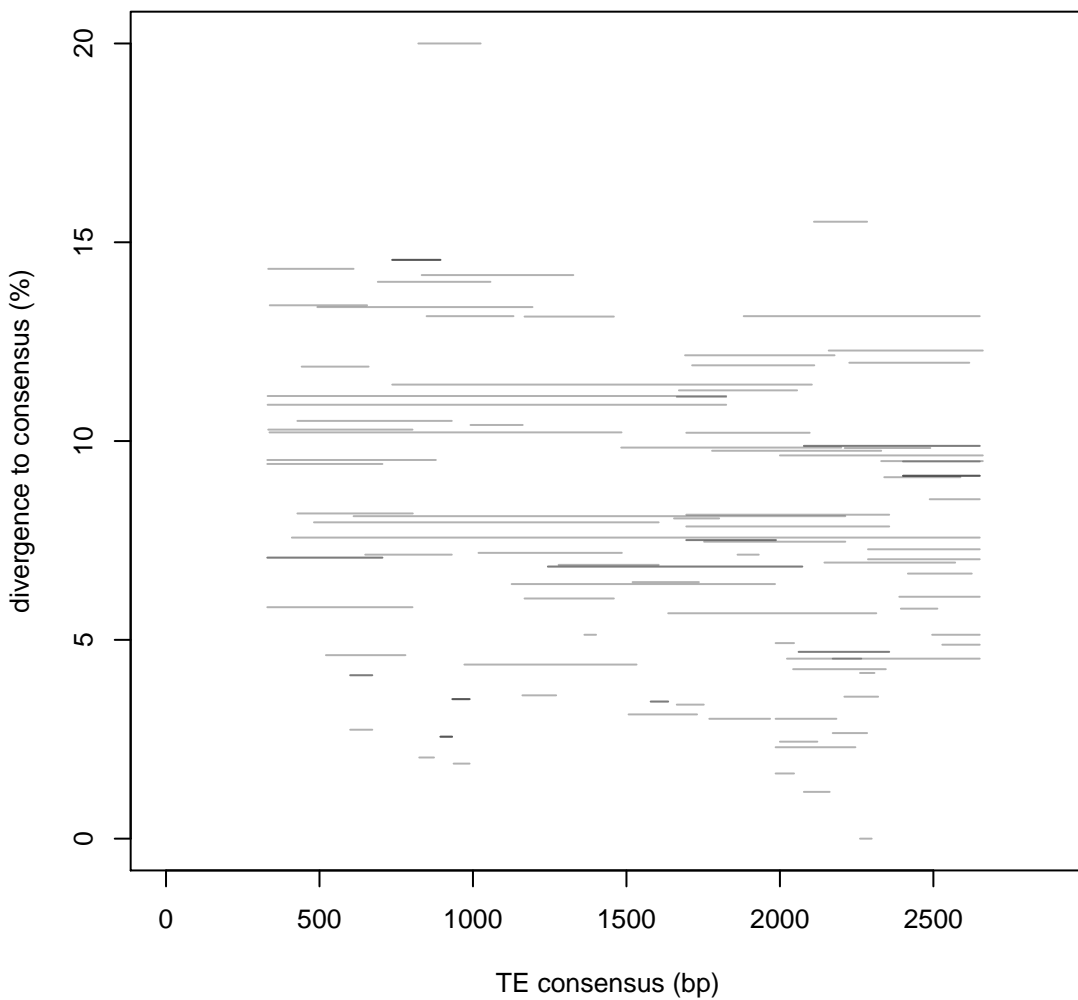
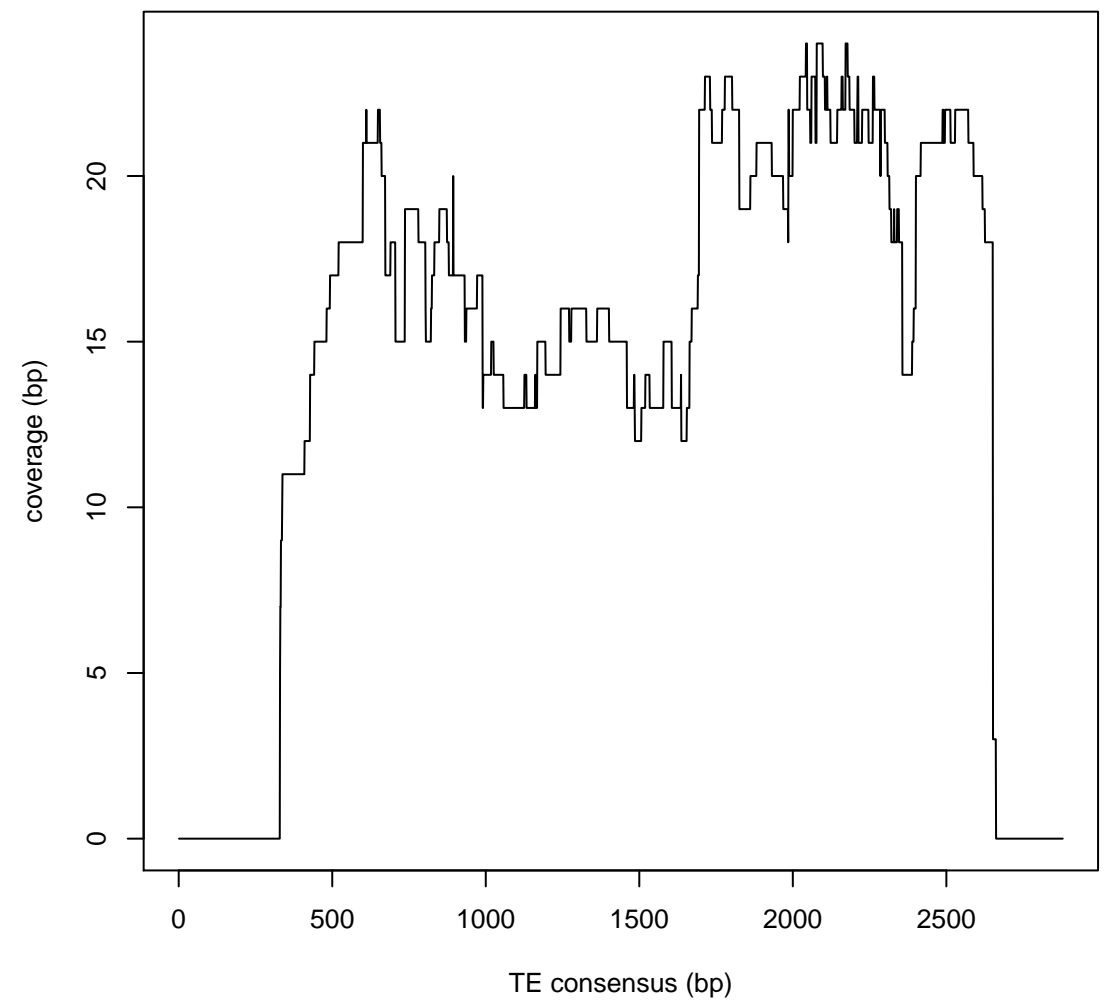


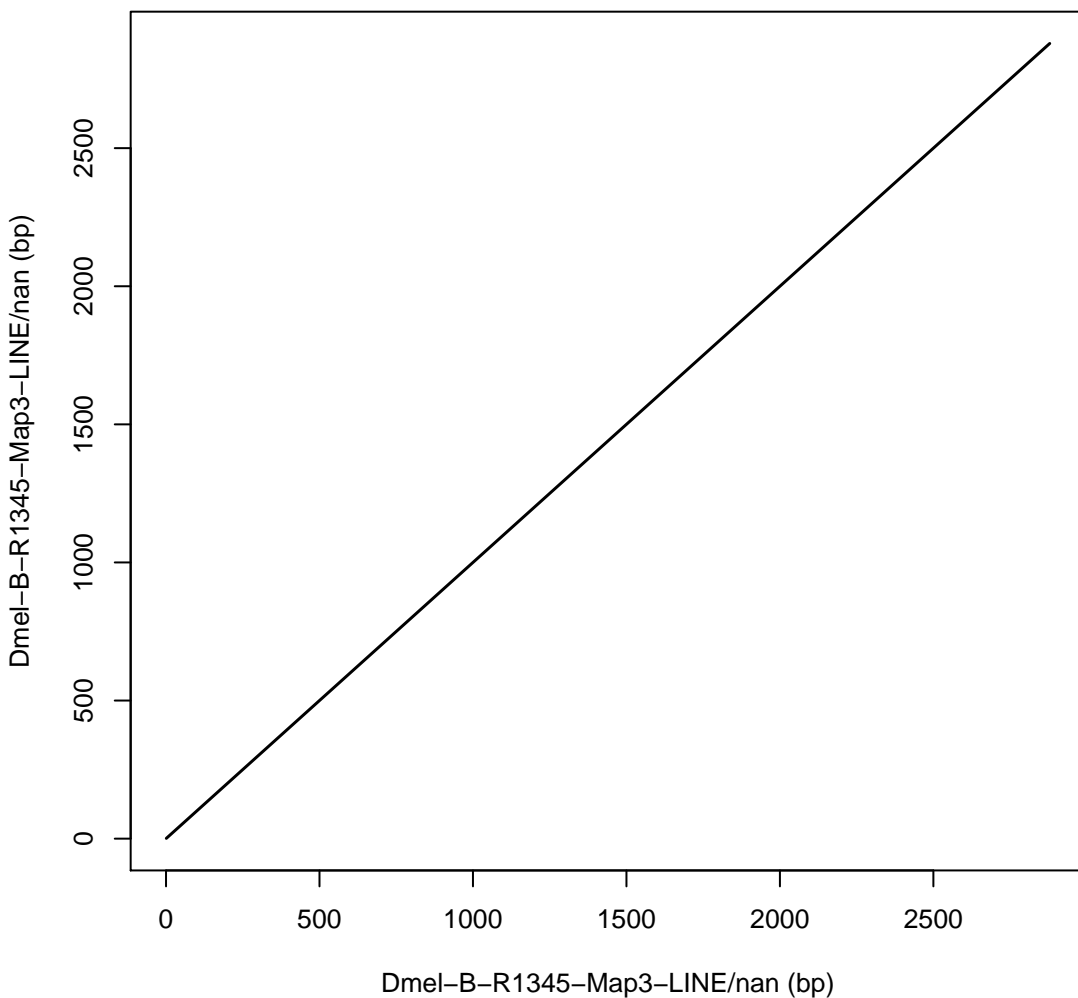
TE: Dmel-B-R1345-Map3-LINE/nan  
consensus size: 2879bp; fragments: 109; full length: 0 (>=2591.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

