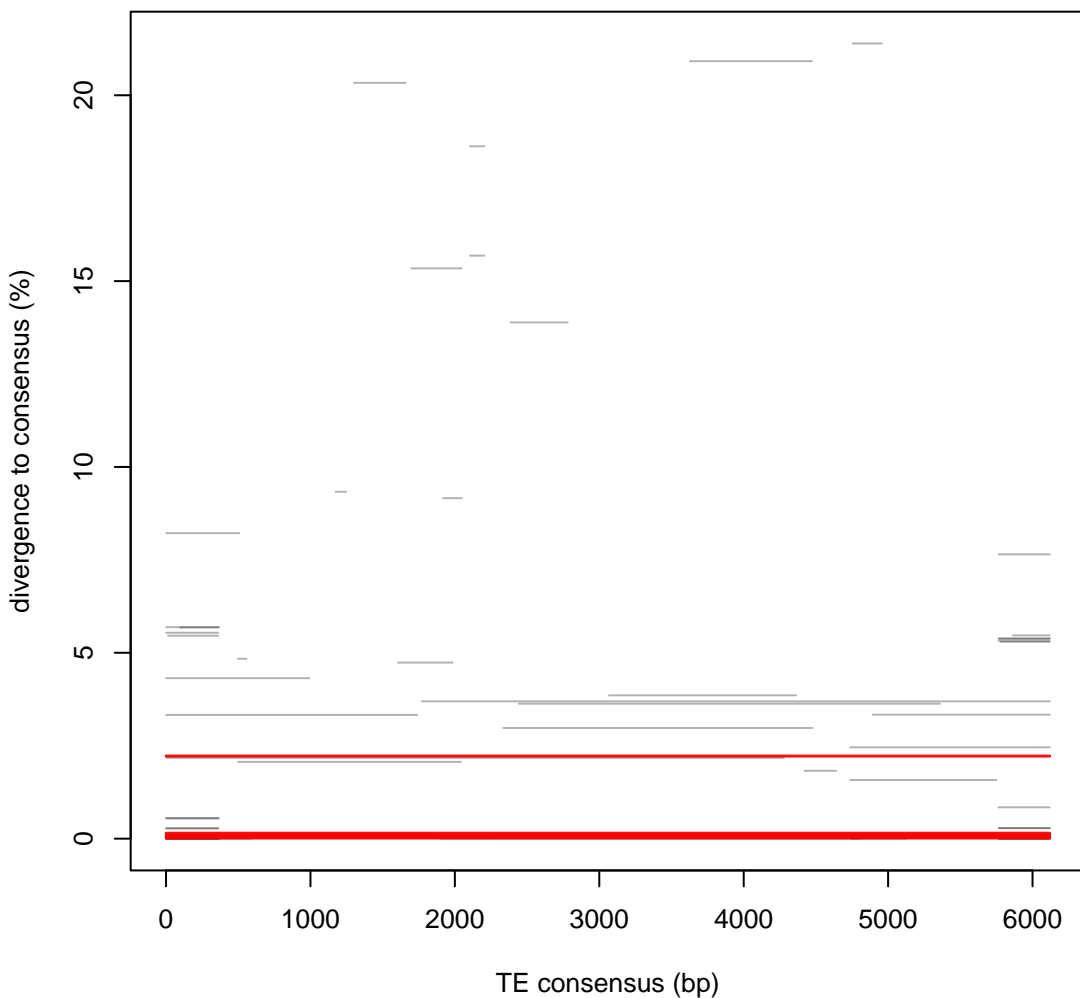
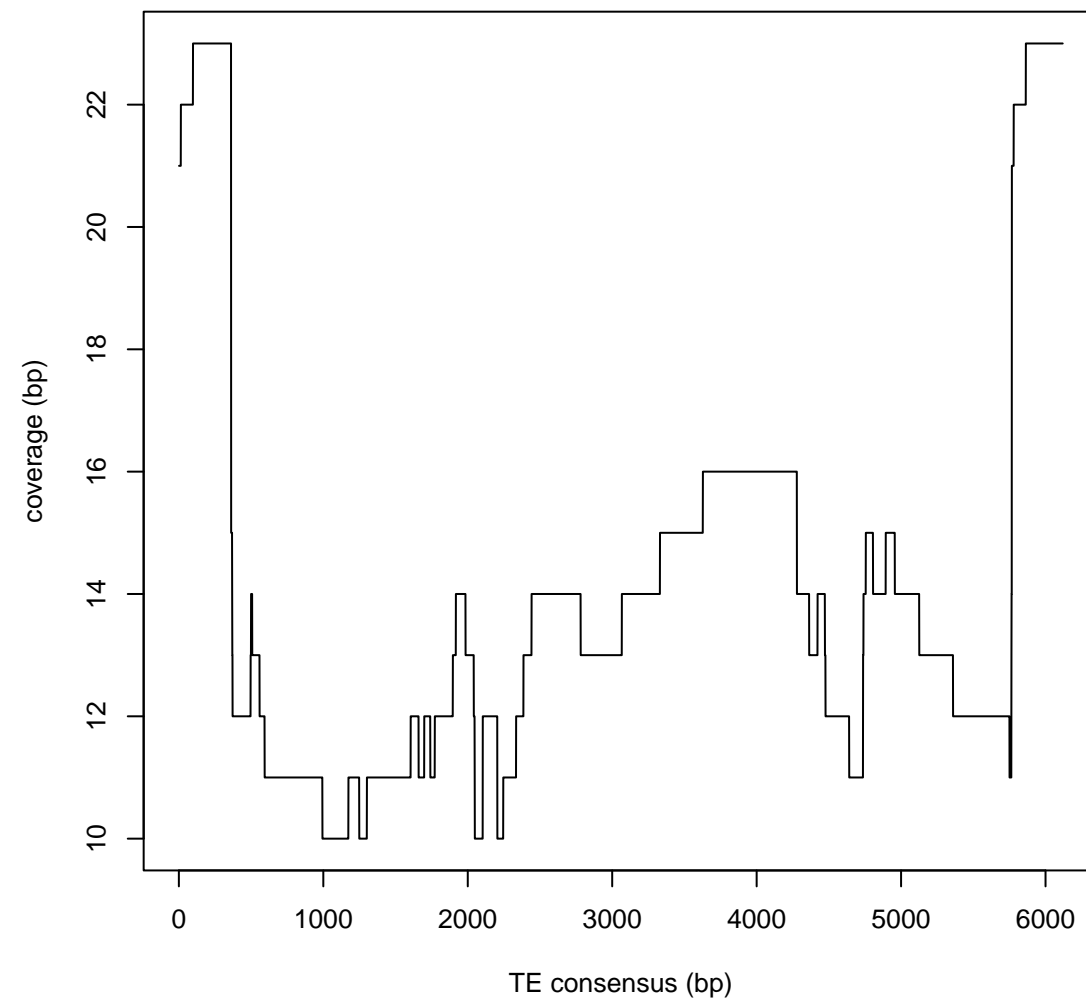


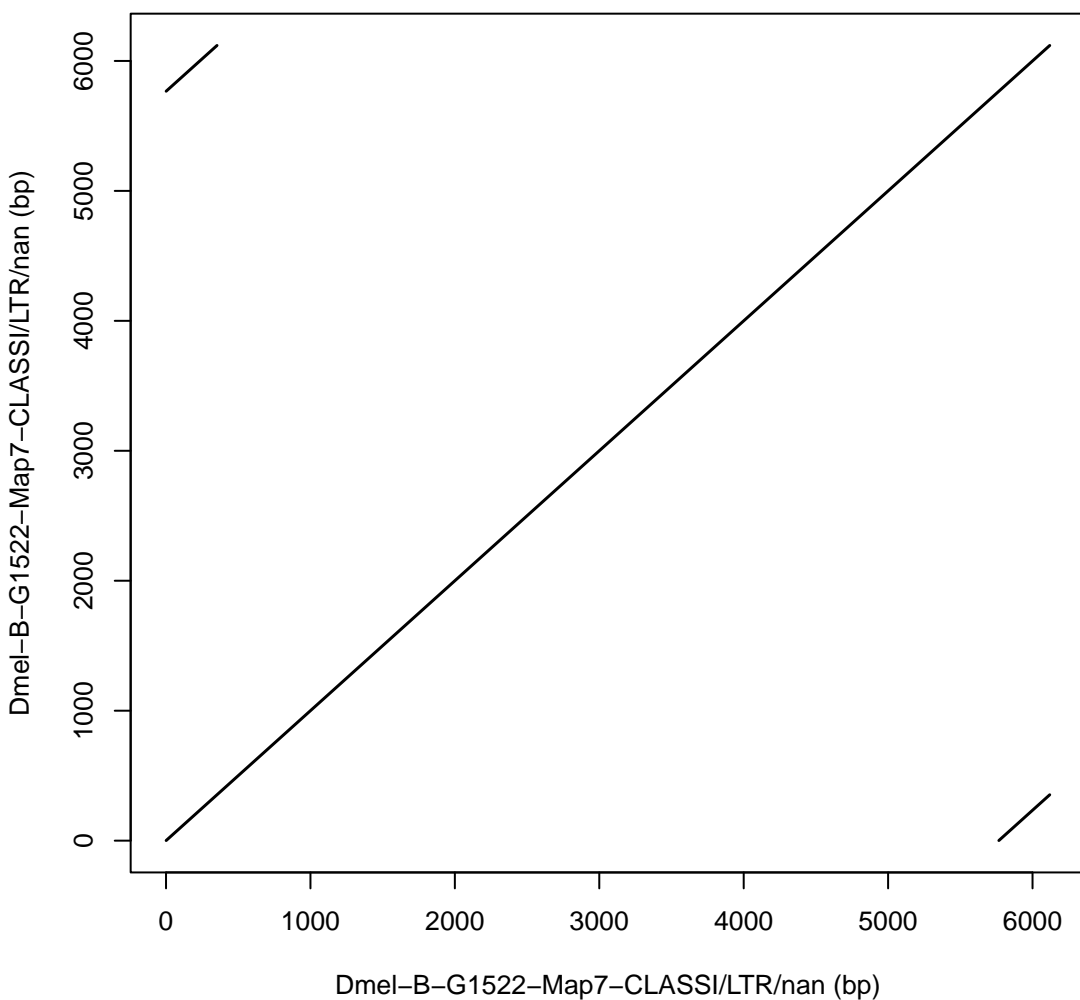
TE: Dmel-B-G1522-Map7-CLASSI/LTR/nan
consensus size: 6119bp; fragments: 59; full length: 6 (≥ 5507.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

