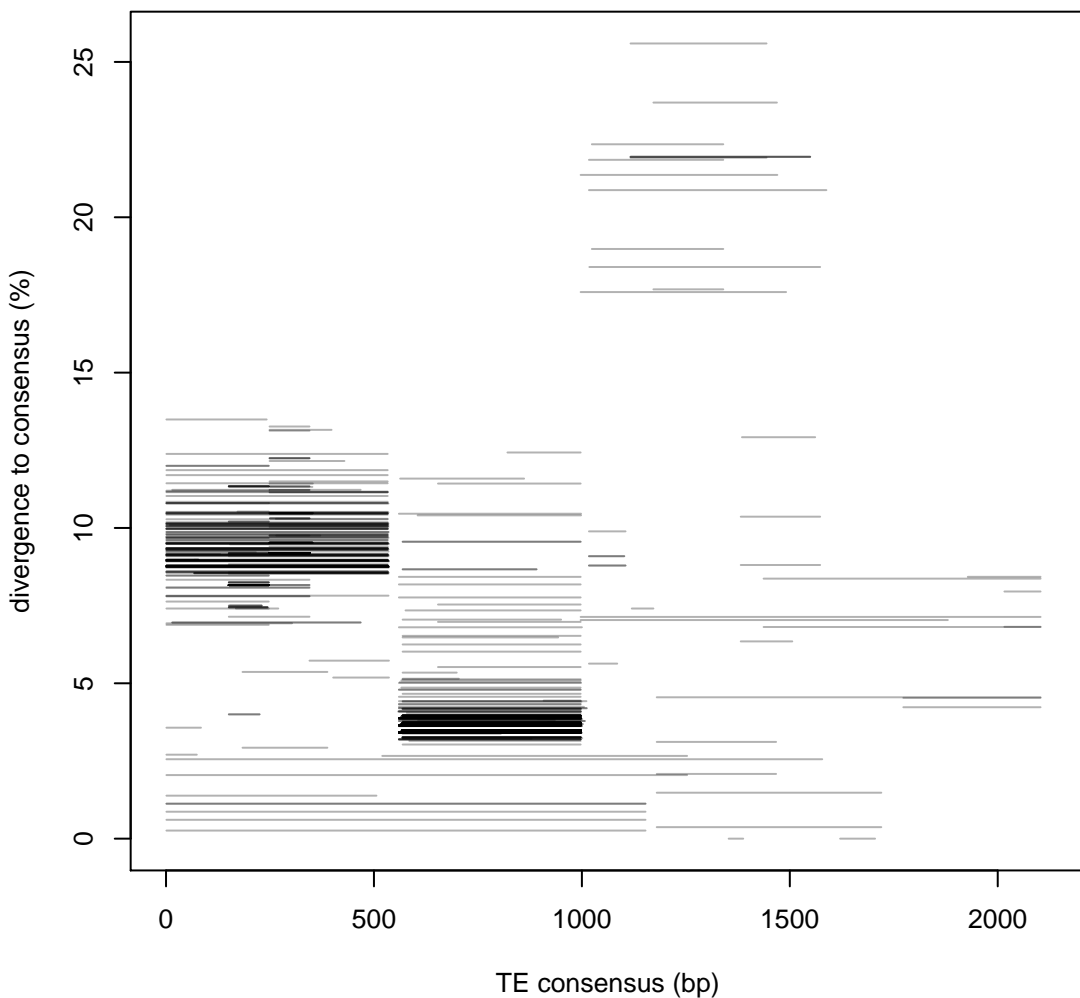
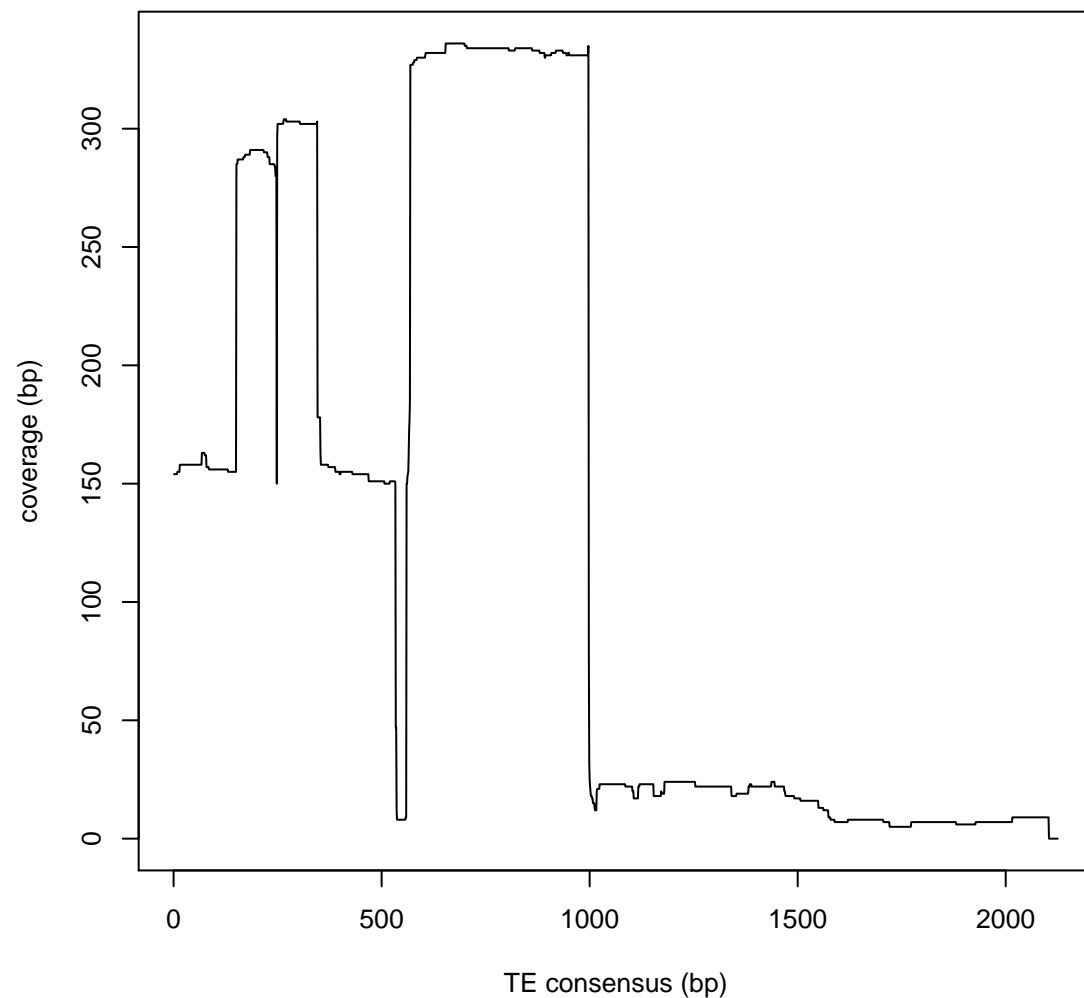


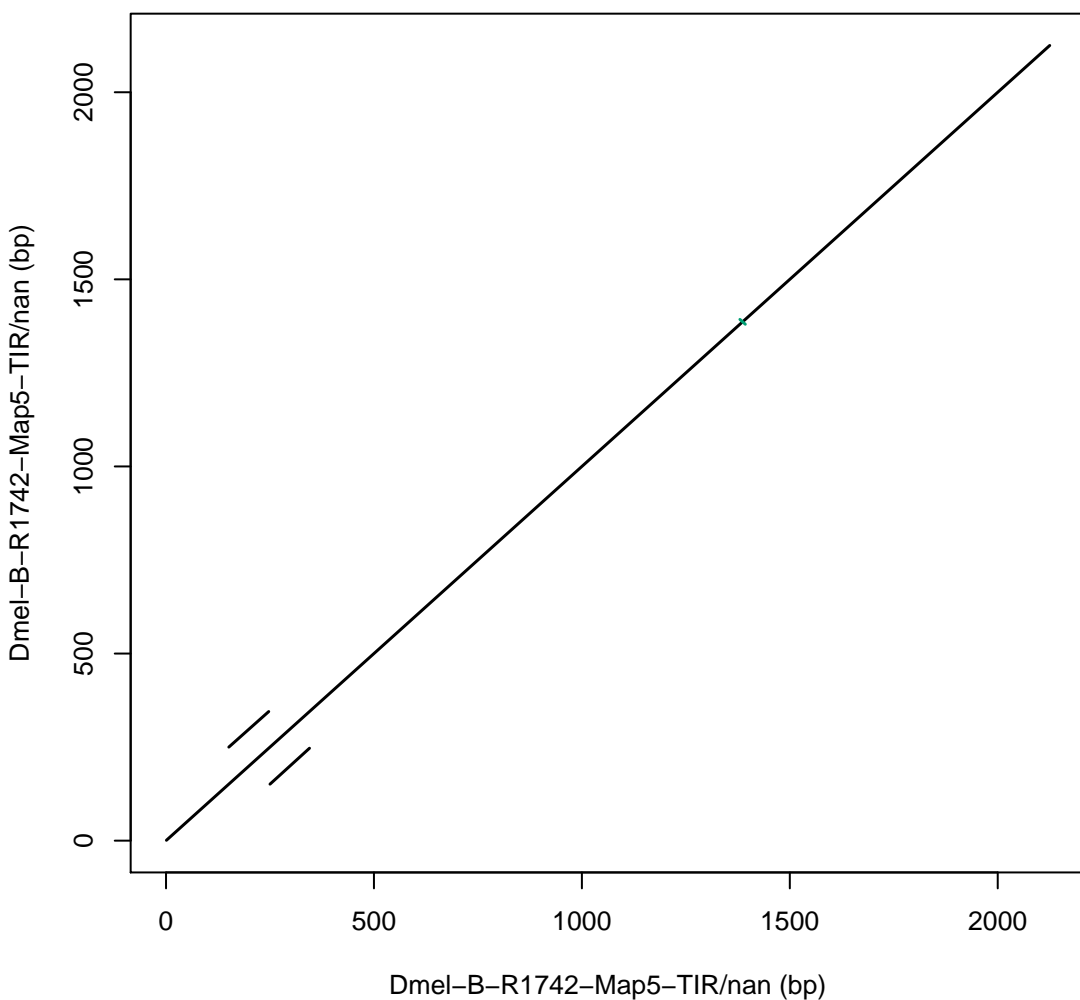
TE: Dmel-B-R1742-Map5-TIR/nan
consensus size: 2125bp; fragments: 830; full length: 0 (>=1912.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

