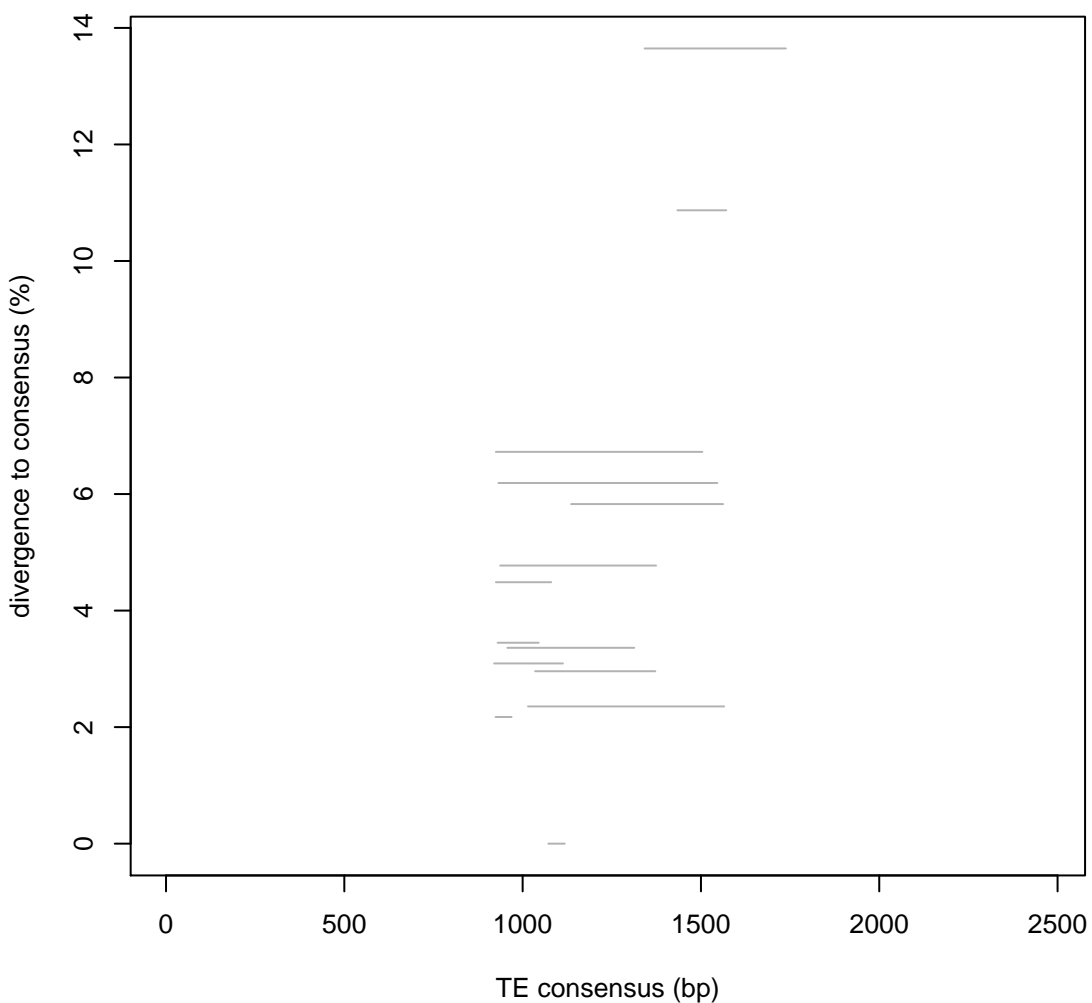
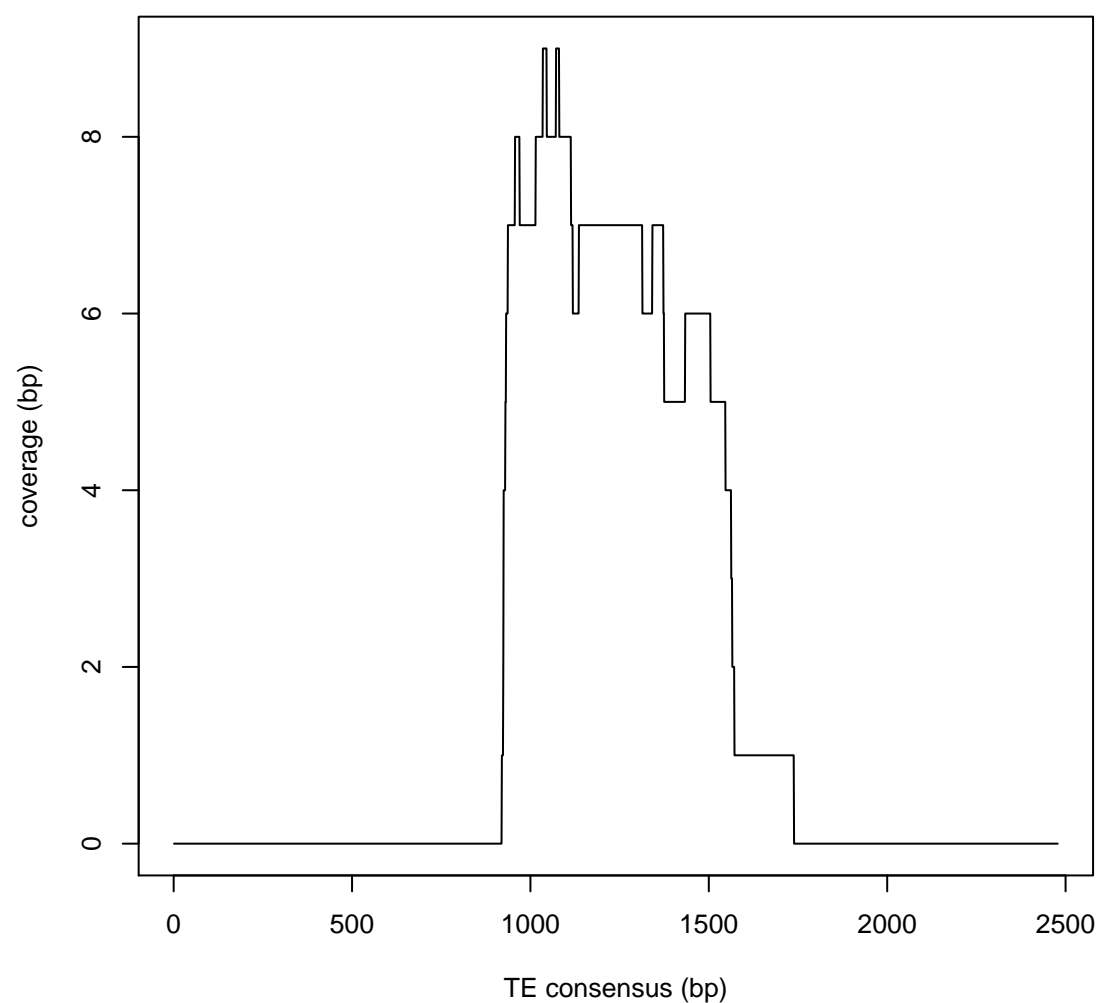


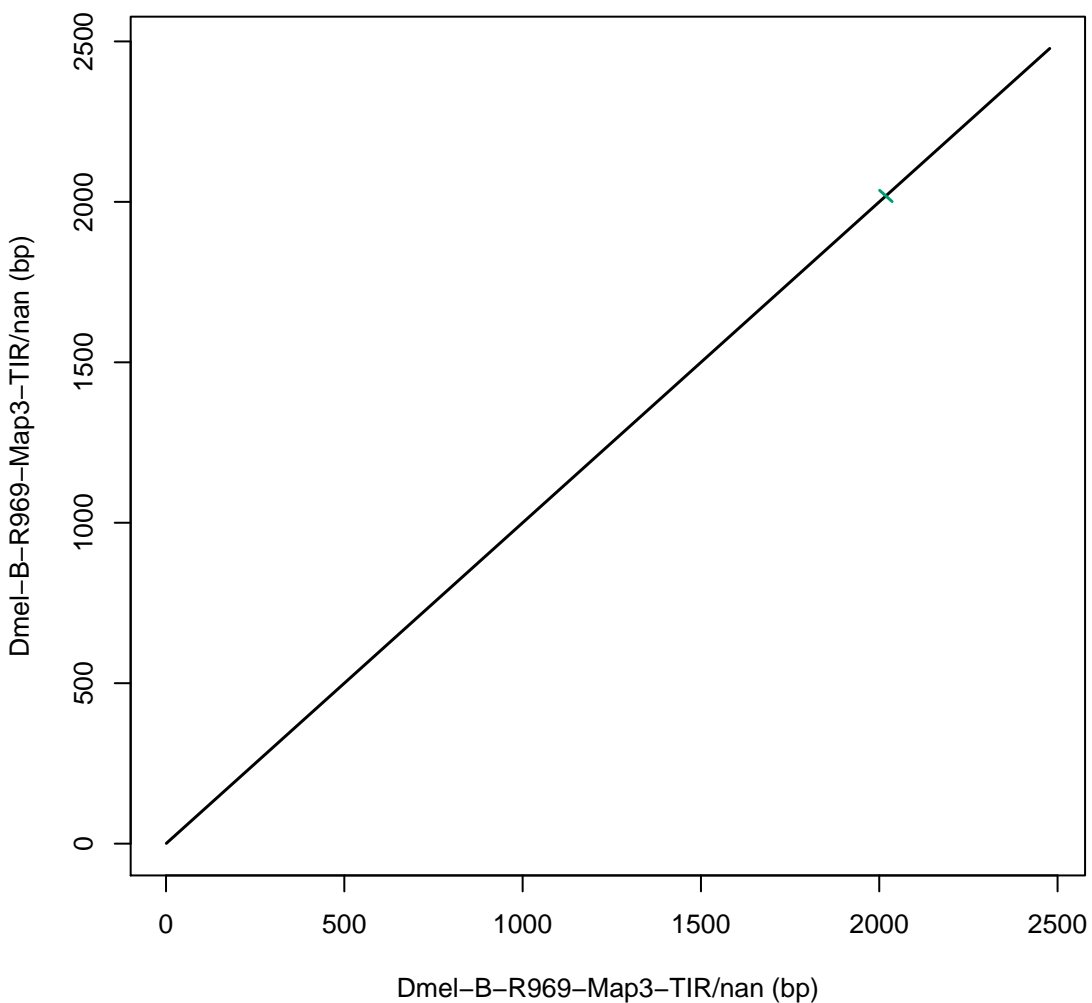
TE: Dmel-B-R969-Map3-TIR/nan
consensus size: 2478bp; fragments: 14; full length: 0 (>=2230.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

