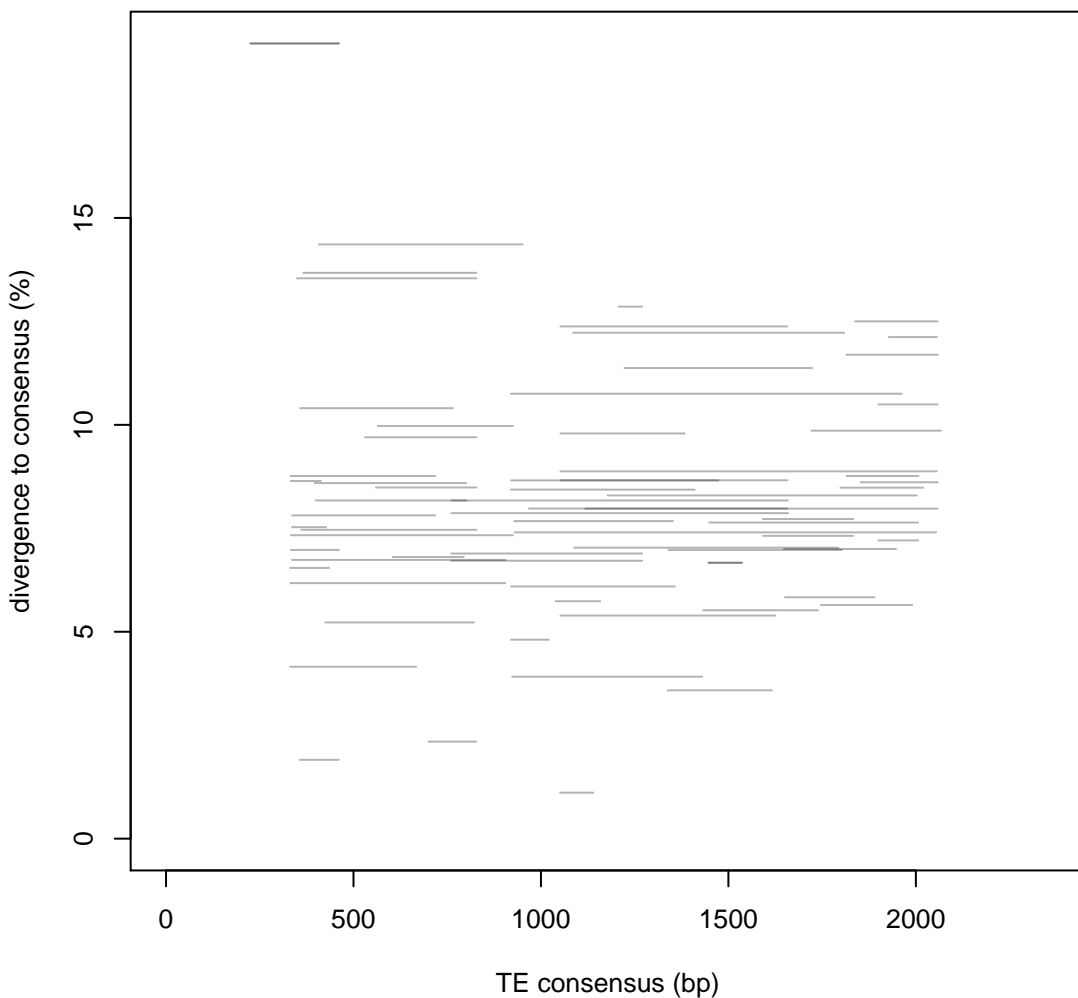
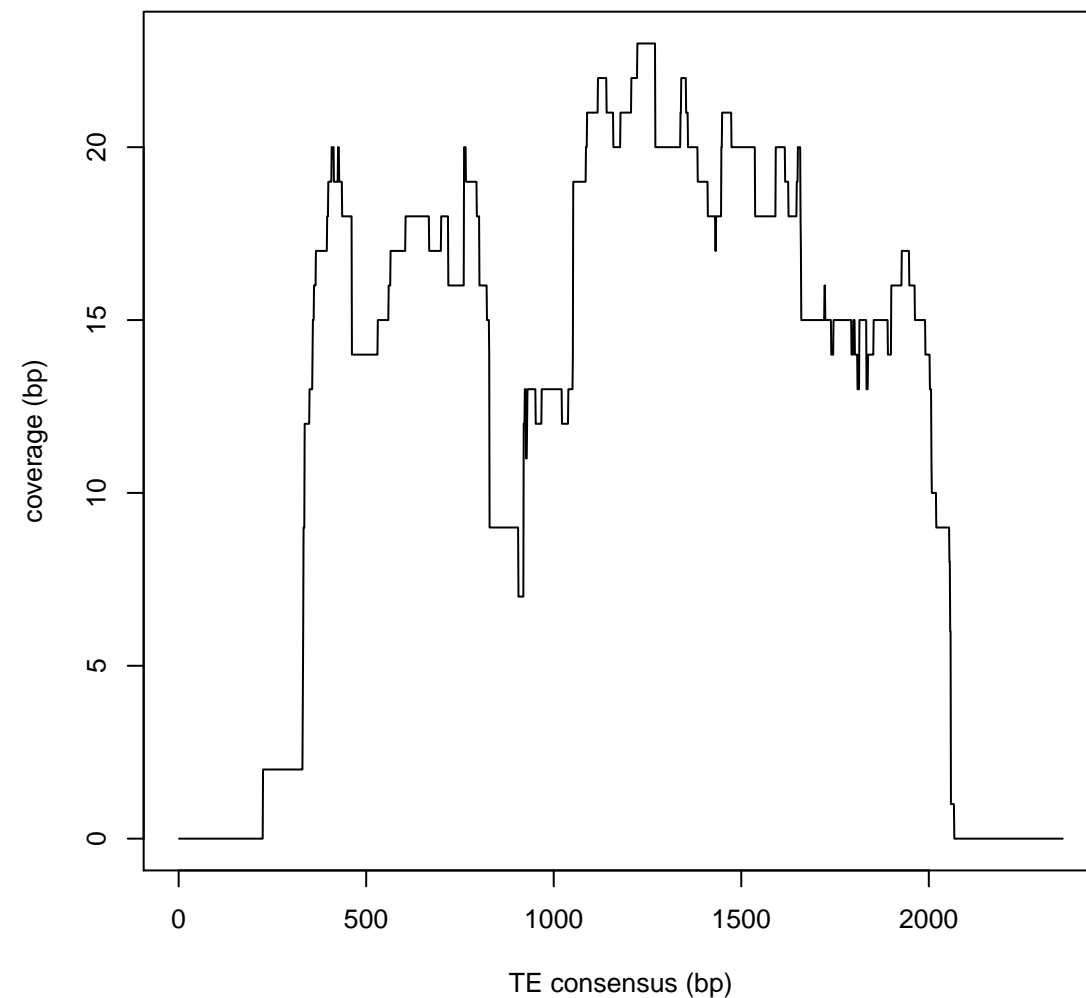


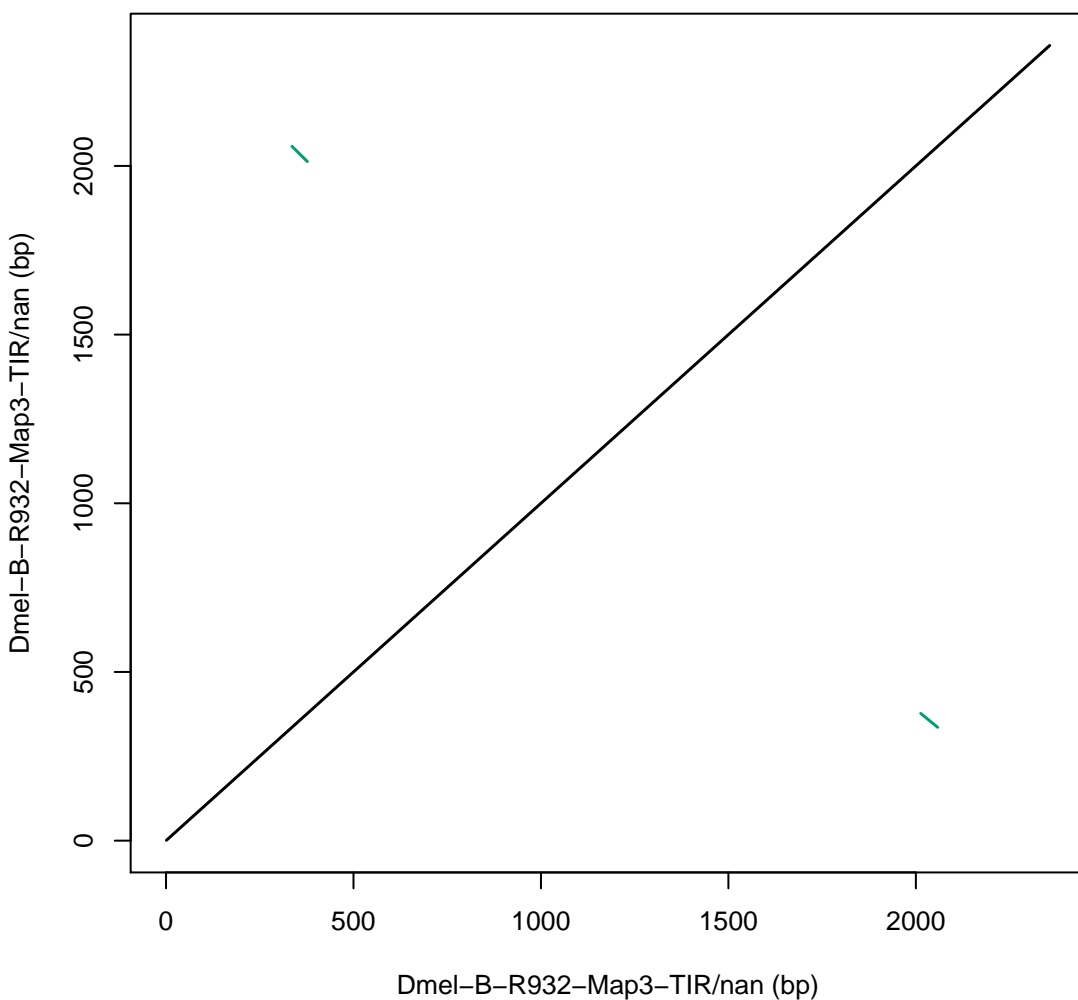
TE: Dmel-B-R932-Map3-TIR/nan
consensus size: 2357bp; fragments: 72; full length: 0 (>=2121.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

