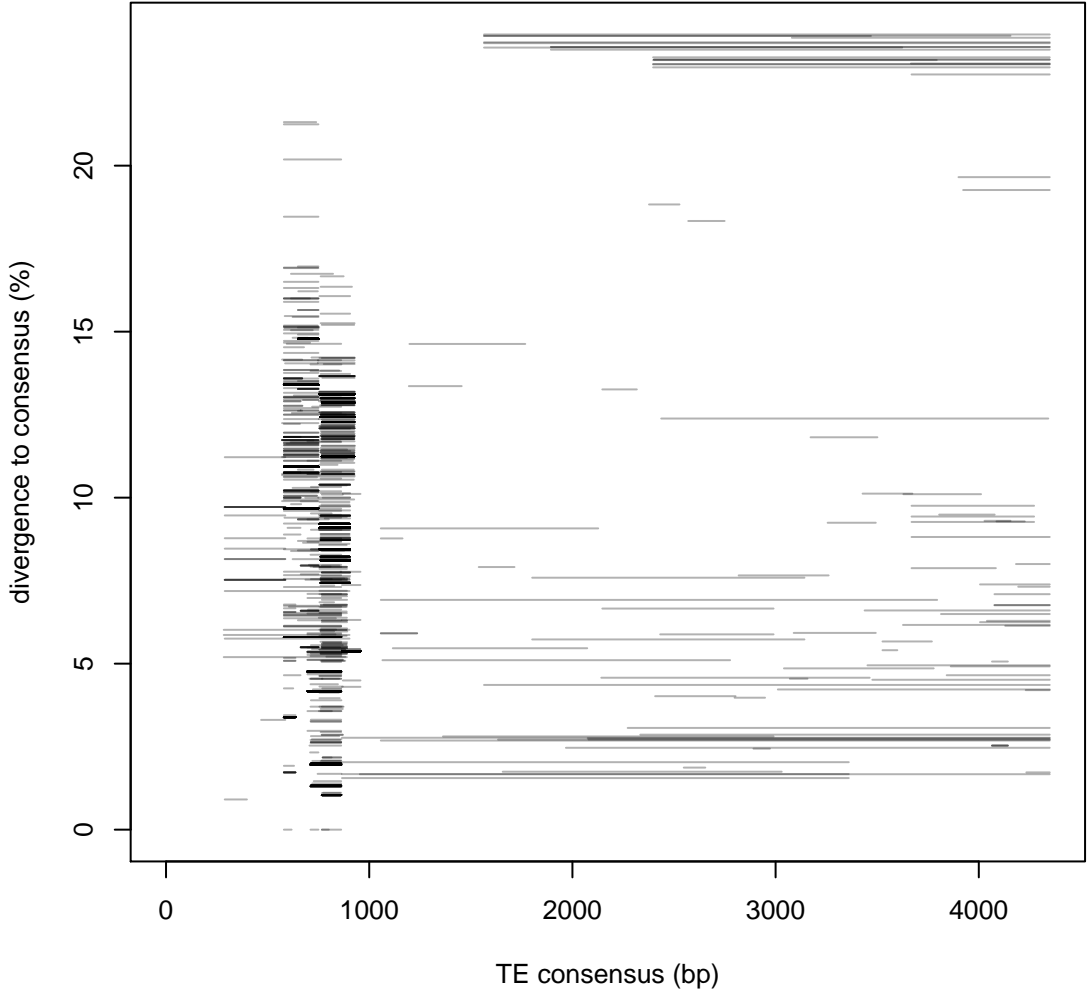
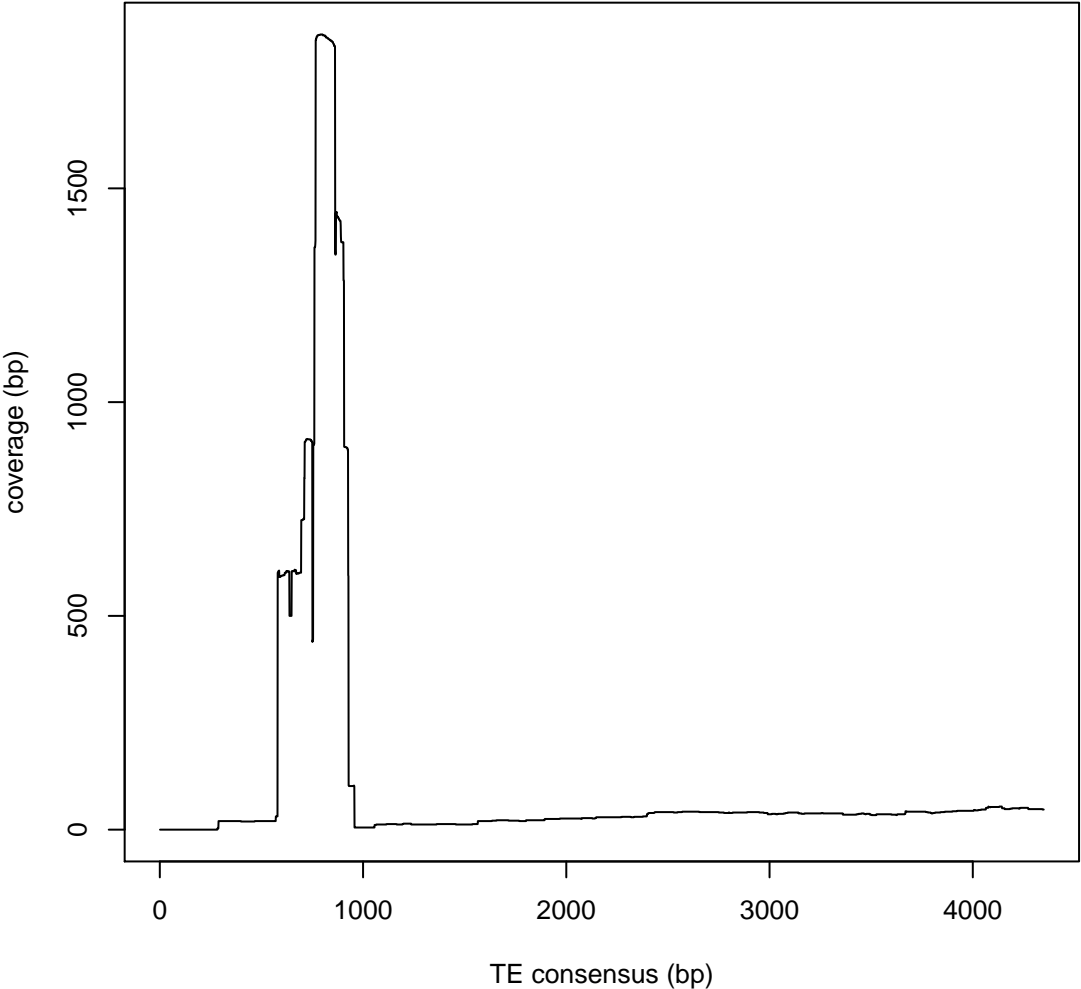


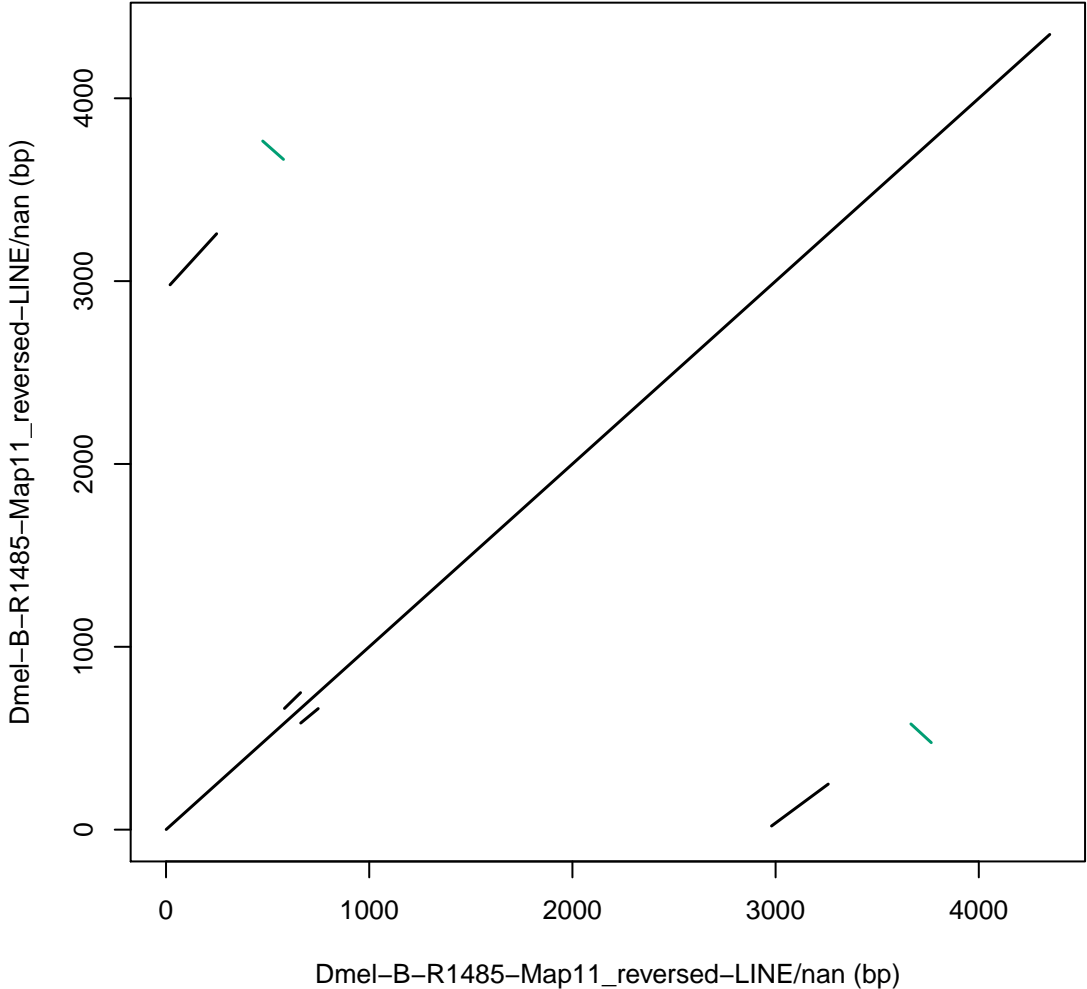
TE: Dmel-B-R1485-Map11_reversed-LINE/nan
consensus size: 4349bp; fragments: 2696; full length: 0 (>=3914.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

