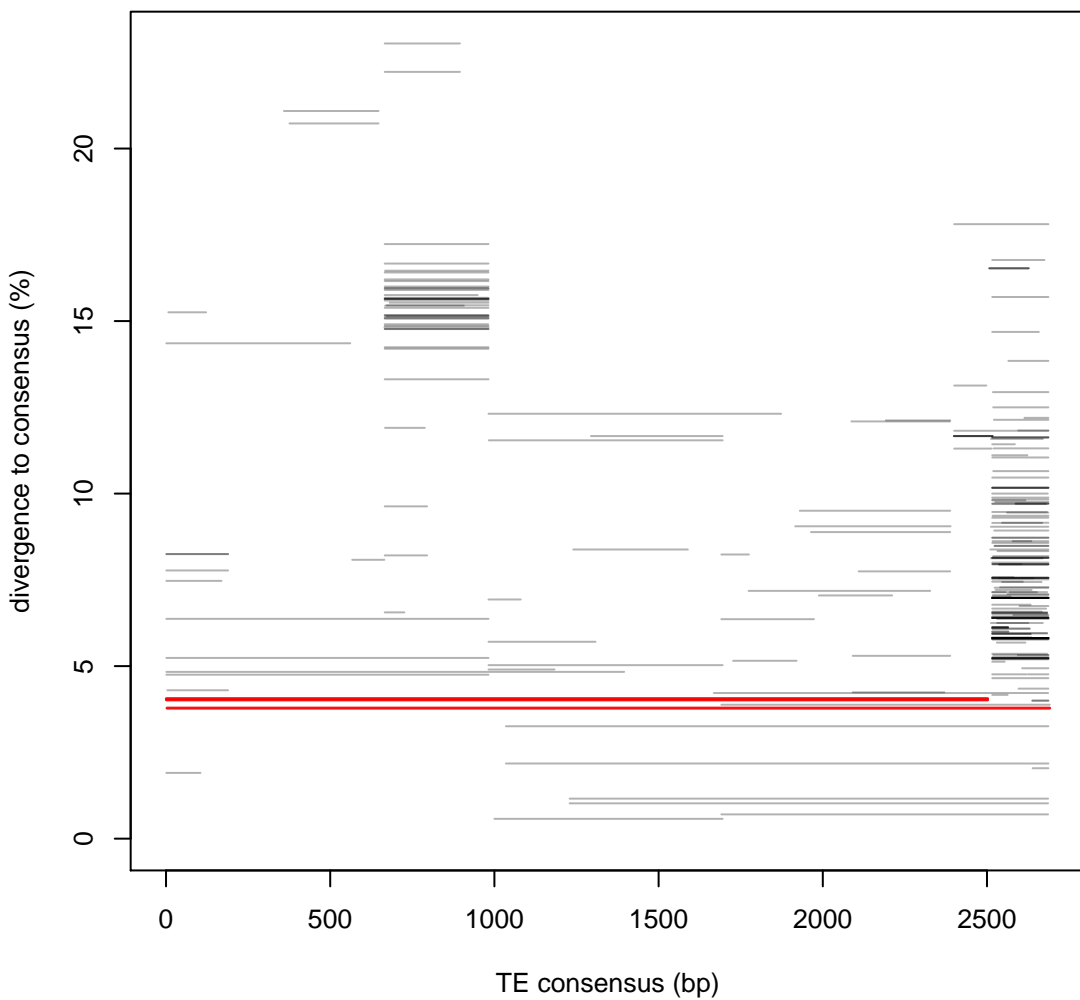
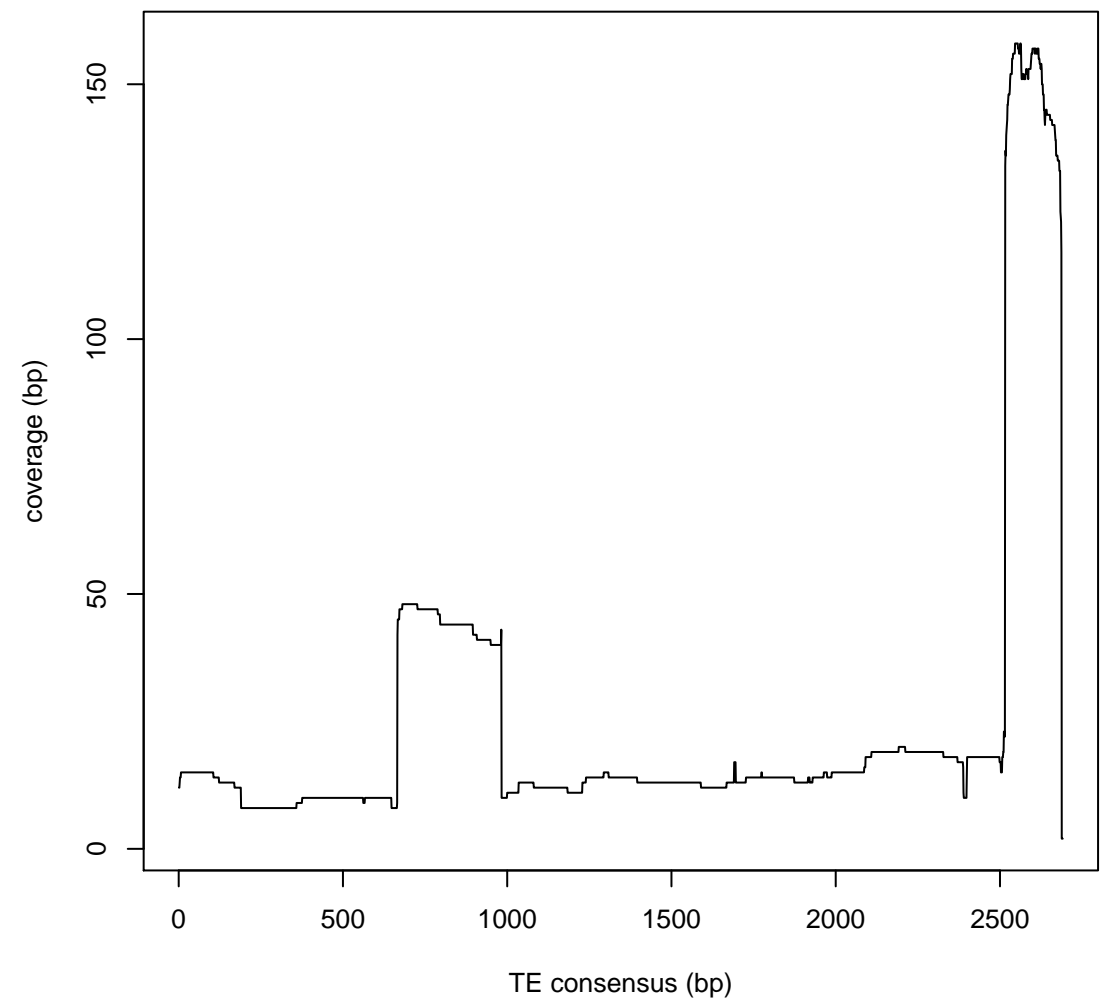


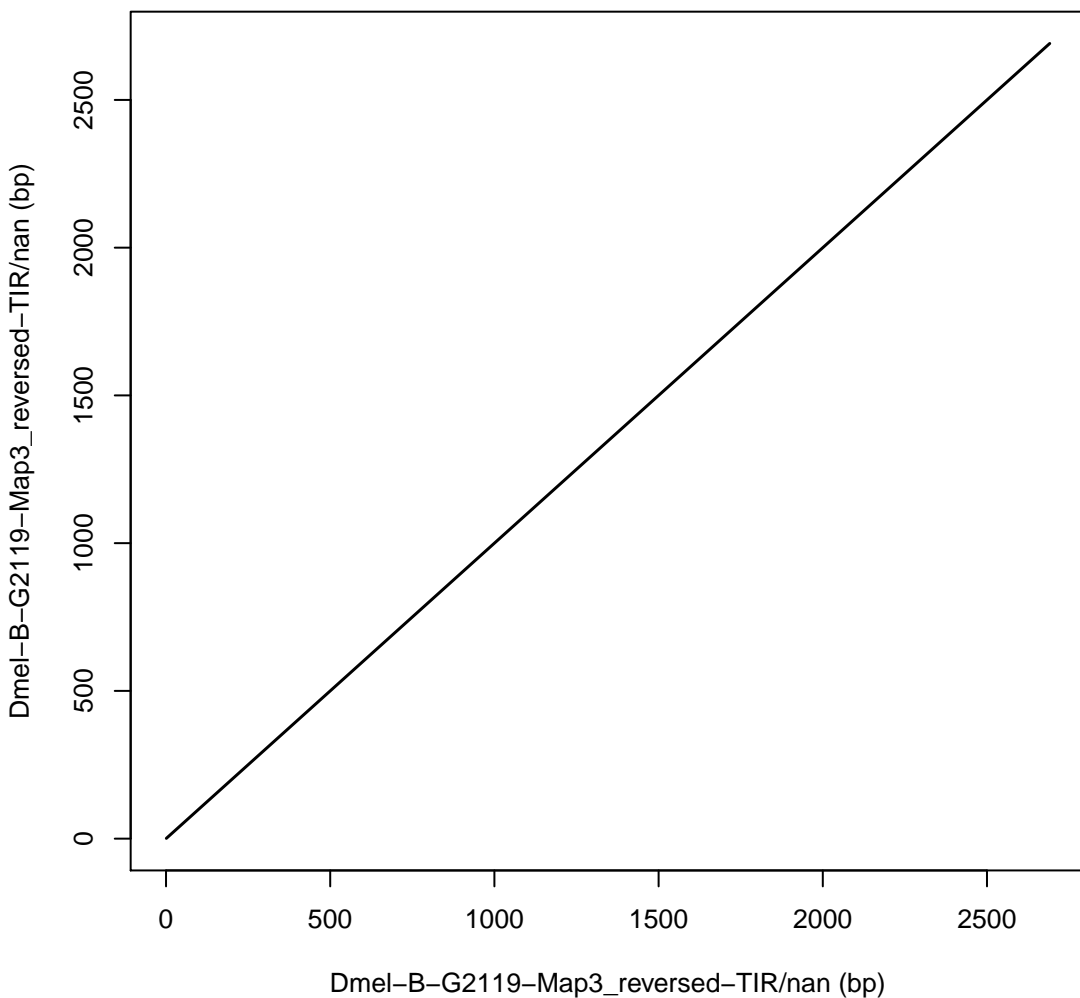
TE: Dmel-B-G2119-Map3_reversed-TIR/nan
consensus size: 2691bp; fragments: 262; full length: 3 (>=2421.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

