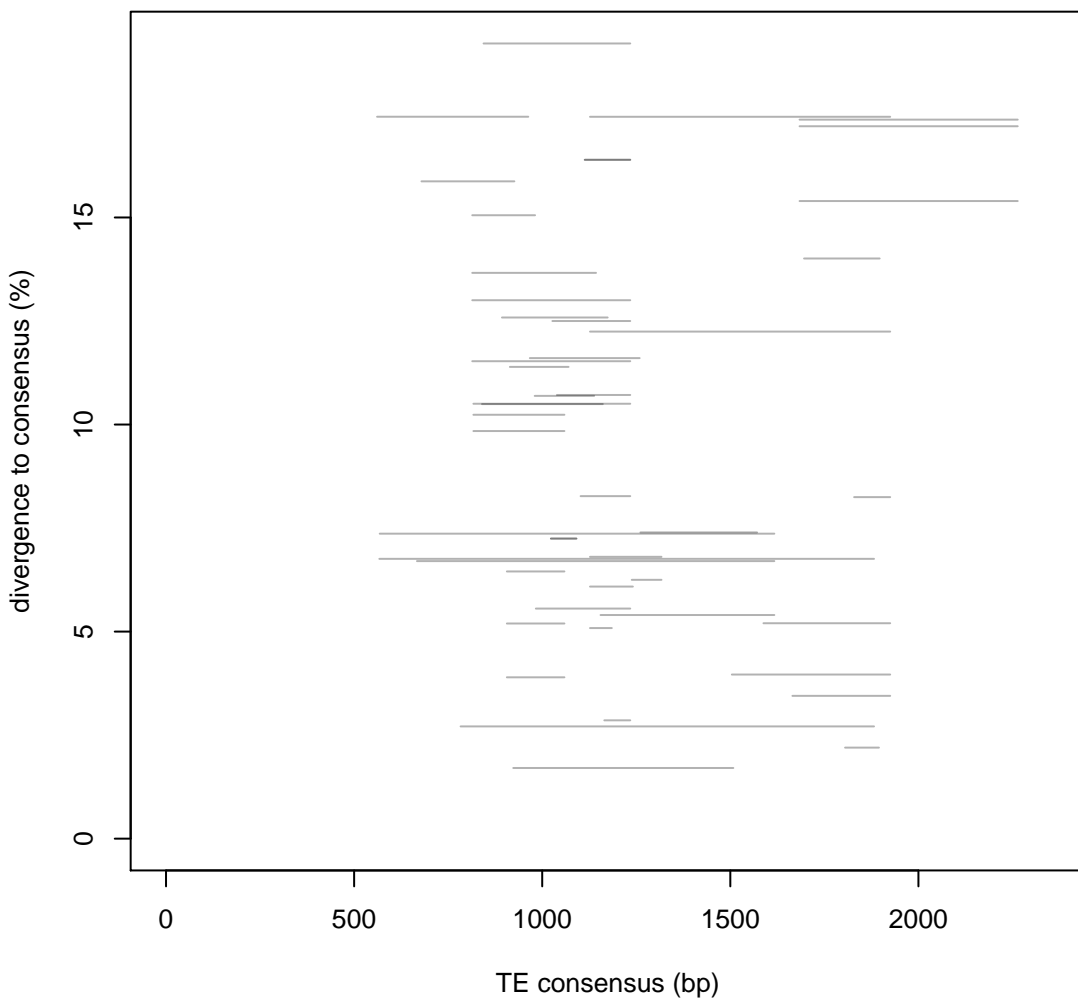
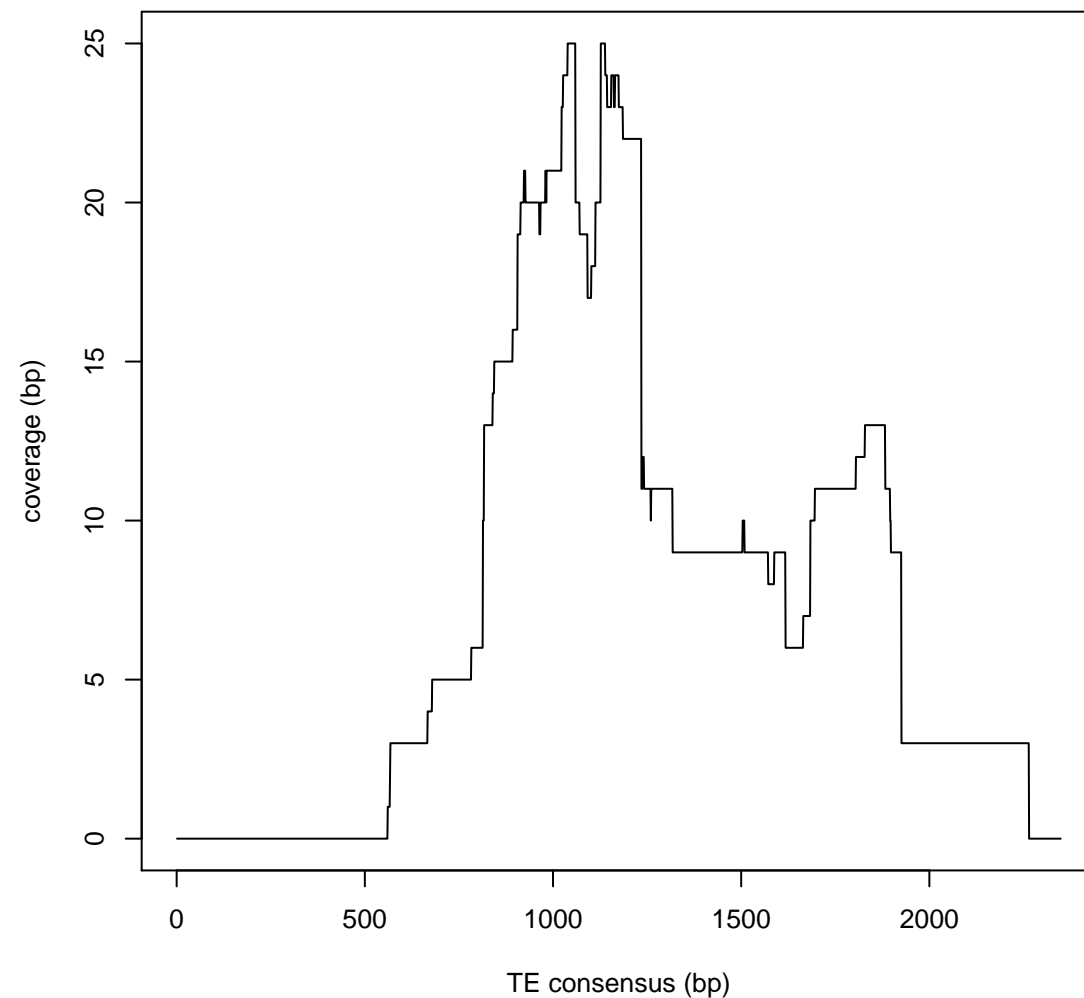


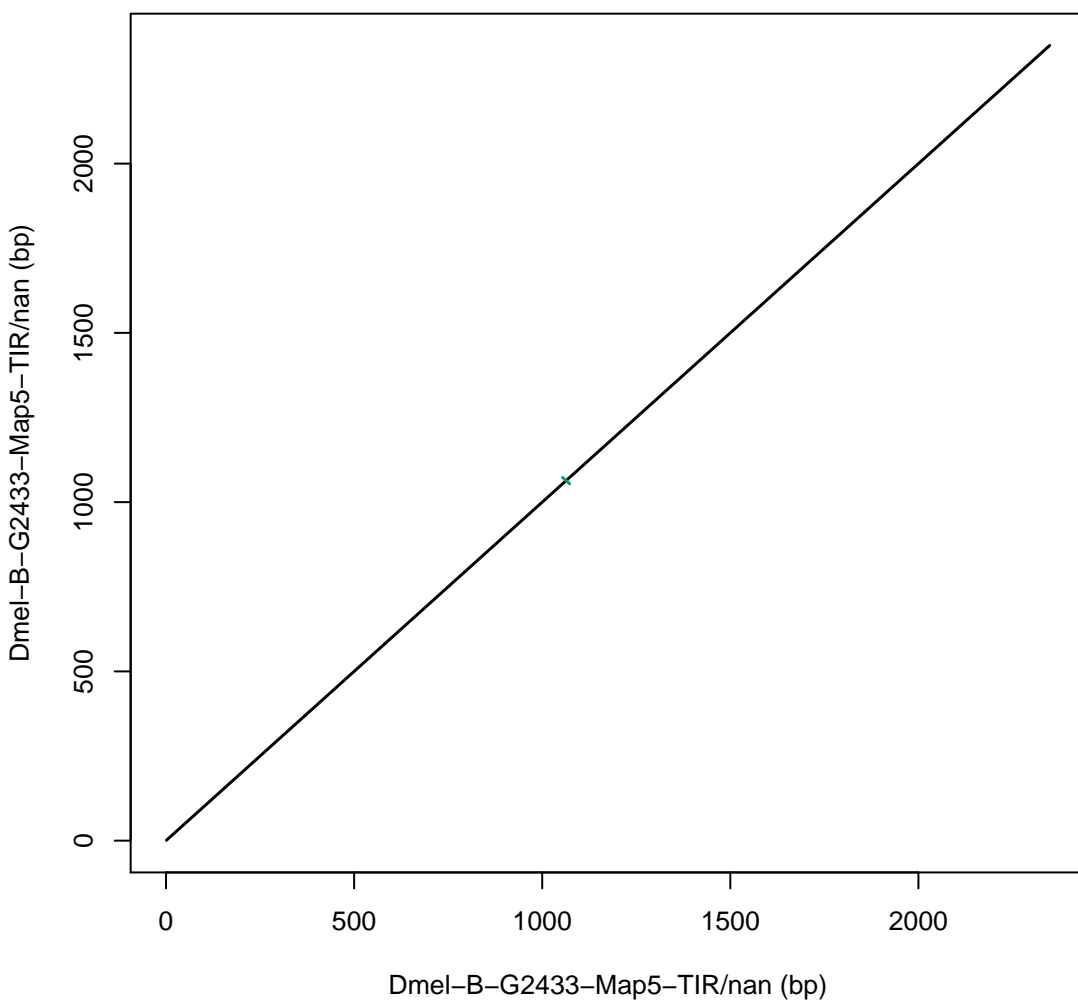
TE: Dmel-B-G2433-Map5-TIR/nan
consensus size: 2349bp; fragments: 49; full length: 0 (>=2114.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

