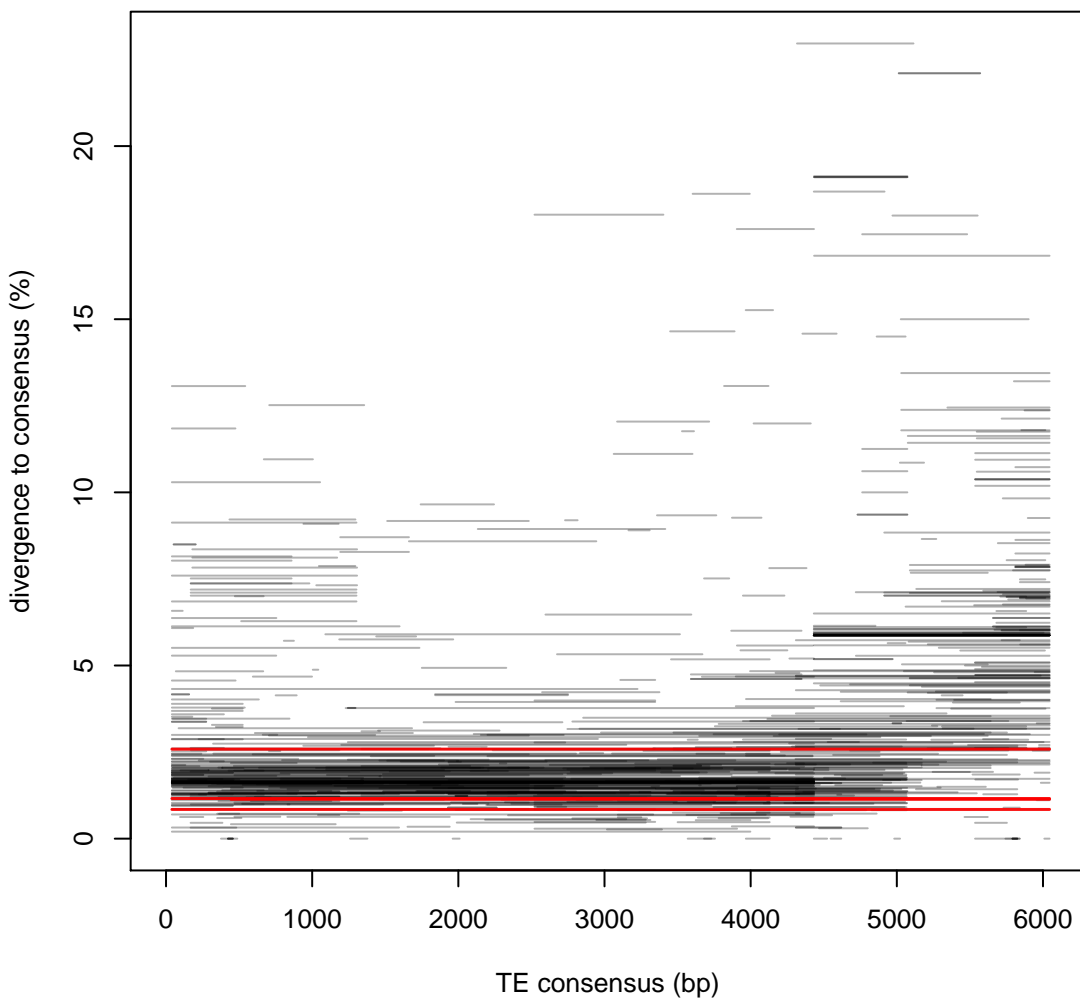
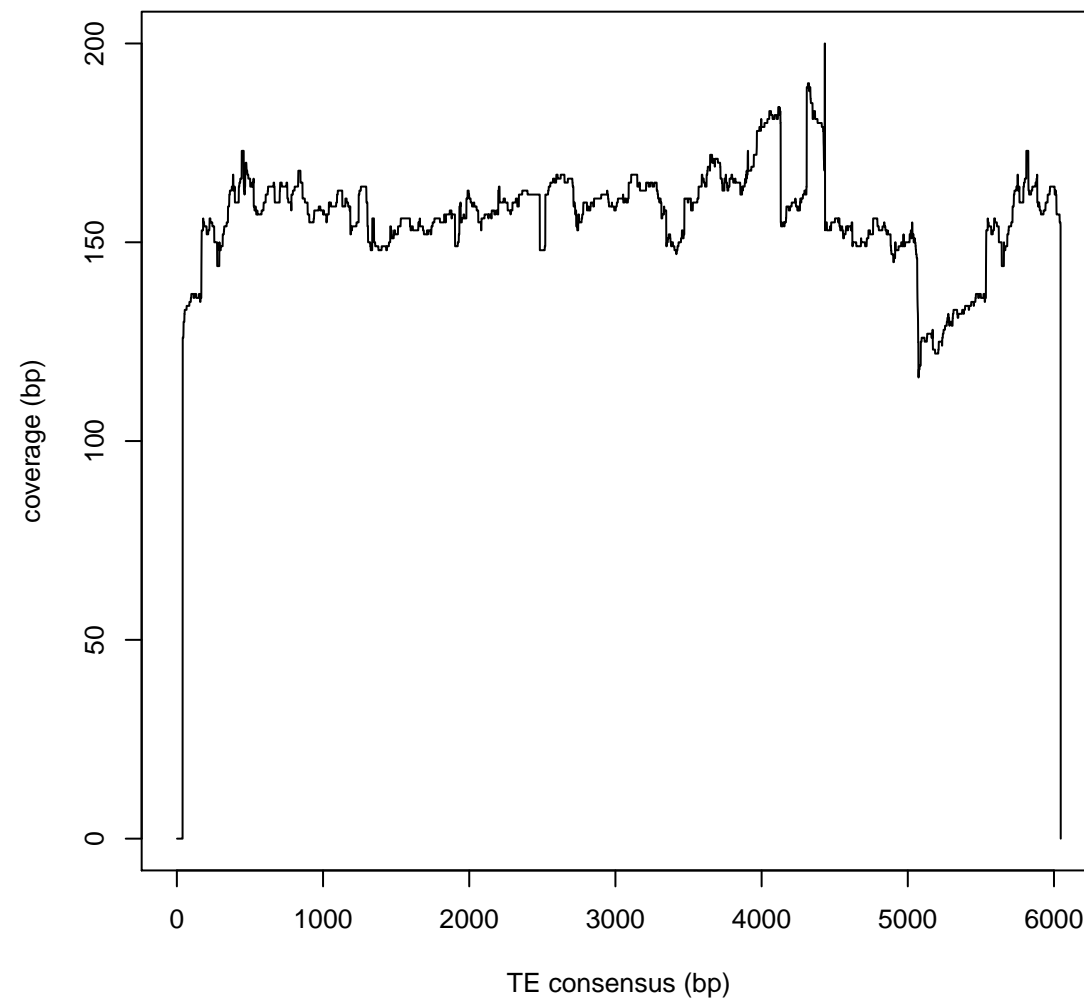


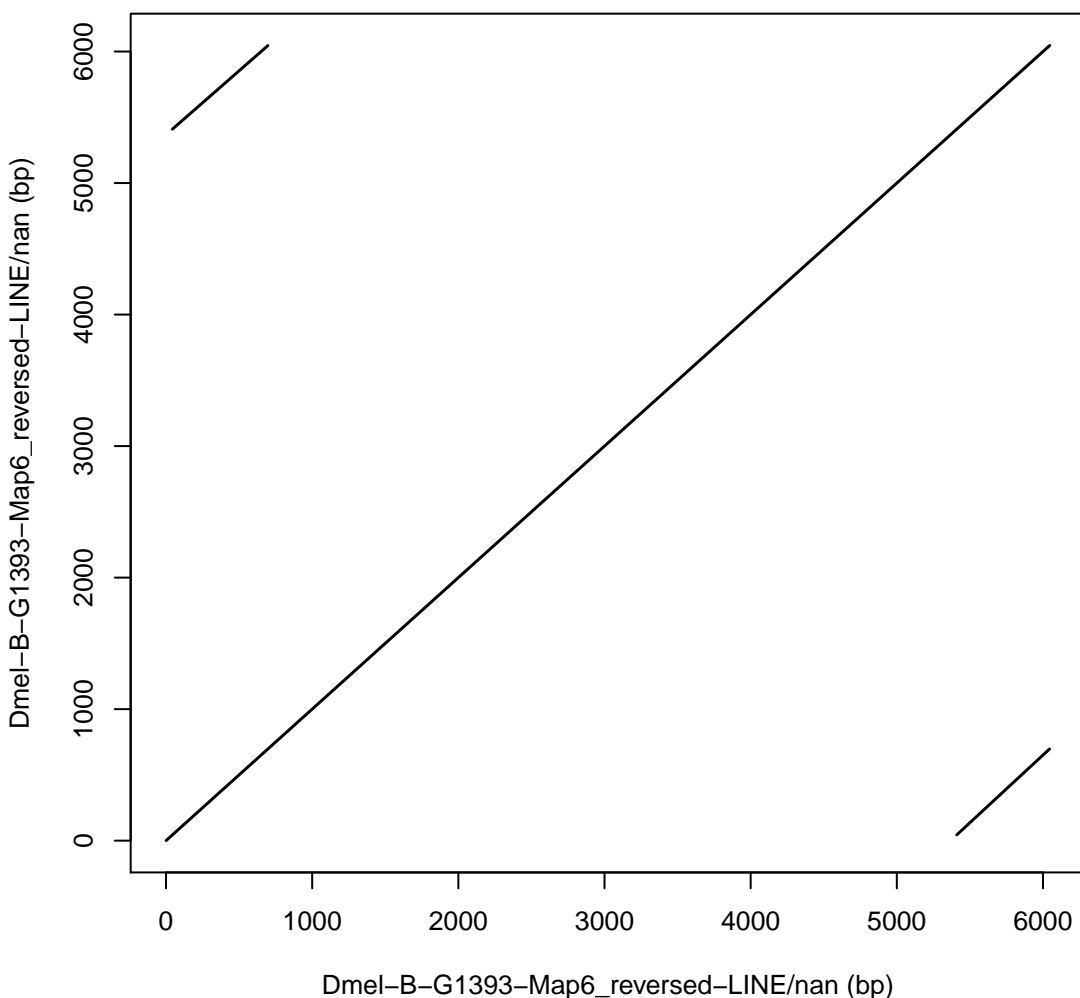
TE: Dmel-B-G1393-Map6_reversed-LINE/nan
consensus size: 6046bp; fragments: 779; full length: 4 (>=5441.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

