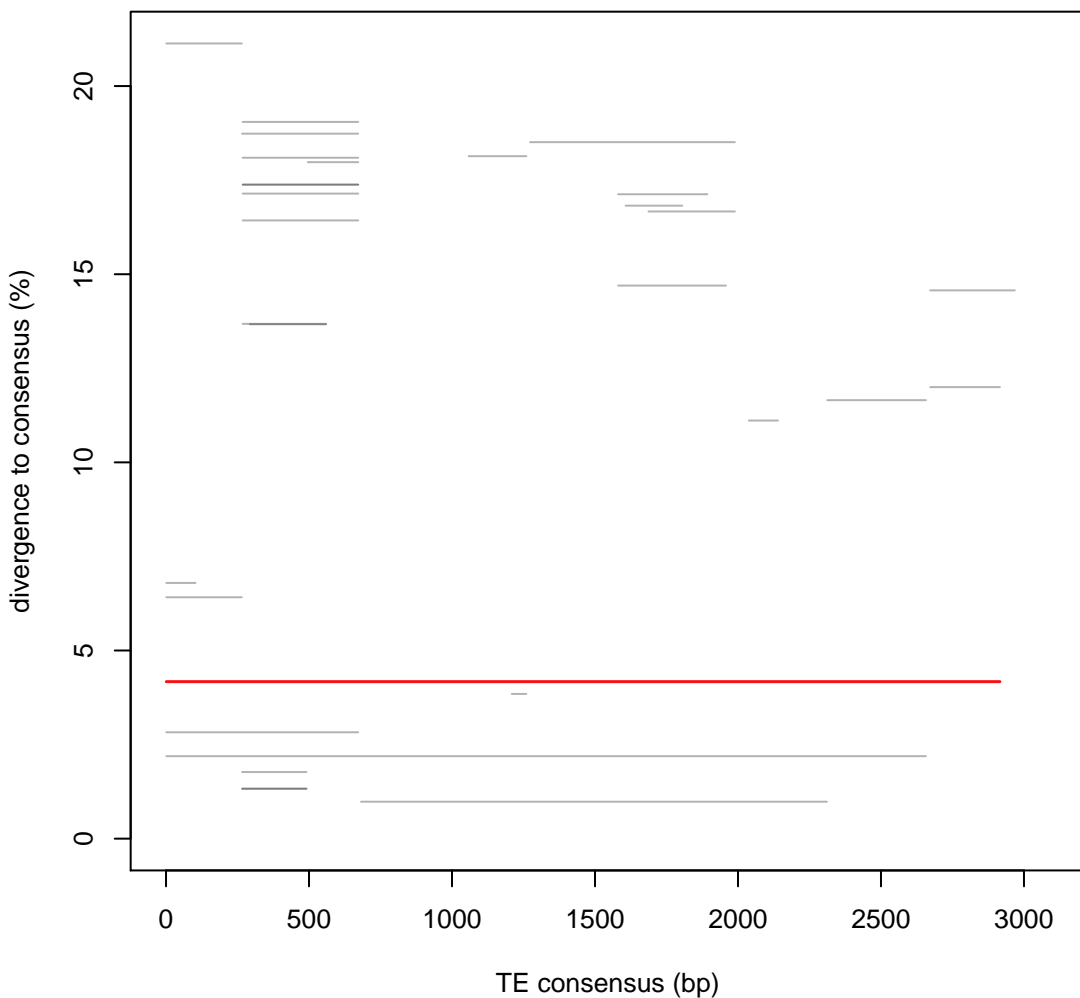
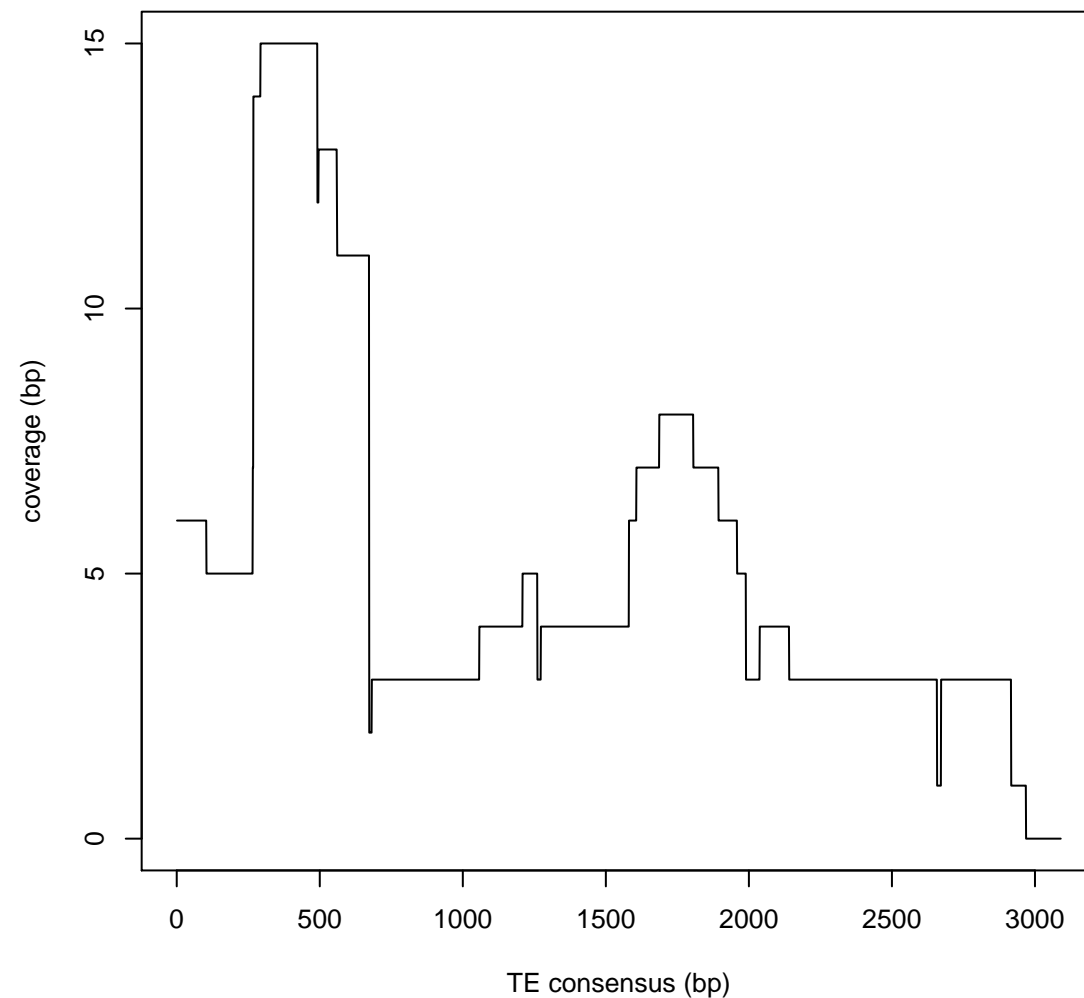


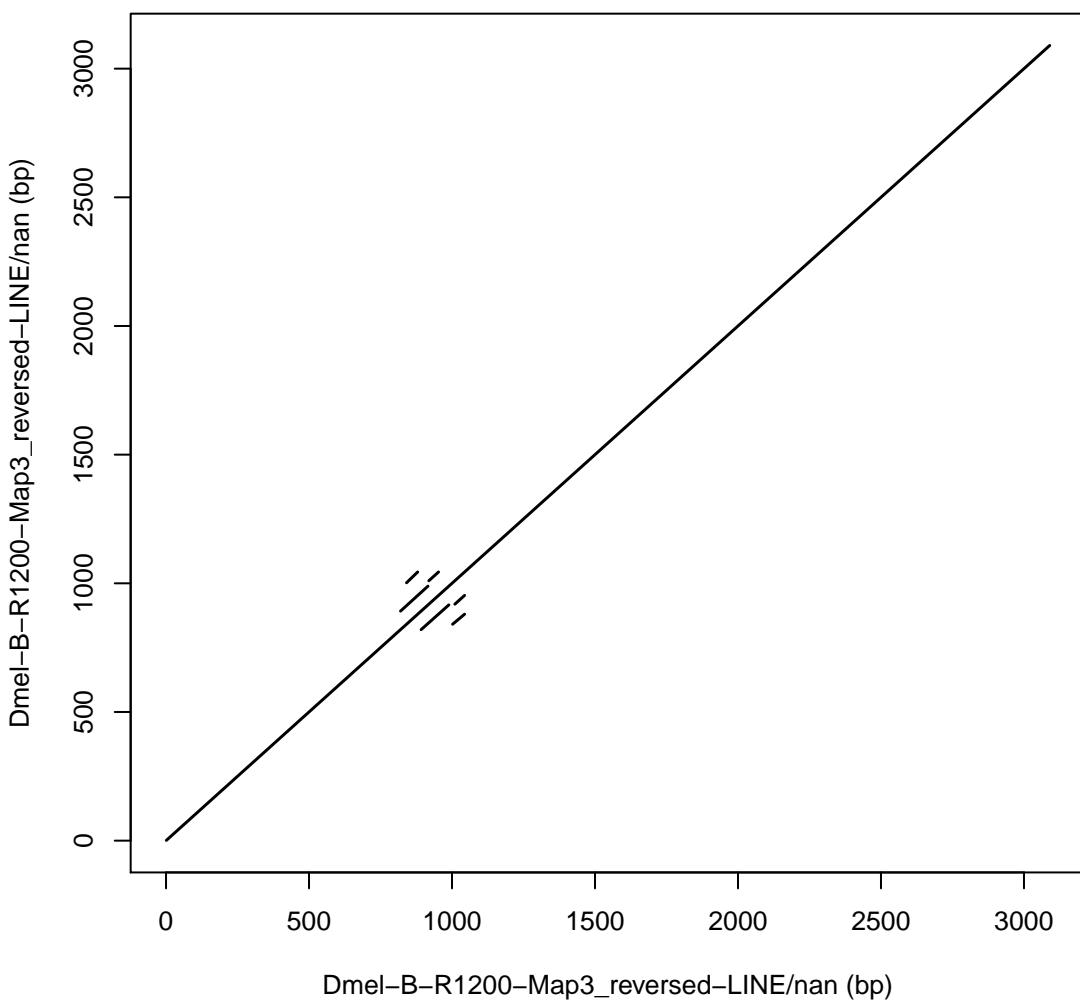
TE: Dmel-B-R1200-Map3_reversed-LINE/nan
consensus size: 3090bp; fragments: 31; full length: 1 (>=2781bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

