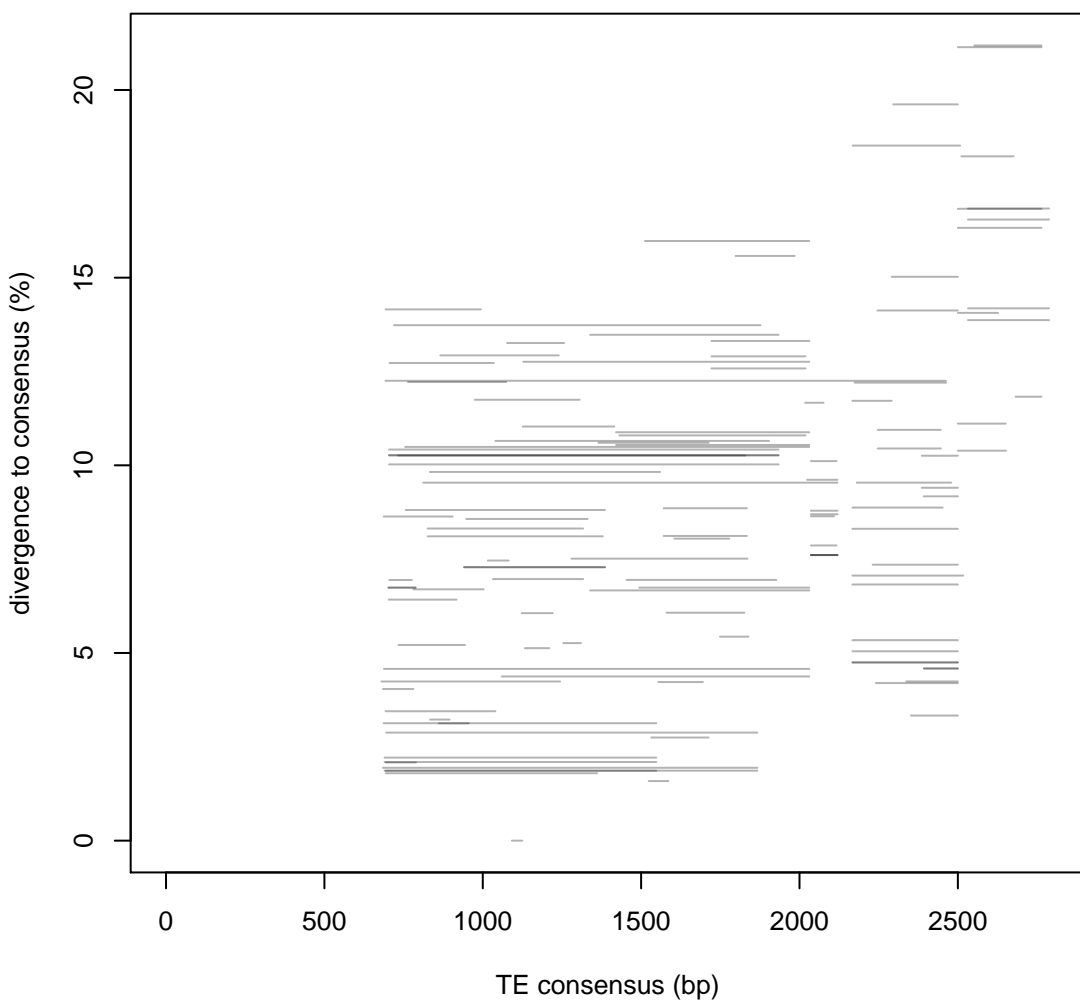
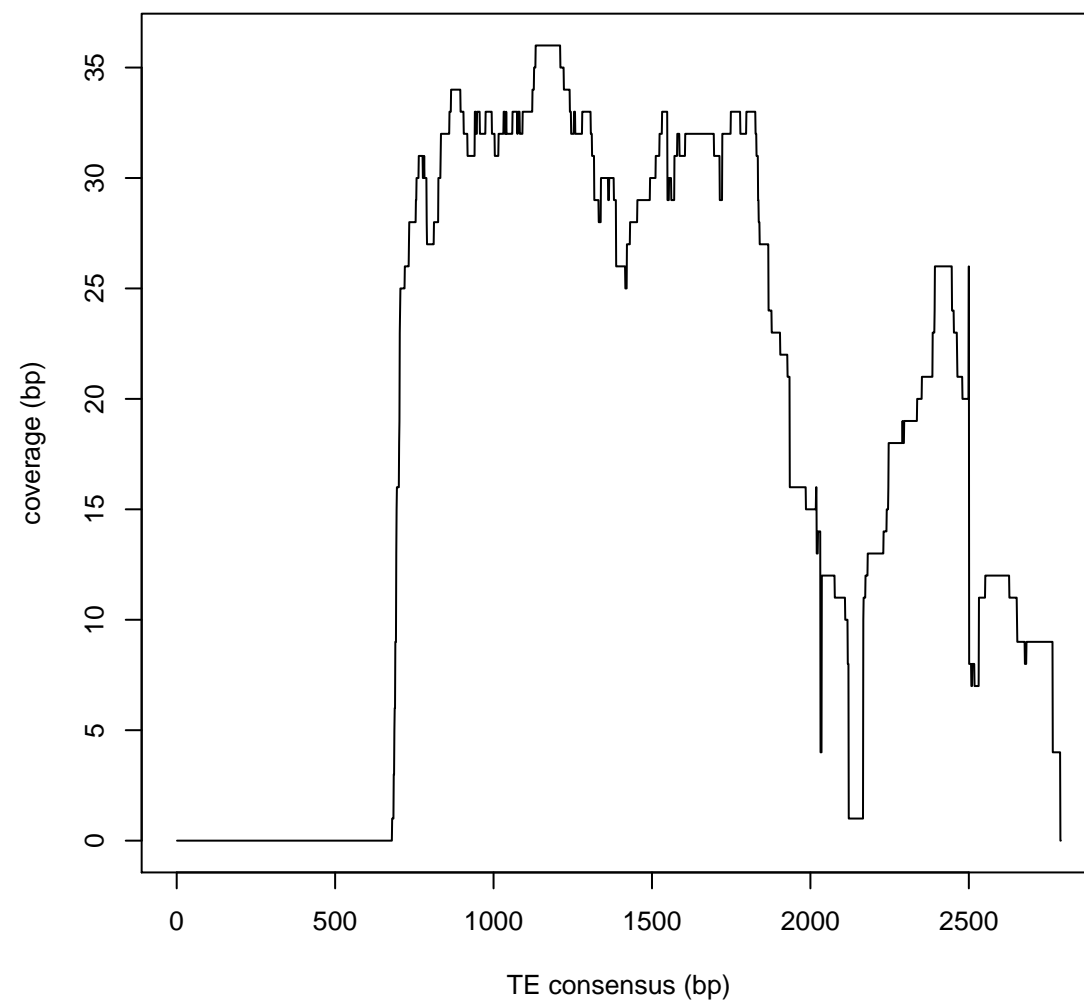


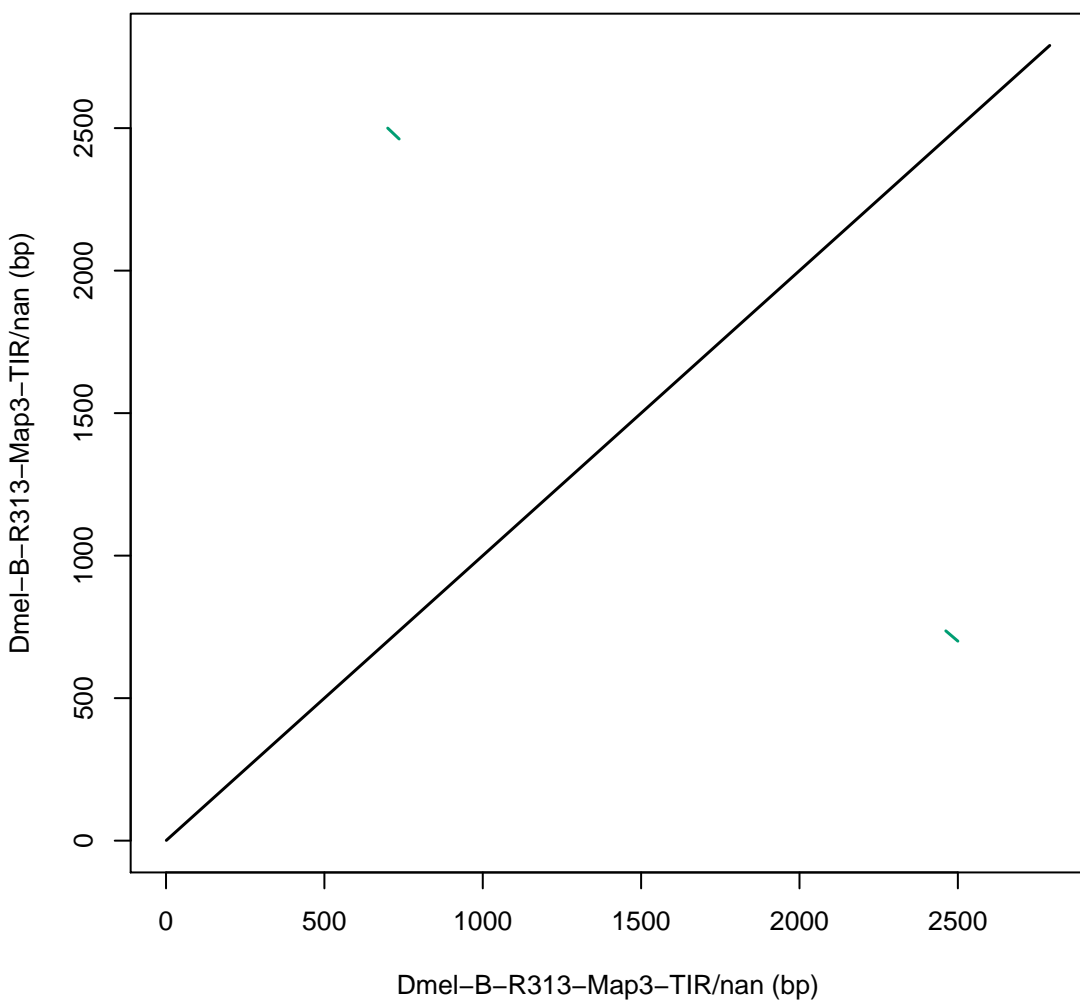
TE: Dmel-B-R313-Map3-TIR/nan
consensus size: 2790bp; fragments: 125; full length: 0 (>=2511bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

