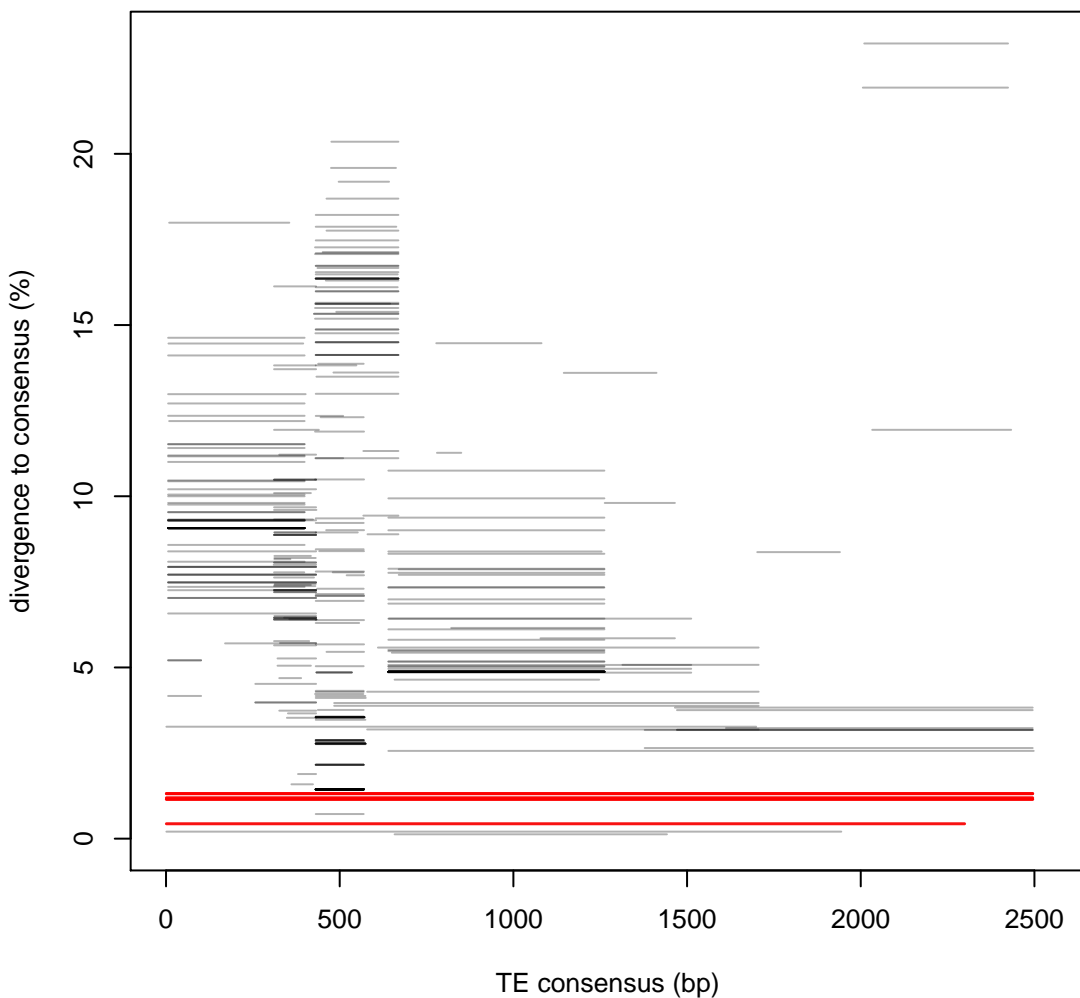
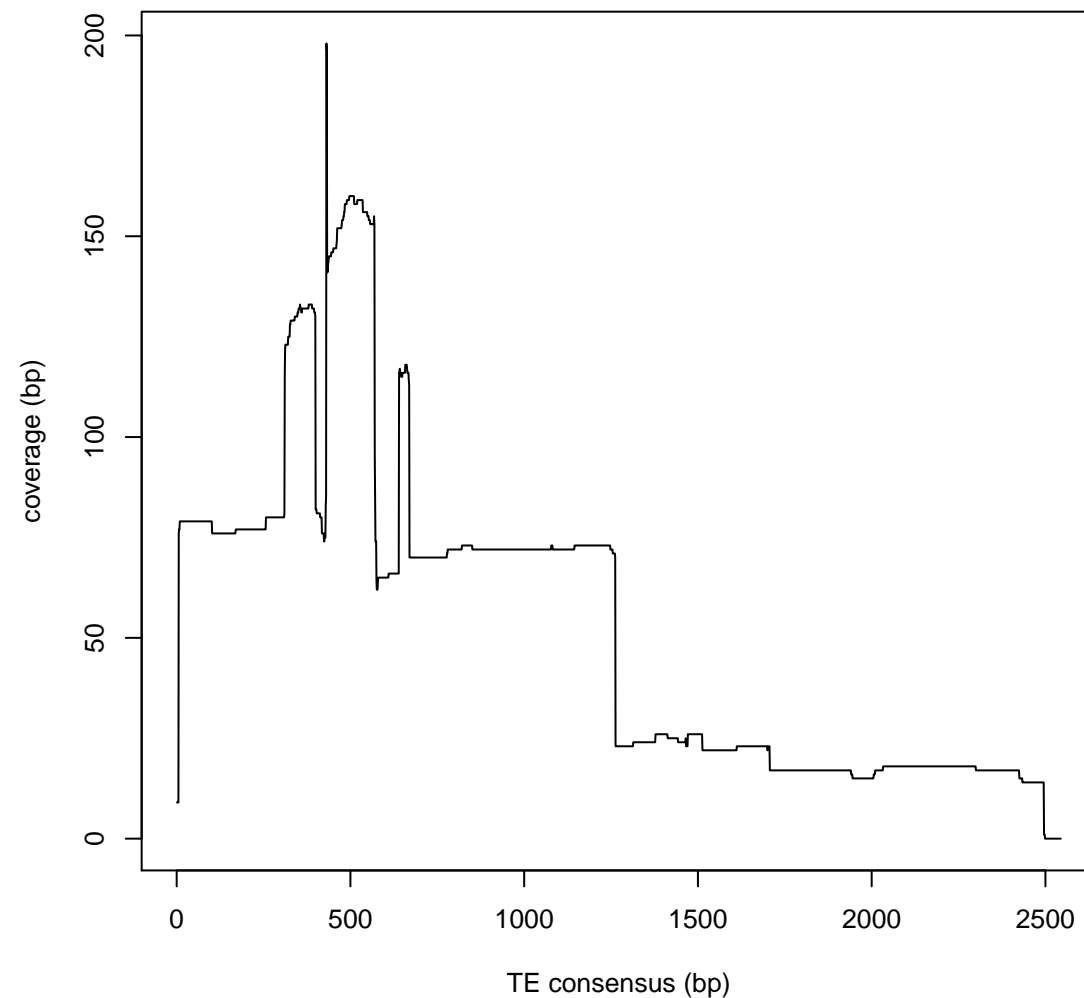


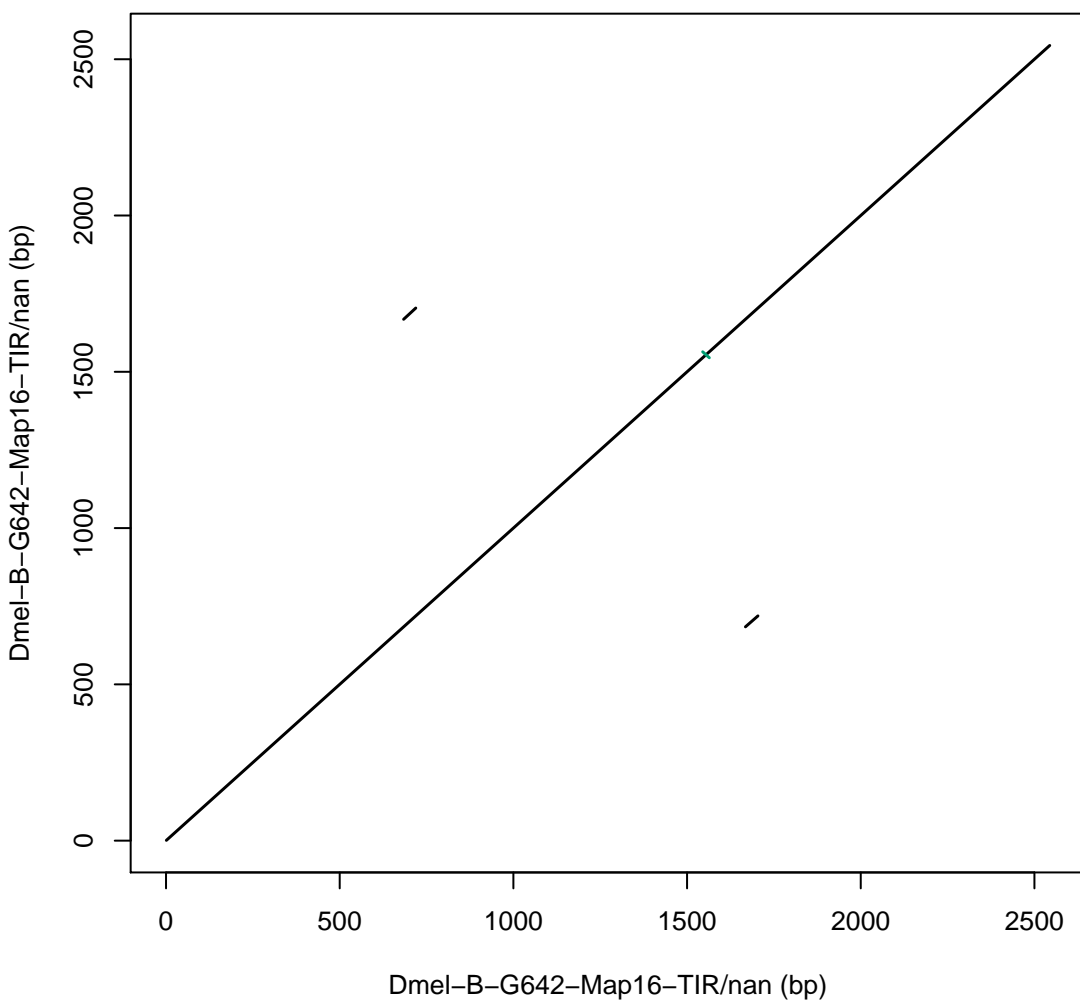
TE: Dmel-B-G642-Map16-TIR/nan
consensus size: 2544bp; fragments: 370; full length: 7 (>=2289.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

