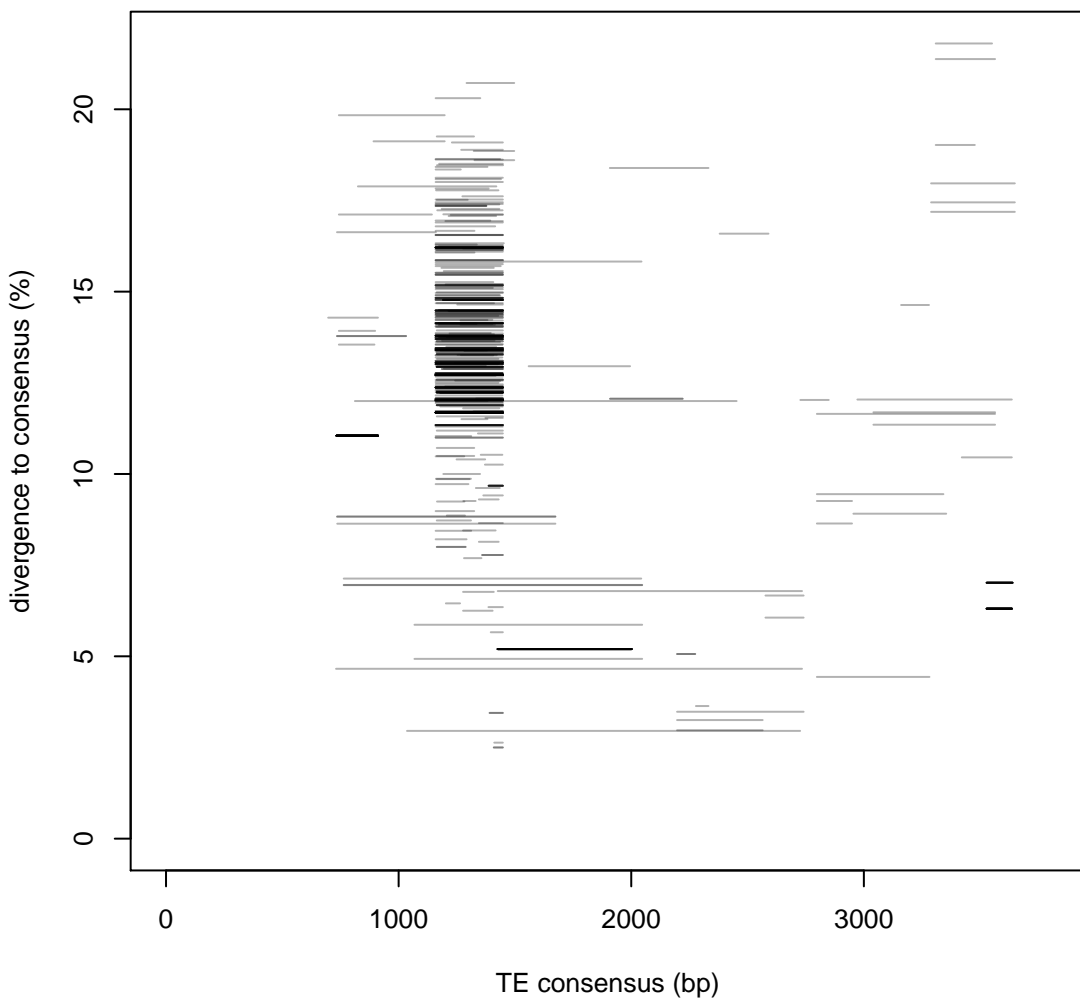
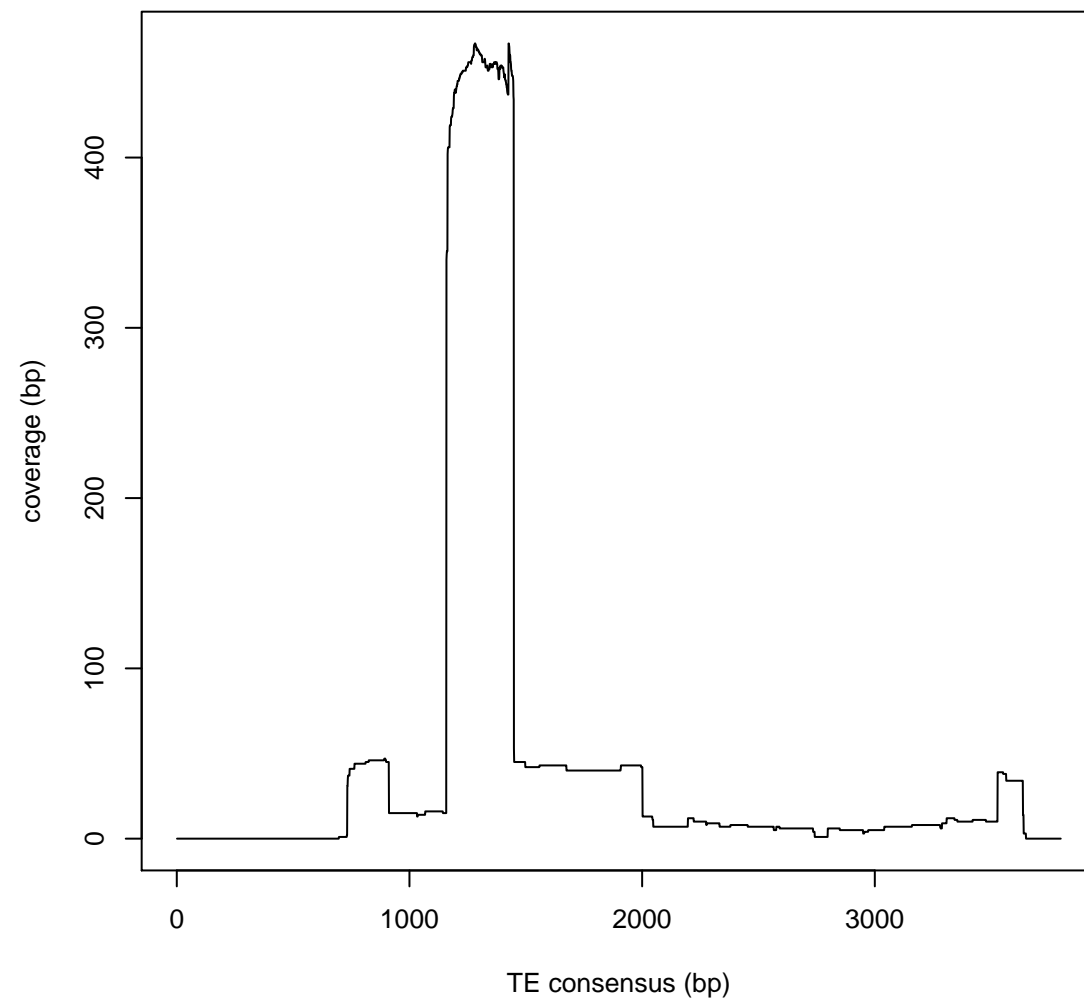


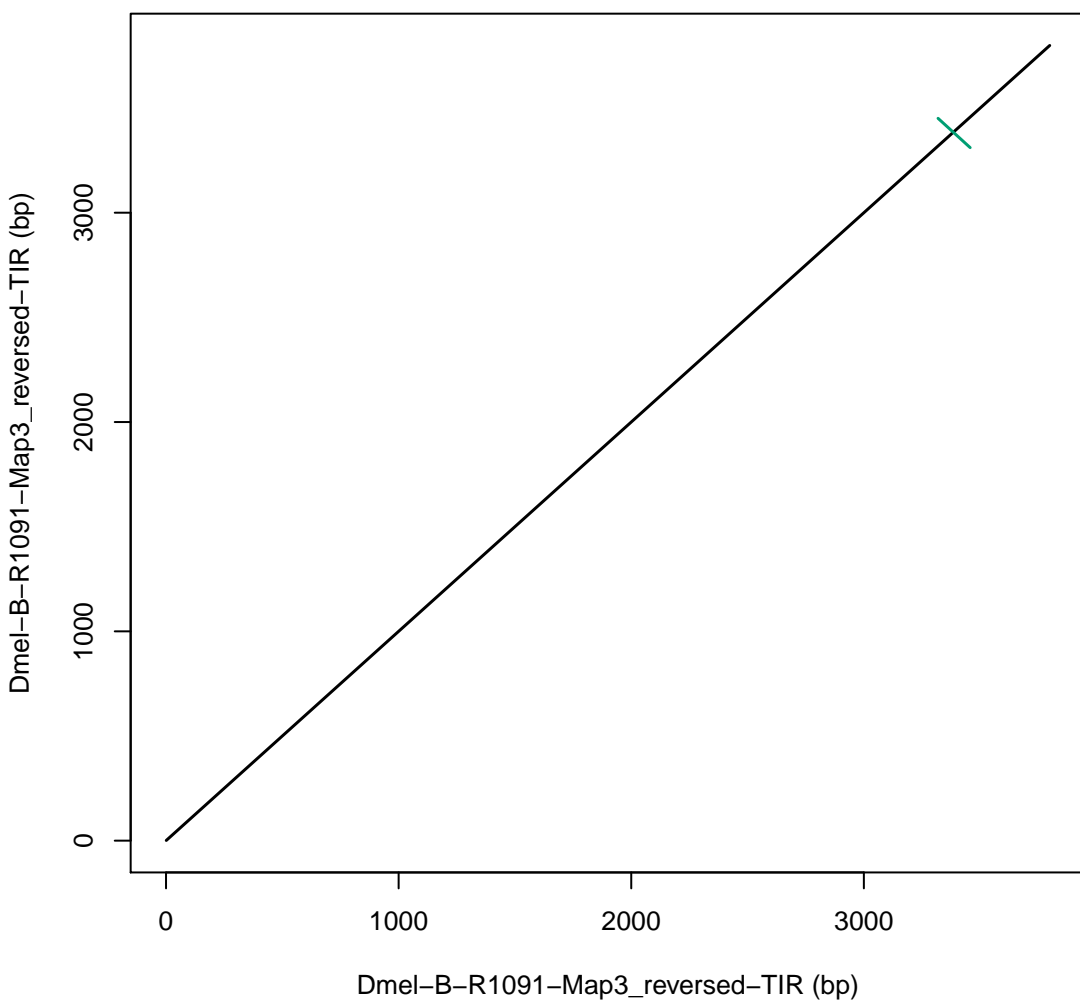
TE: Dmel-B-R1091-Map3_reversed-TIR
consensus size: 3799bp; fragments: 628; full length: 0 (>=3419.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

