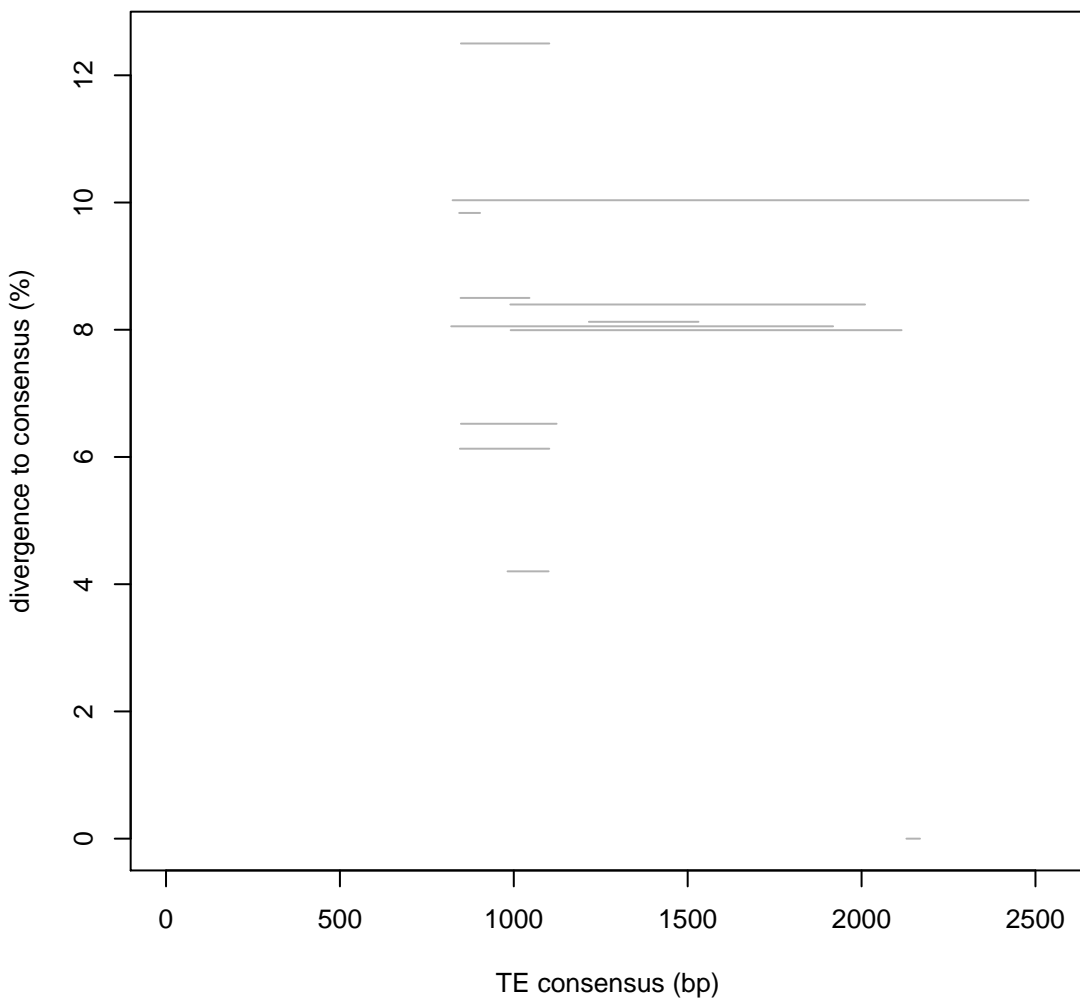
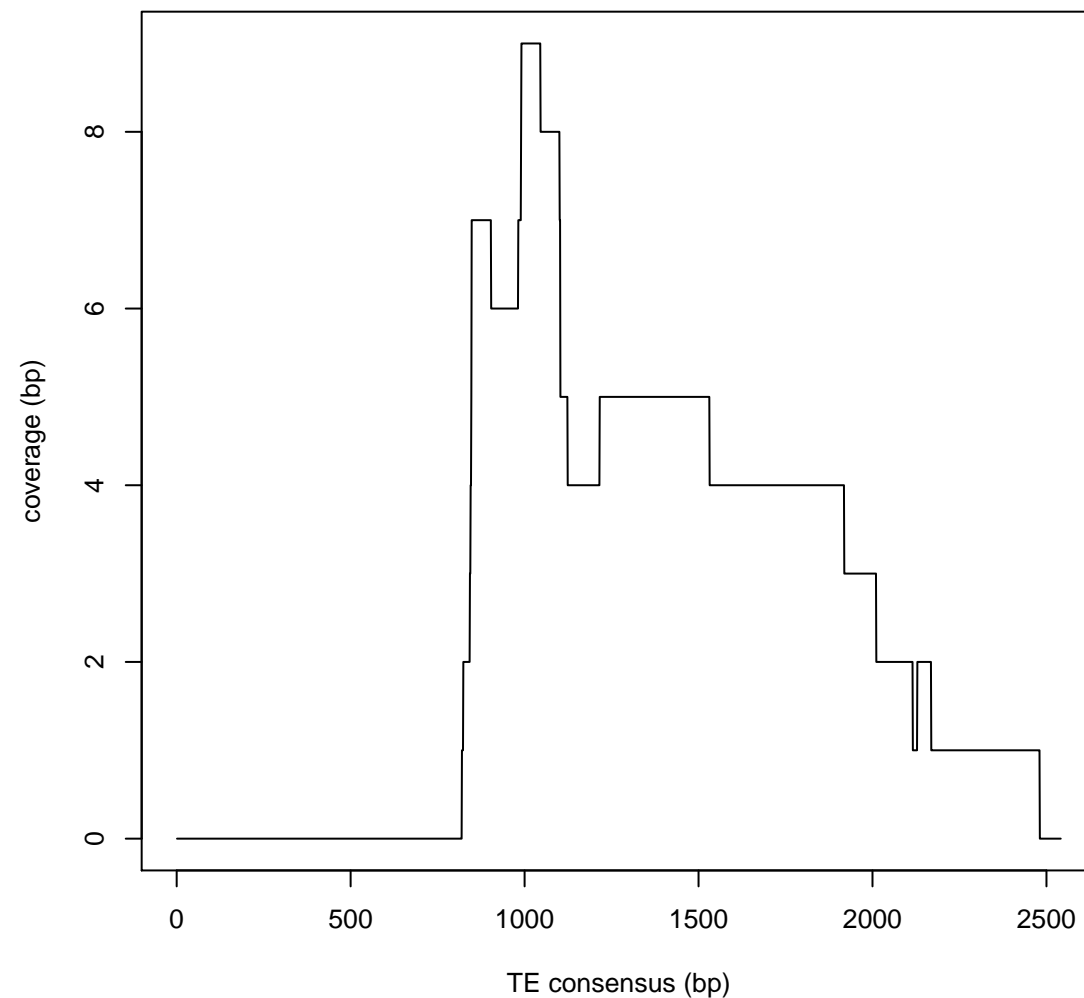


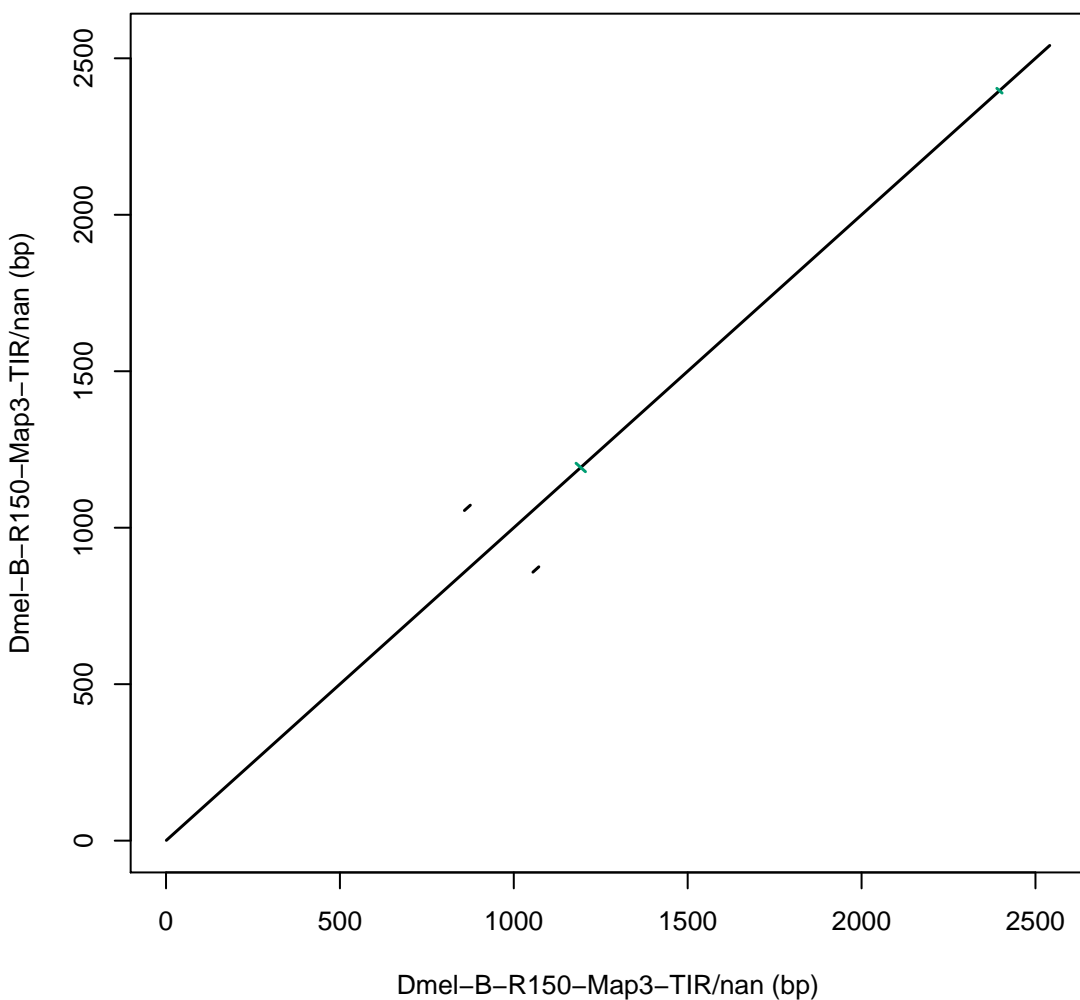
TE: Dmel-B-R150-Map3-TIR/nan
consensus size: 2541bp; fragments: 12; full length: 0 (>=2286.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

