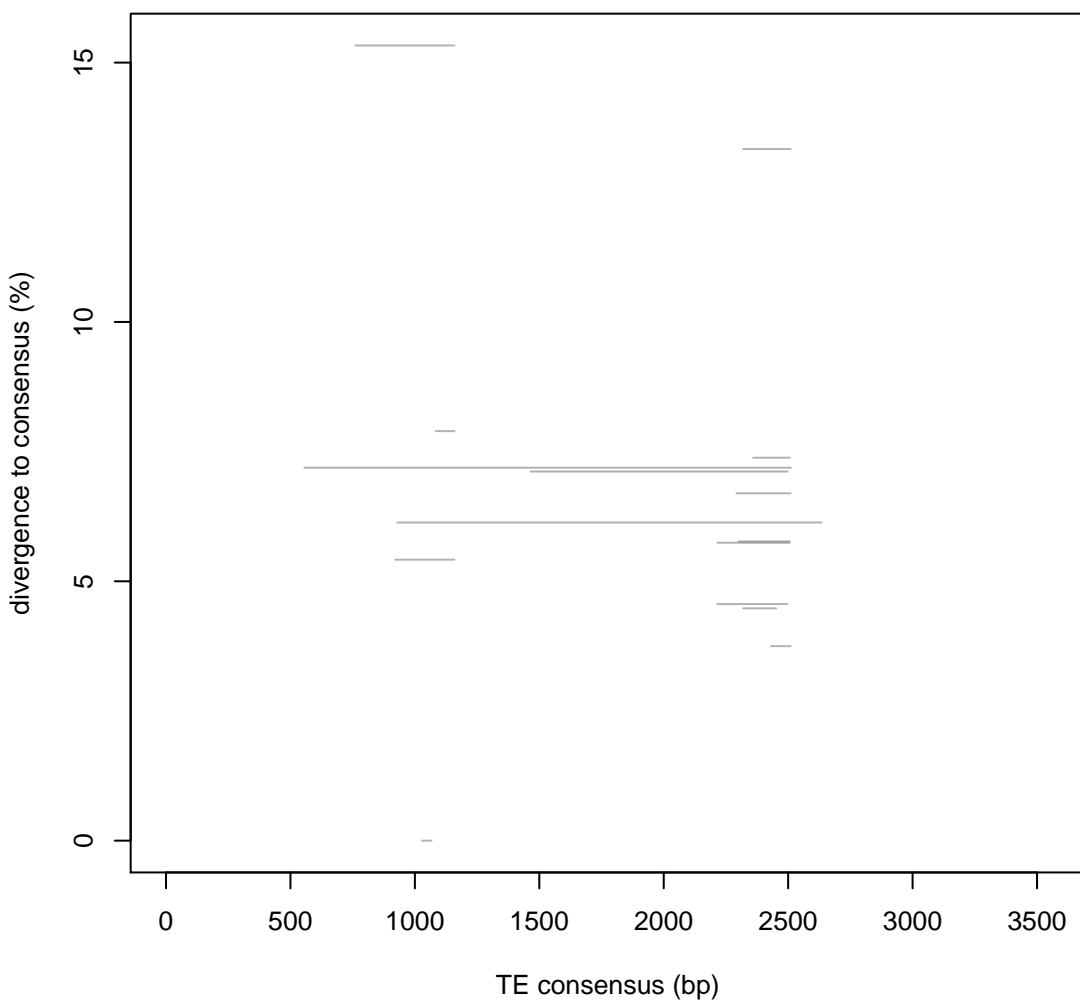
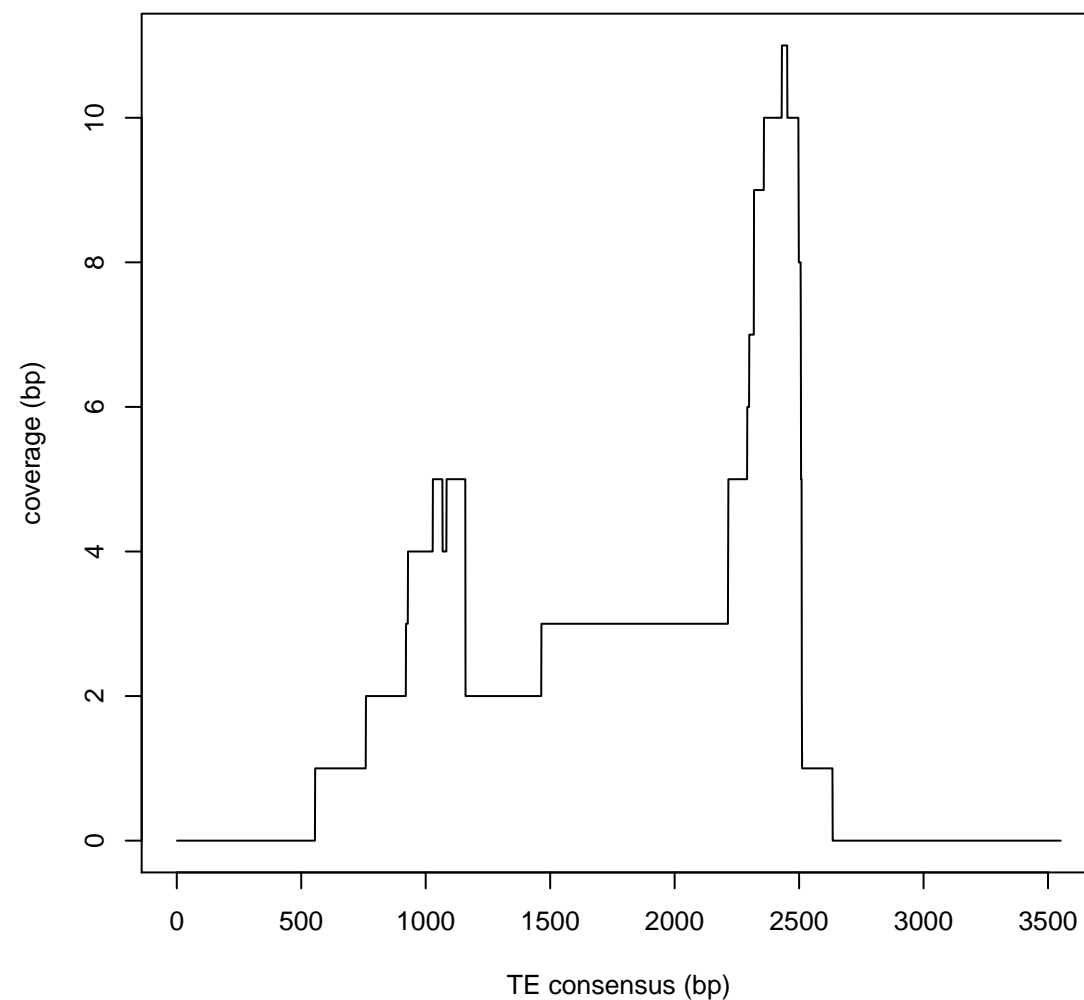


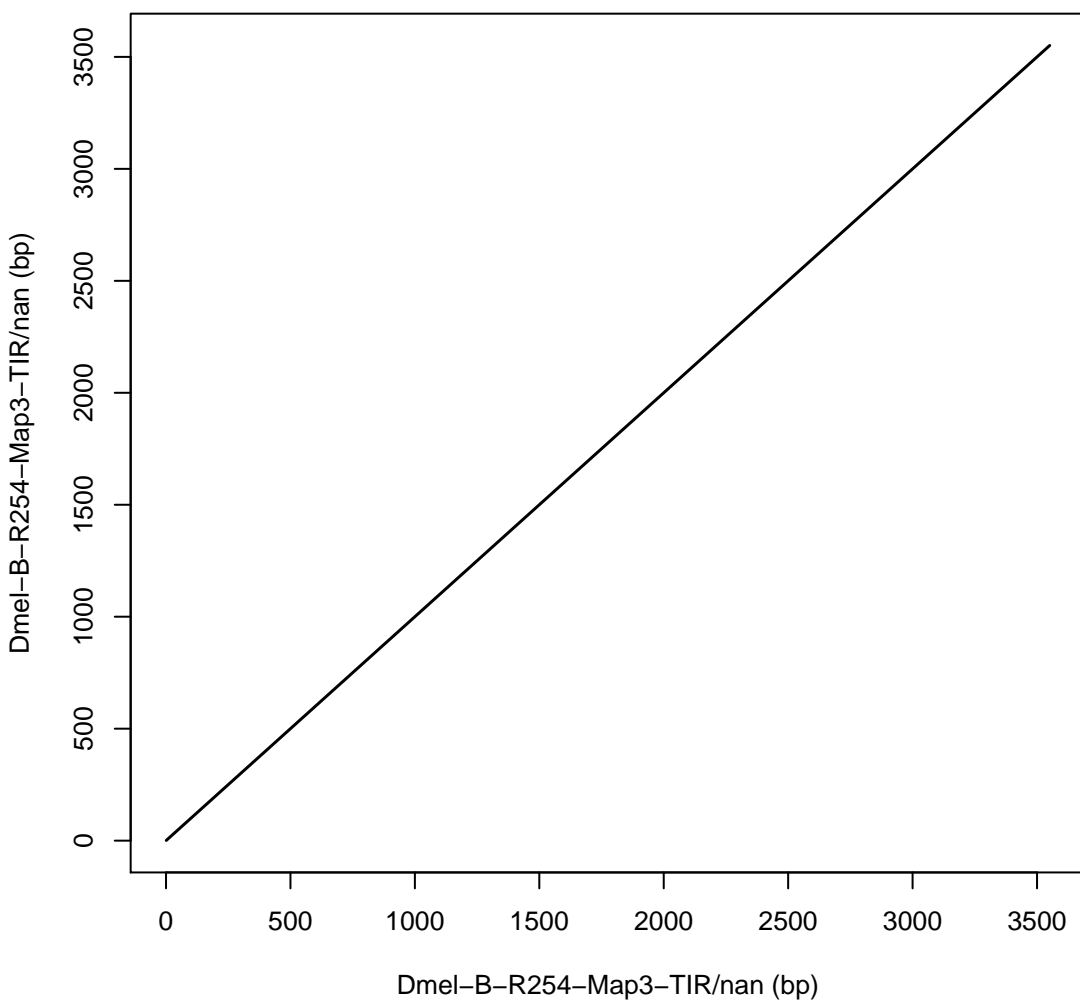
TE: Dmel-B-R254-Map3-TIR/nan
consensus size: 3551bp; fragments: 15; full length: 0 (>=3195.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

