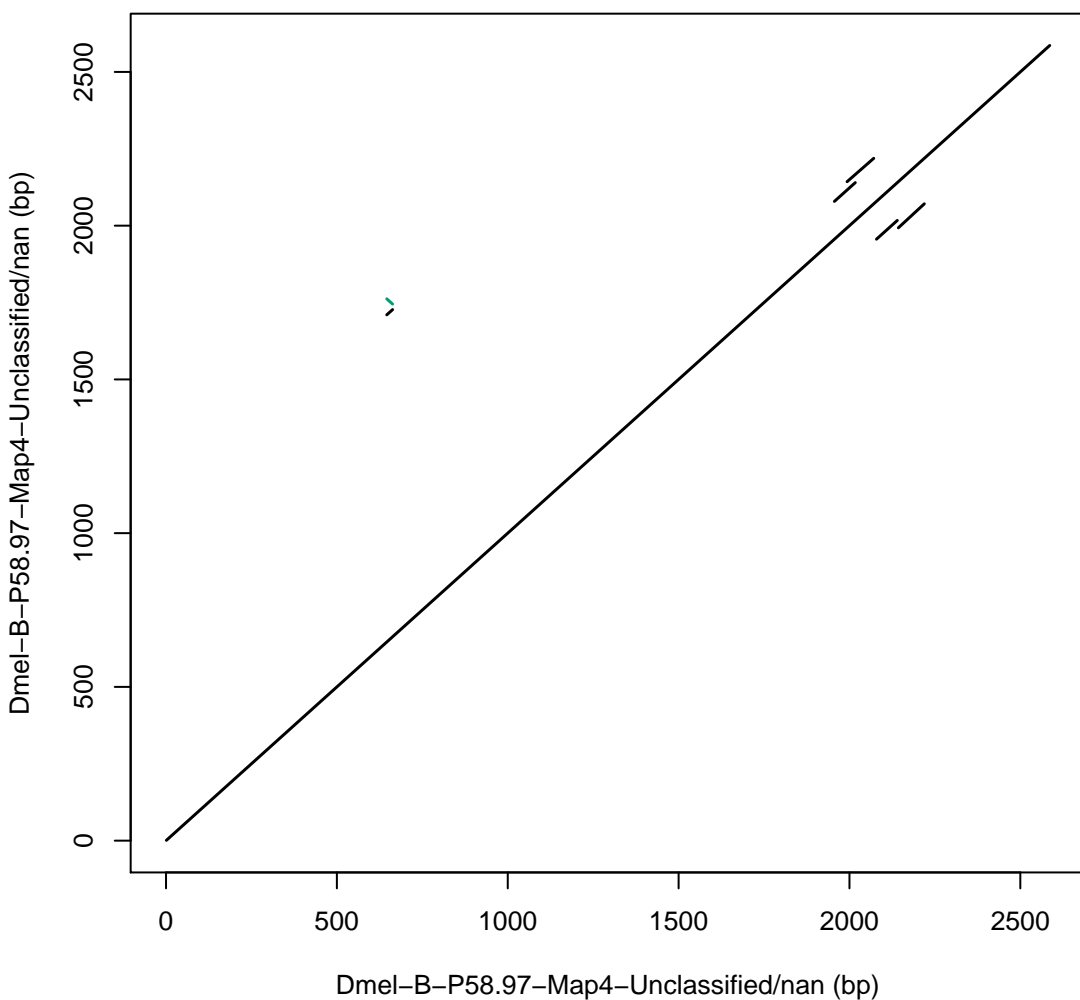
TE: Dmel-B-P58.97-Map4-Unclassified/nan
consensus size: 2586bp; fragments: 150; full length: 2 (≥ 2327.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

