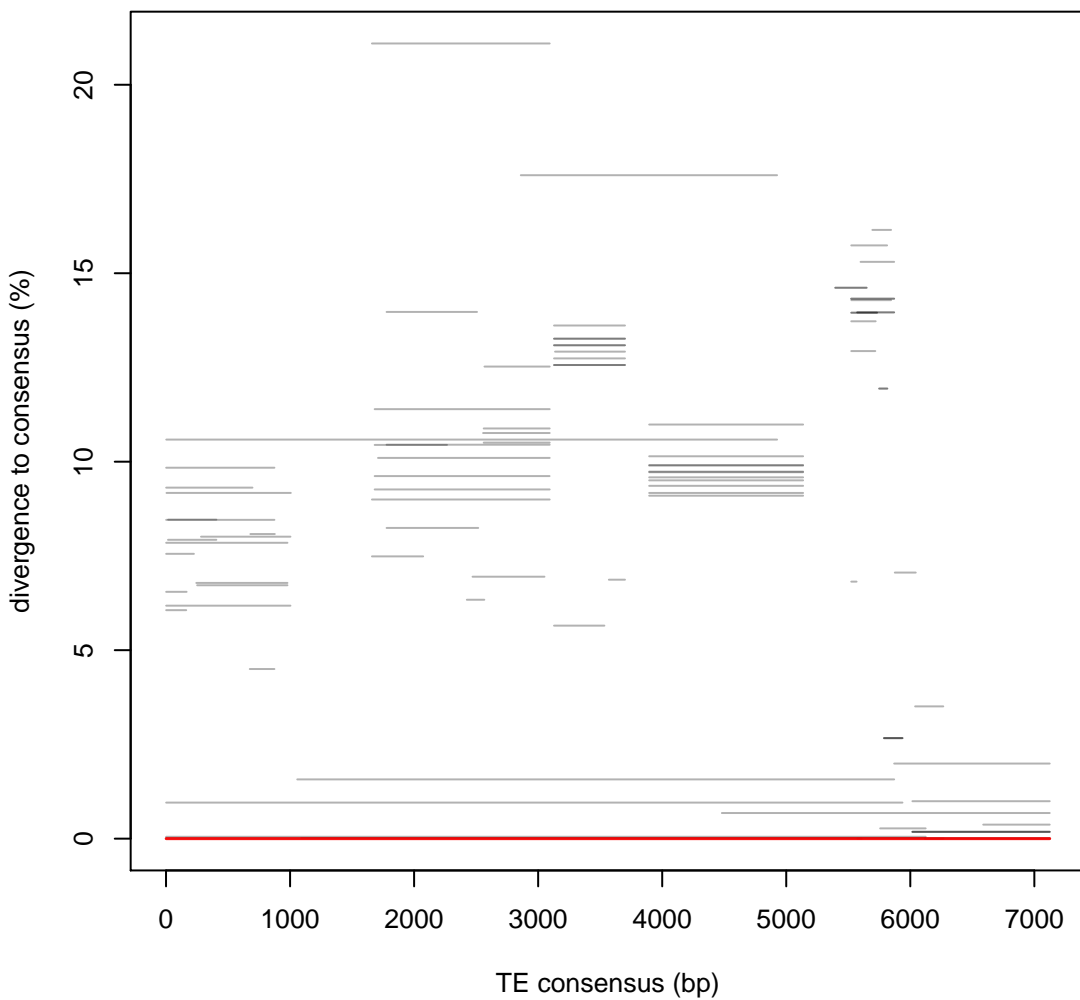
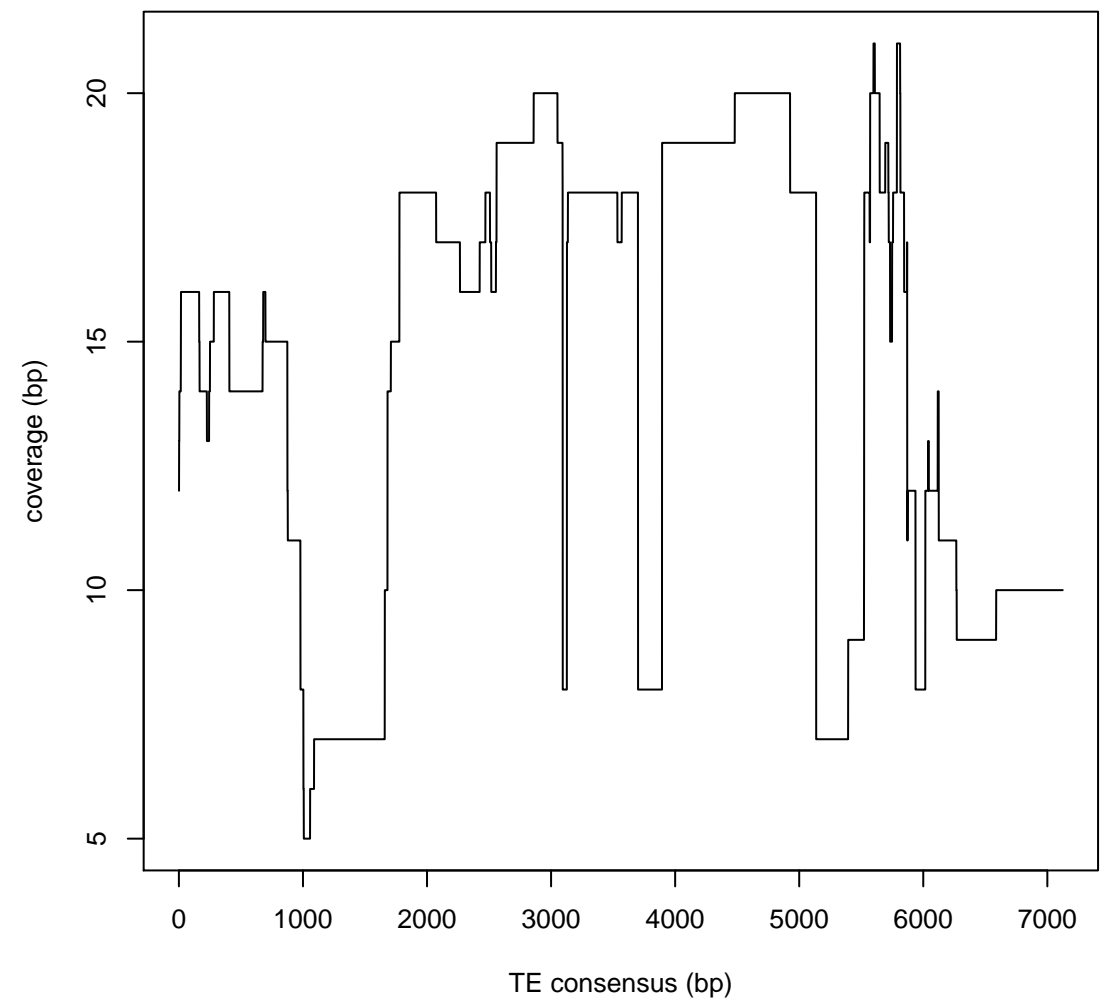


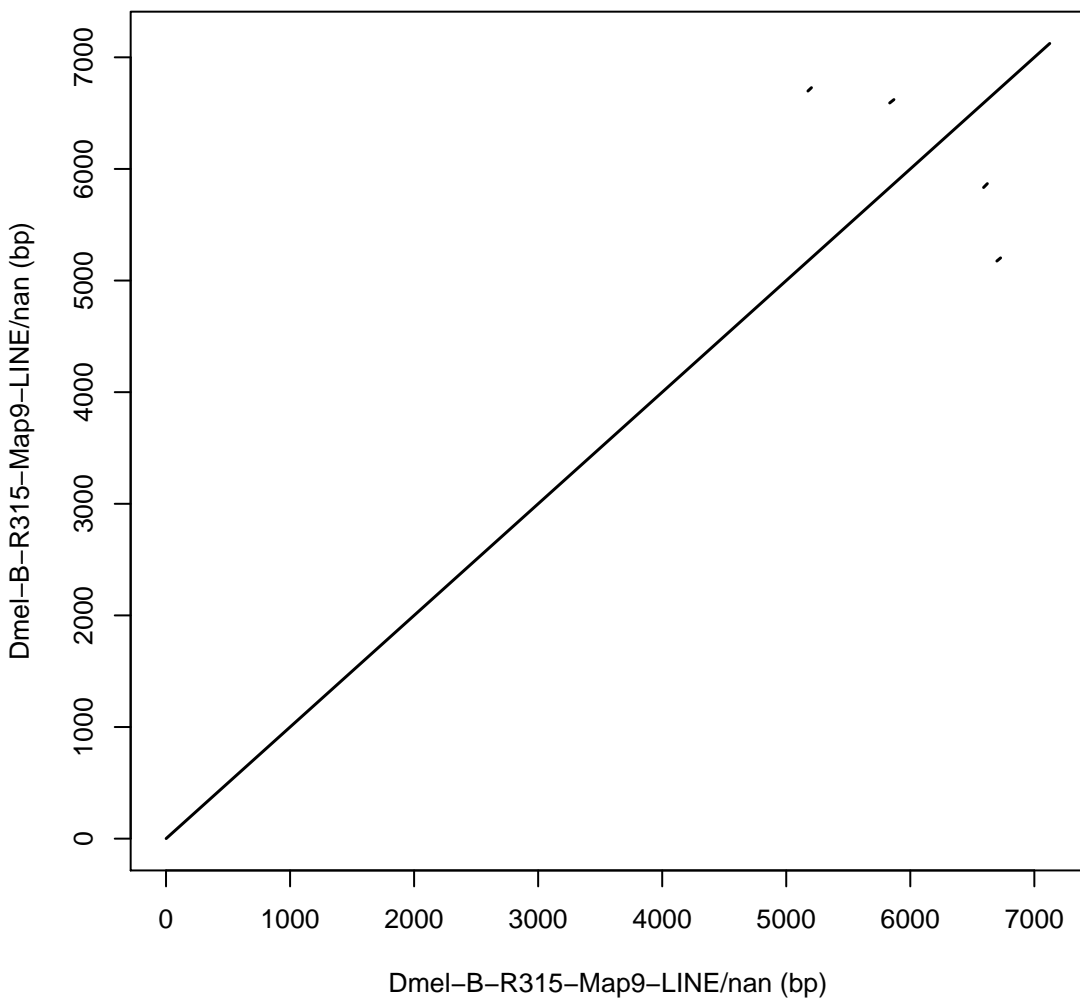
TE: Dmel-B-R315-Map9-LINE/nan
consensus size: 7124bp; fragments: 96; full length: 1 (≥ 6411.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

