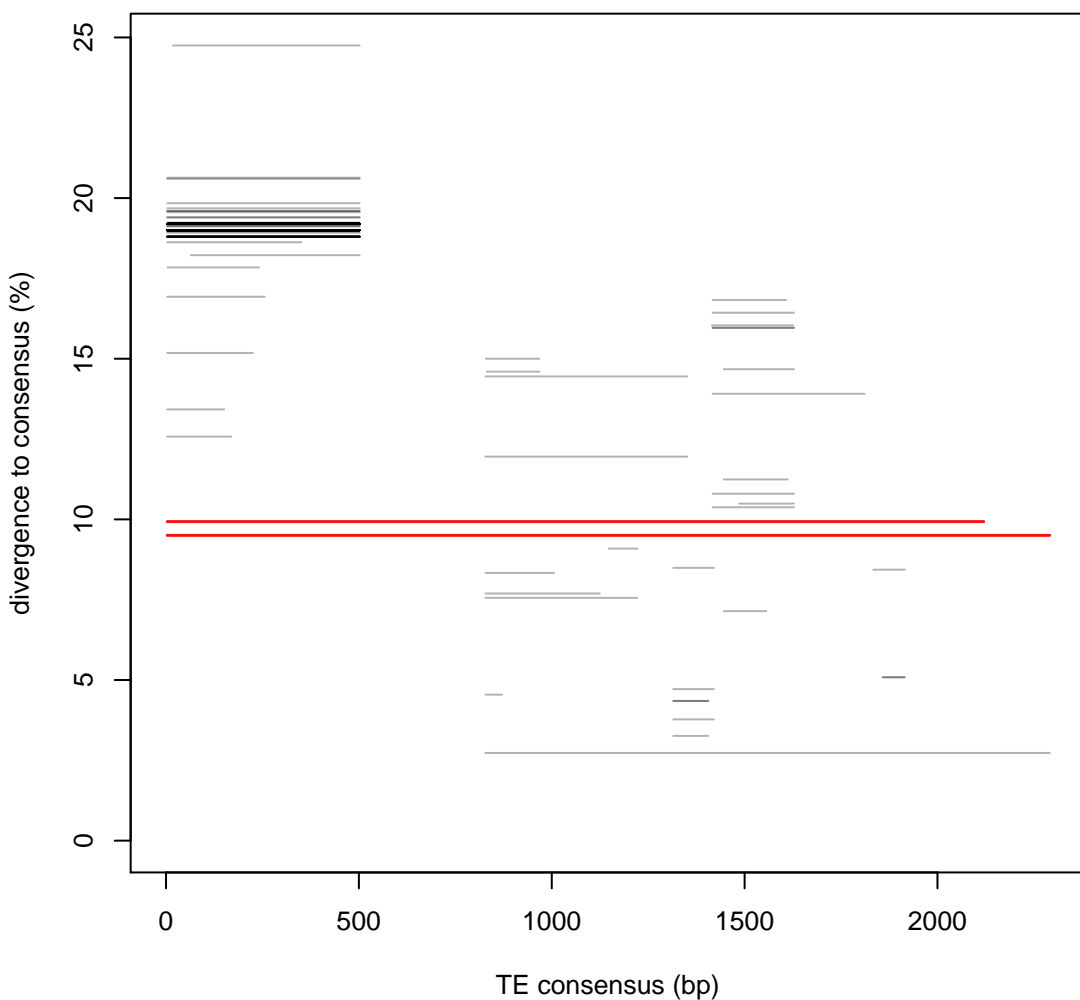
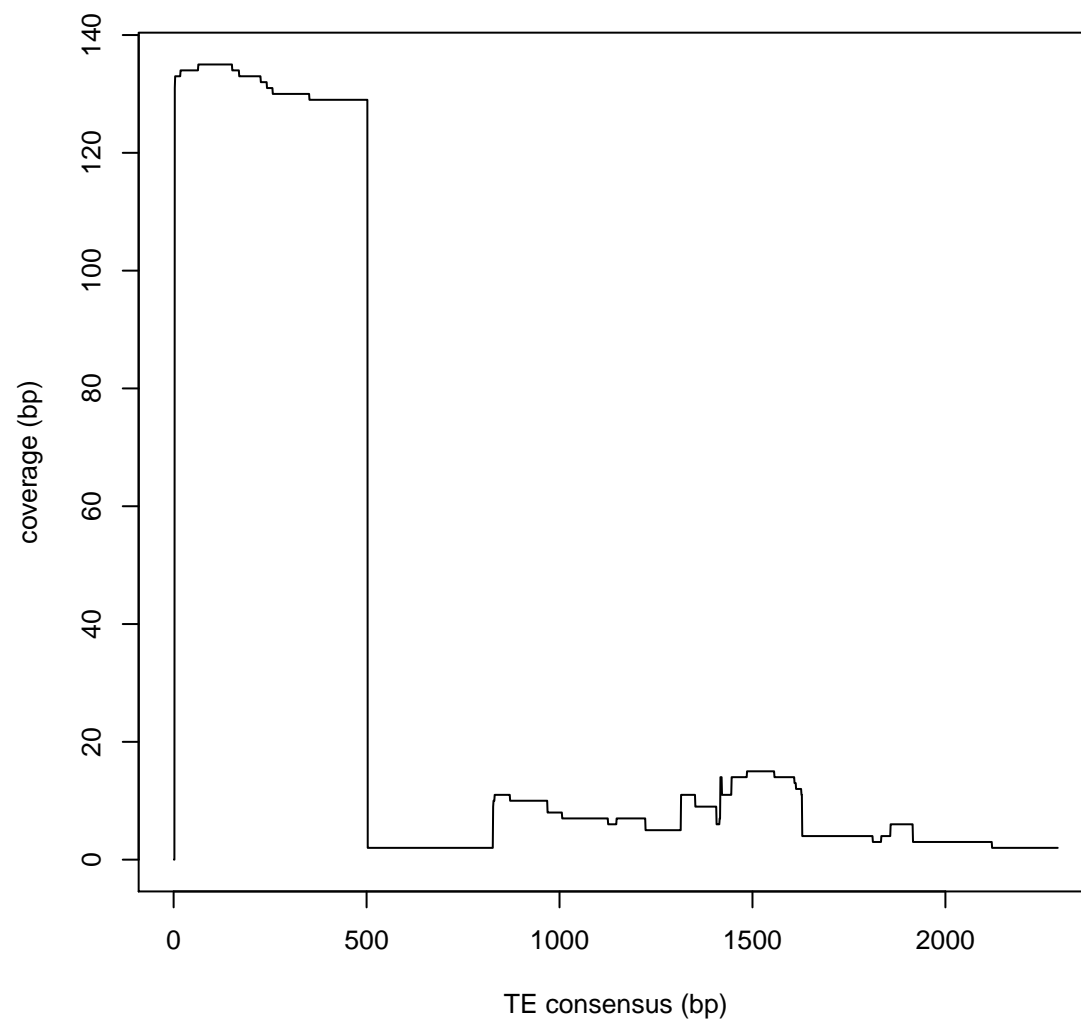


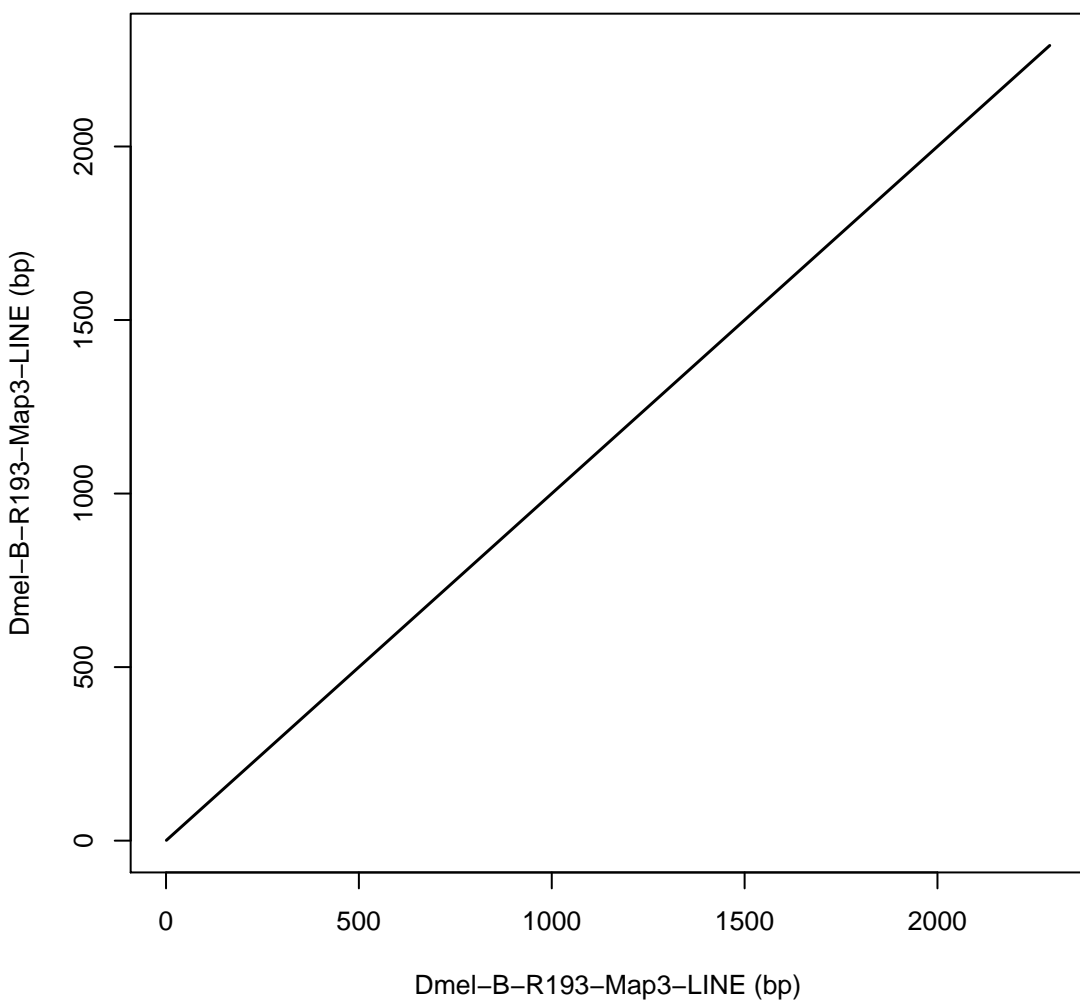
TE: Dmel-B-R193-Map3-LINE
consensus size: 2291bp; fragments: 166; full length: 2 (≥ 2061.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

