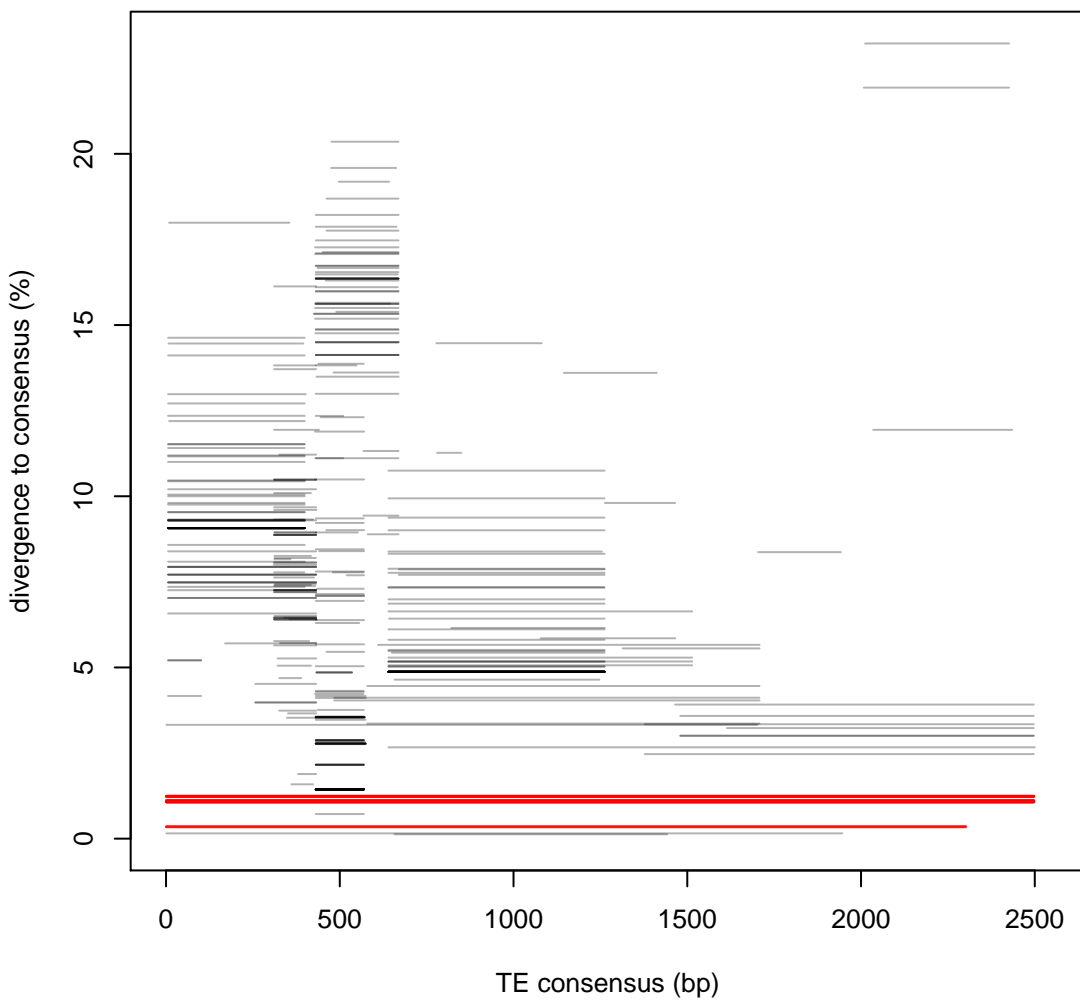
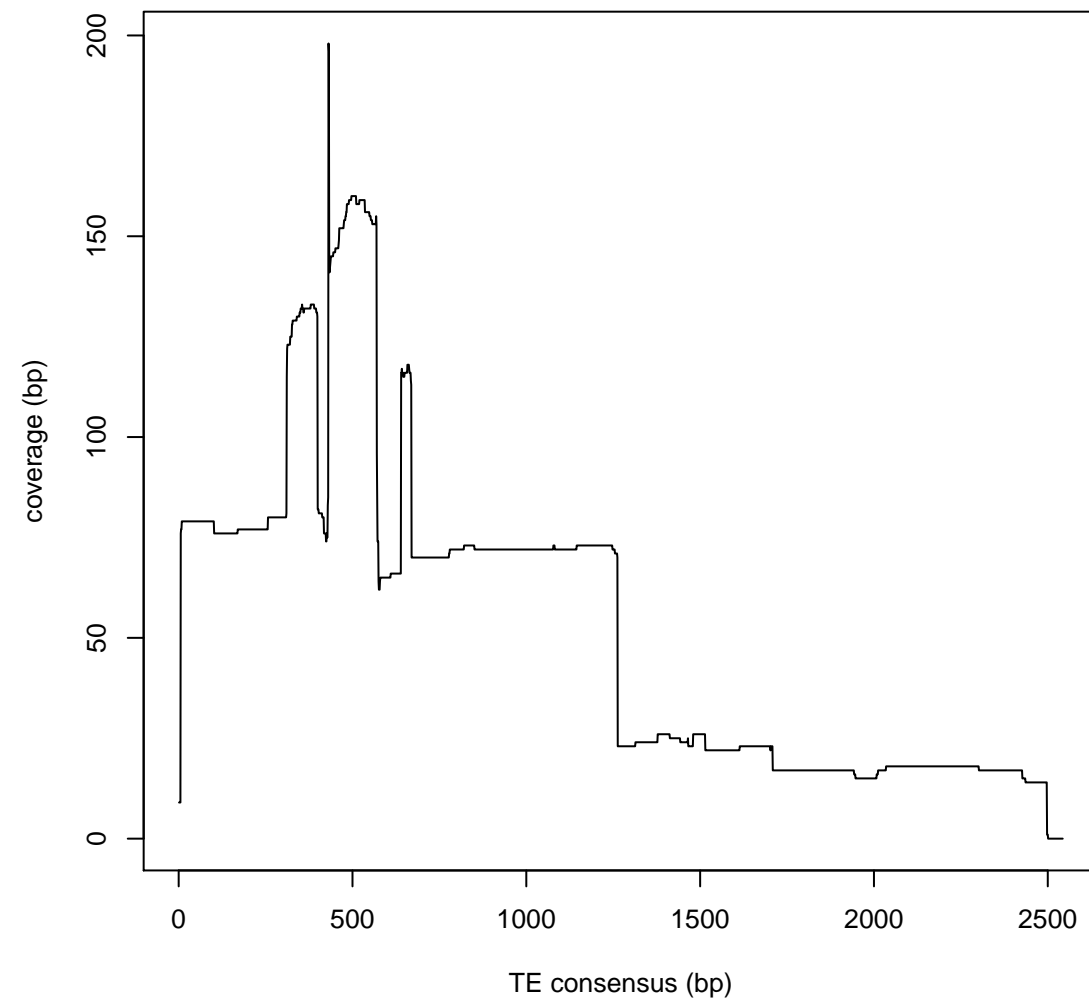


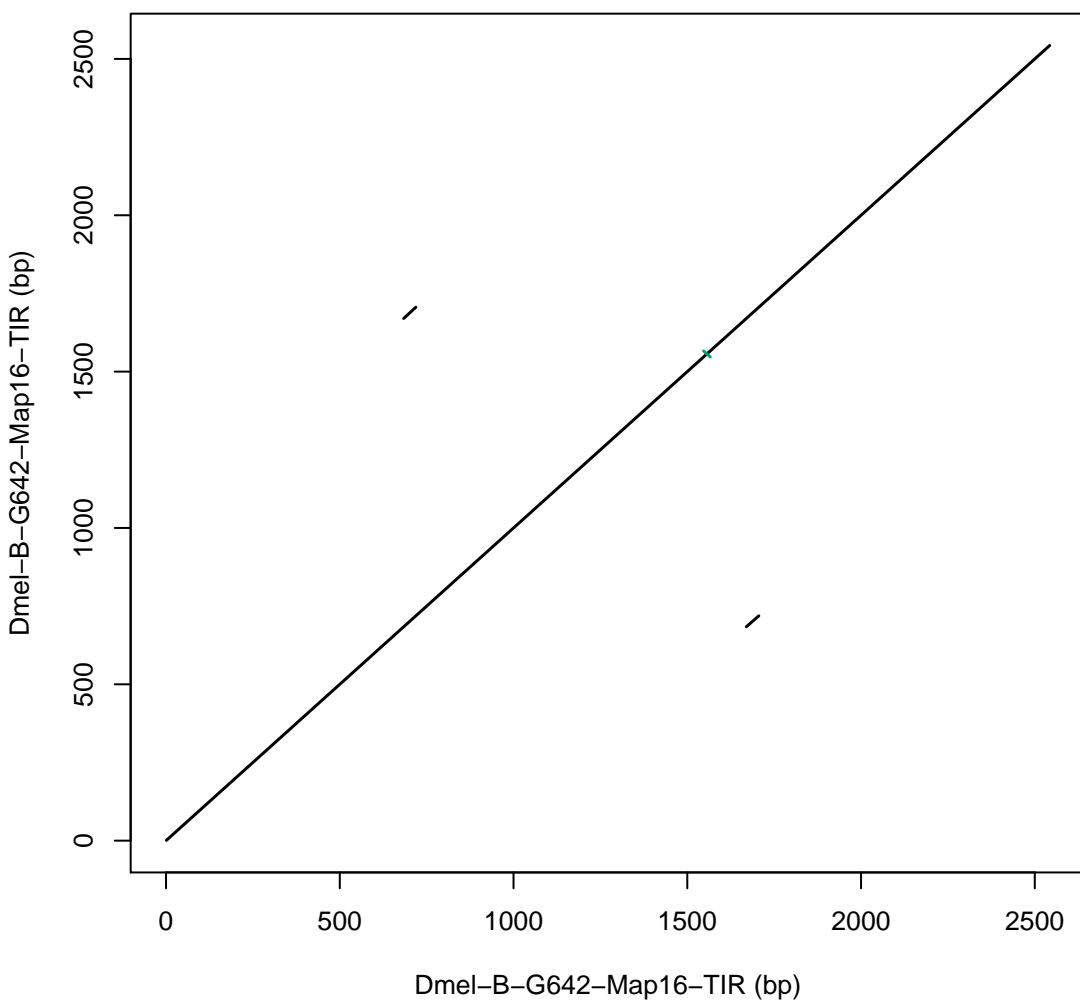
TE: Dmel-B-G642-Map16-TIR
consensus size: 2543bp; fragments: 370; full length: 7 (>=2288.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

