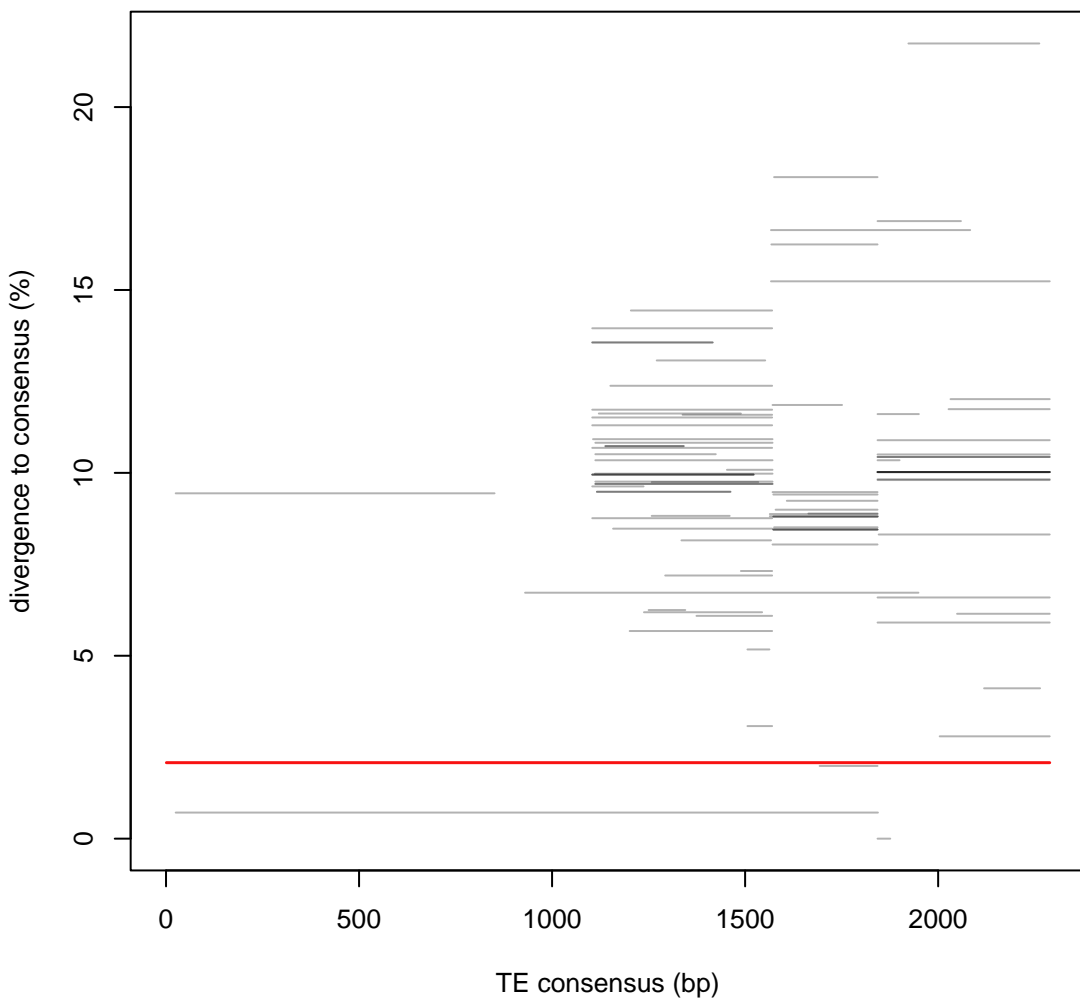
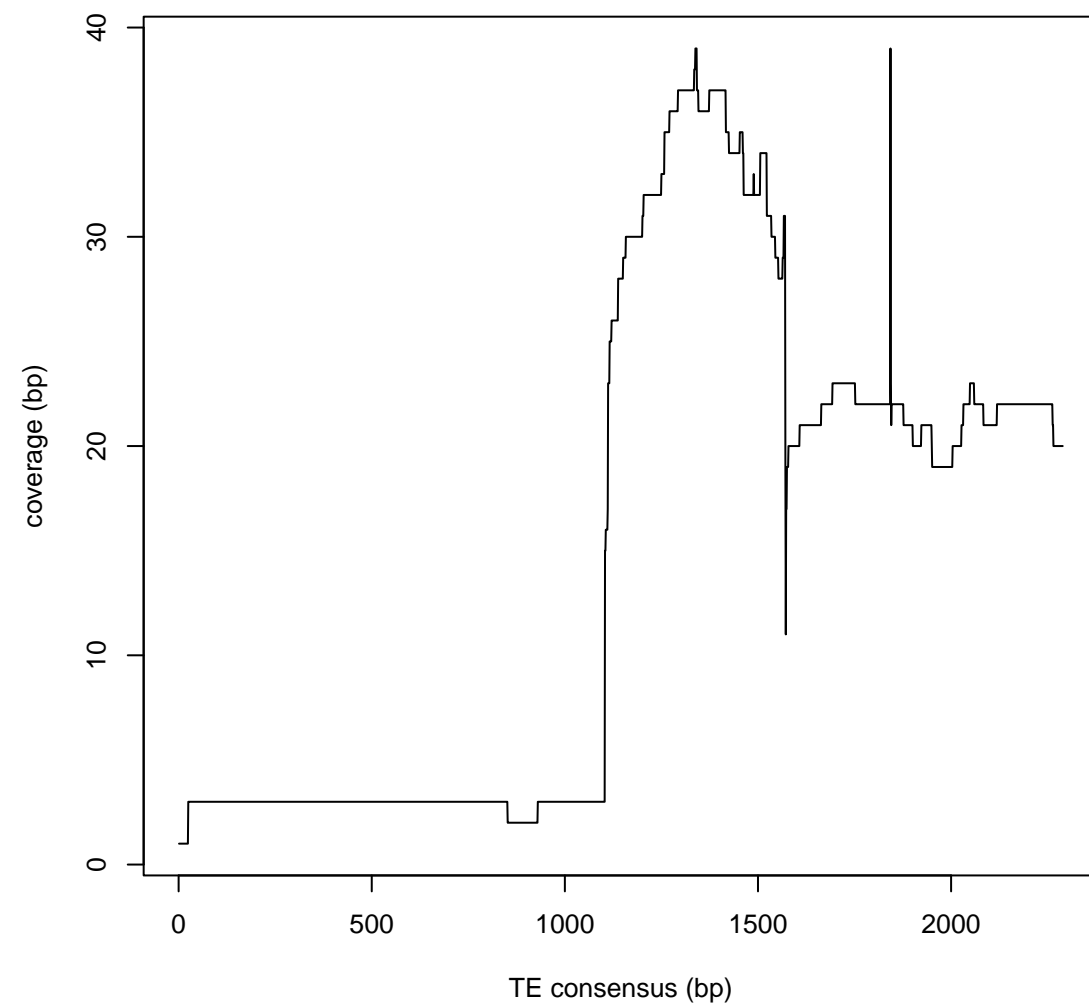


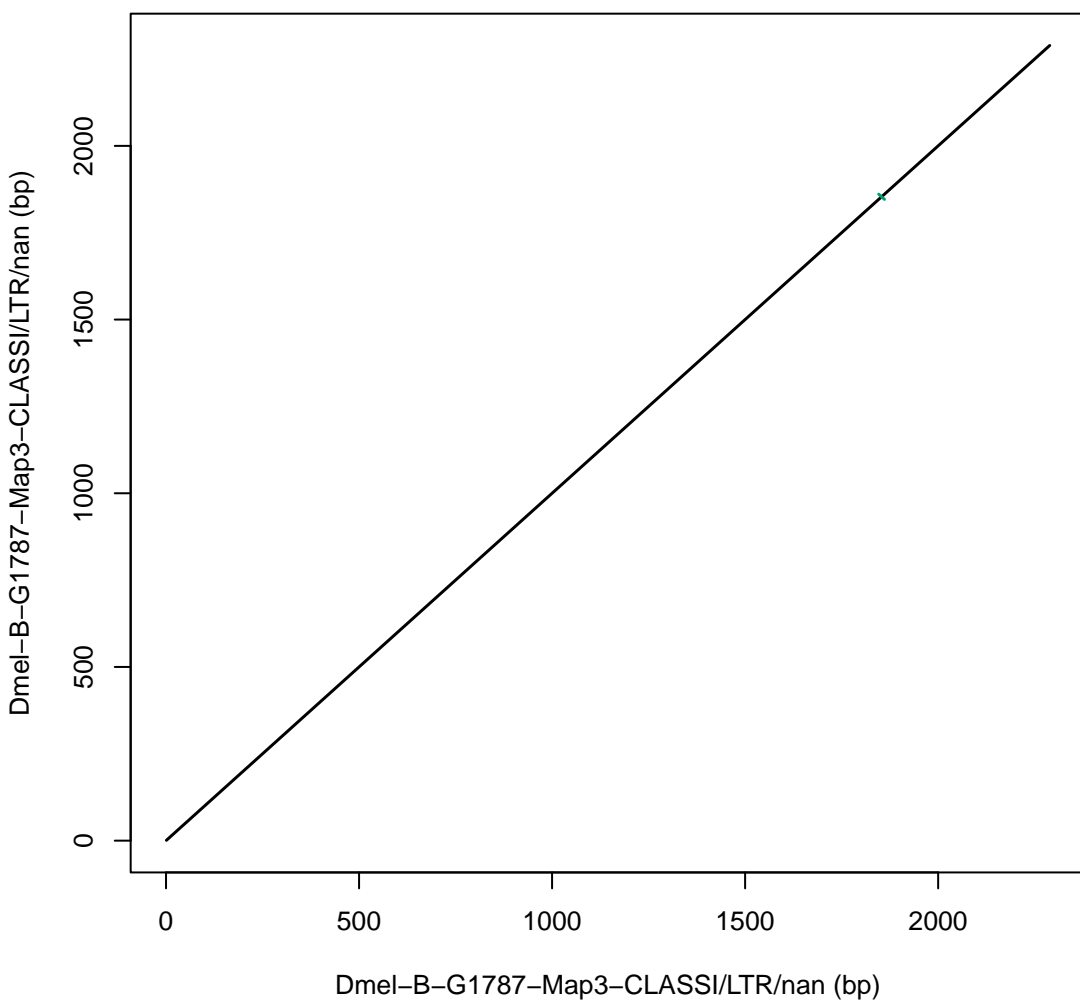
TE: Dmel-B-G1787-Map3-CLASSI/LTR/nan
consensus size: 2289bp; fragments: 90; full length: 1 (>=2060.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

