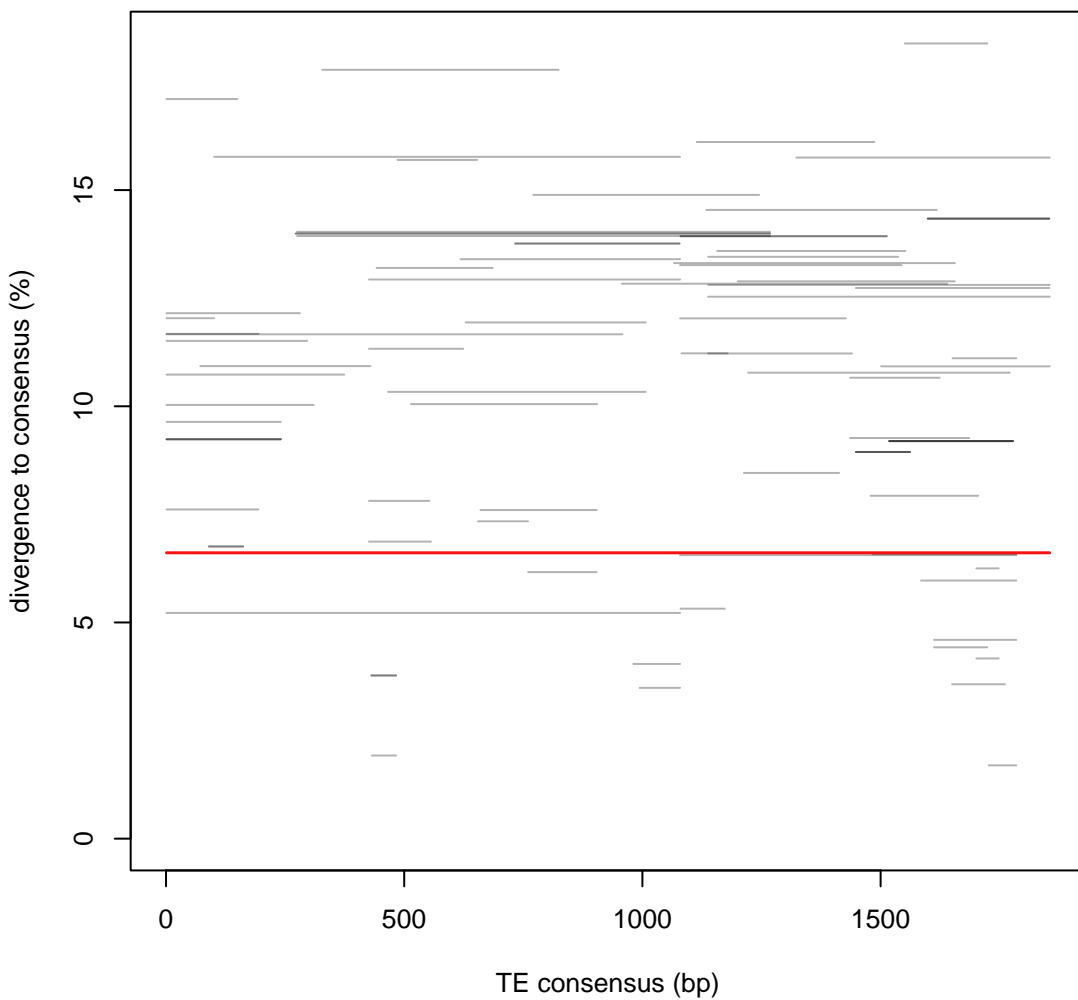
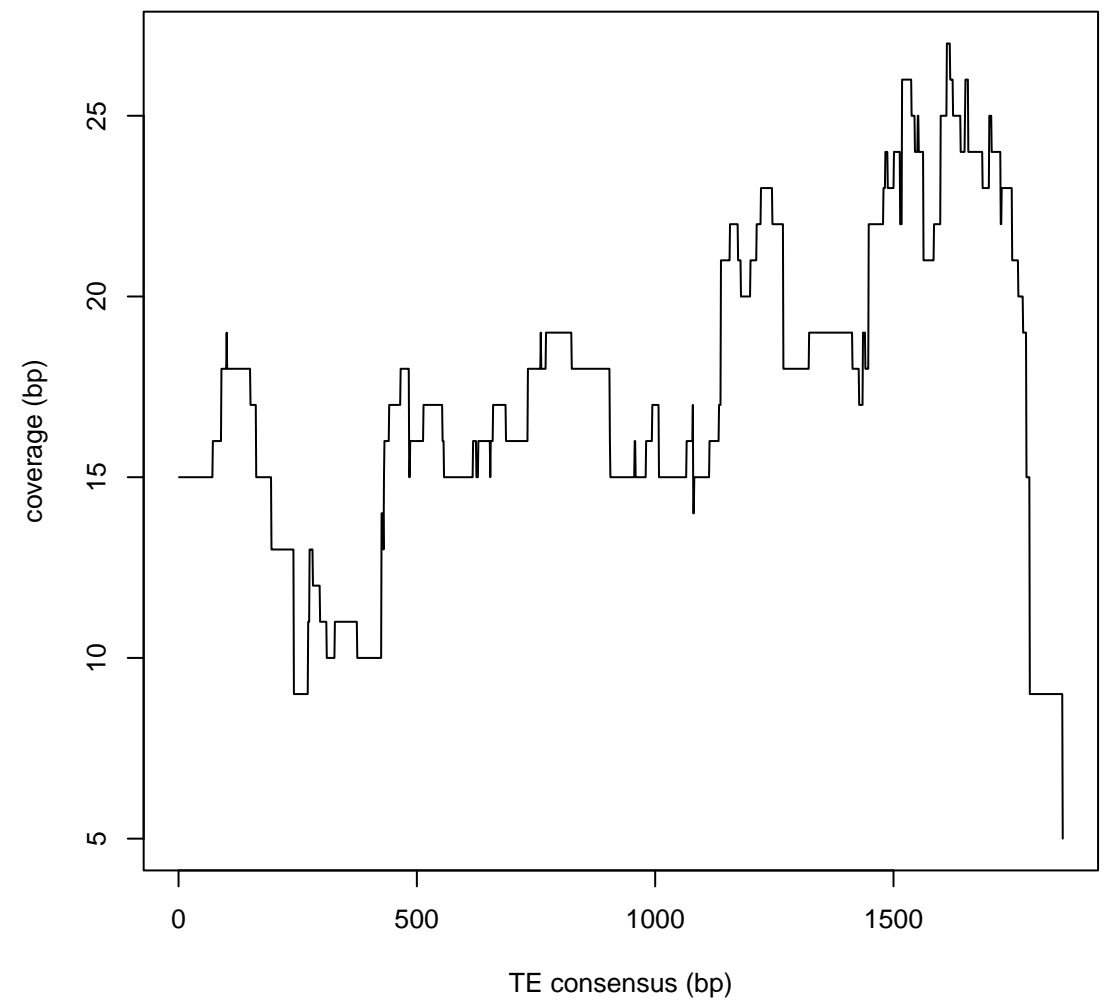


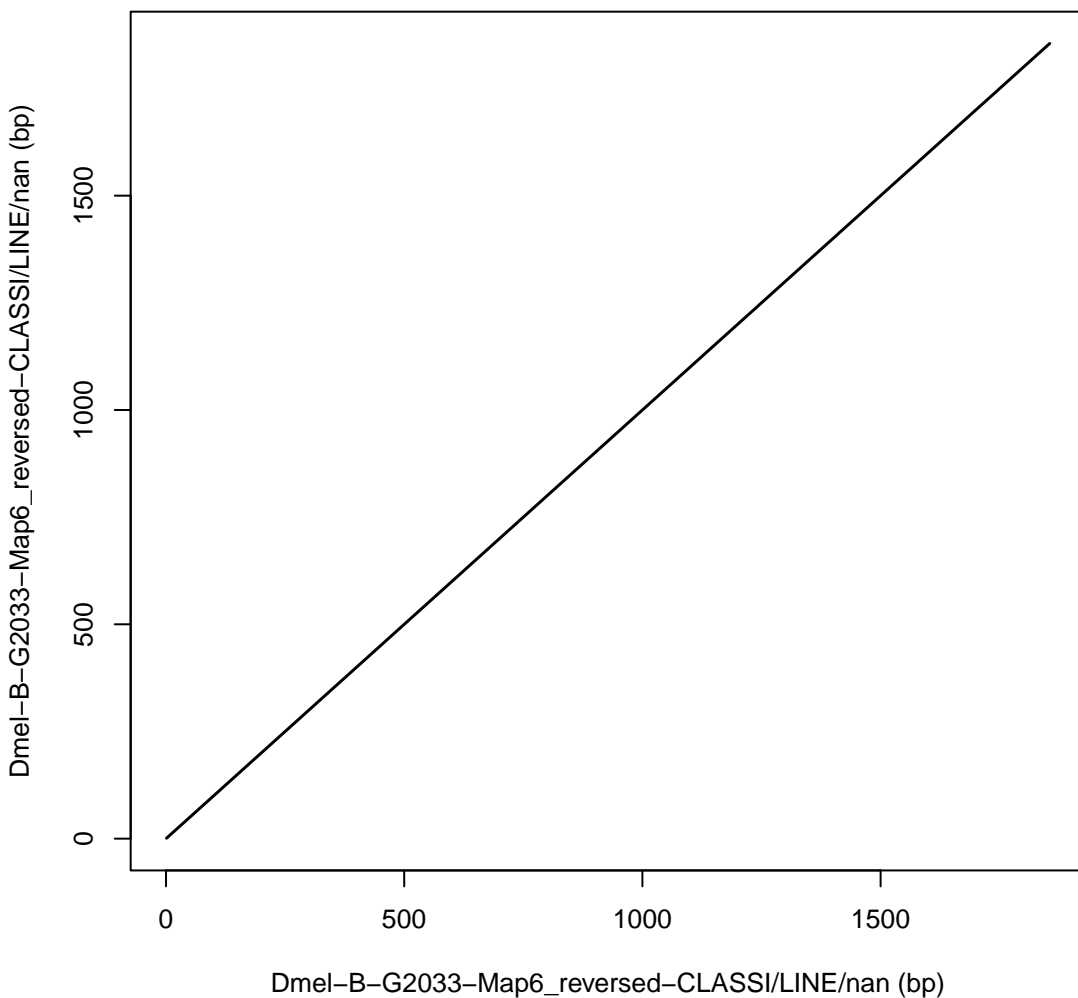
TE: Dmel-B-G2033-Map6_reversed-CLASSI/LINE/nan
consensus size: 1855bp; fragments: 90; full length: 1 (≥ 1669.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

