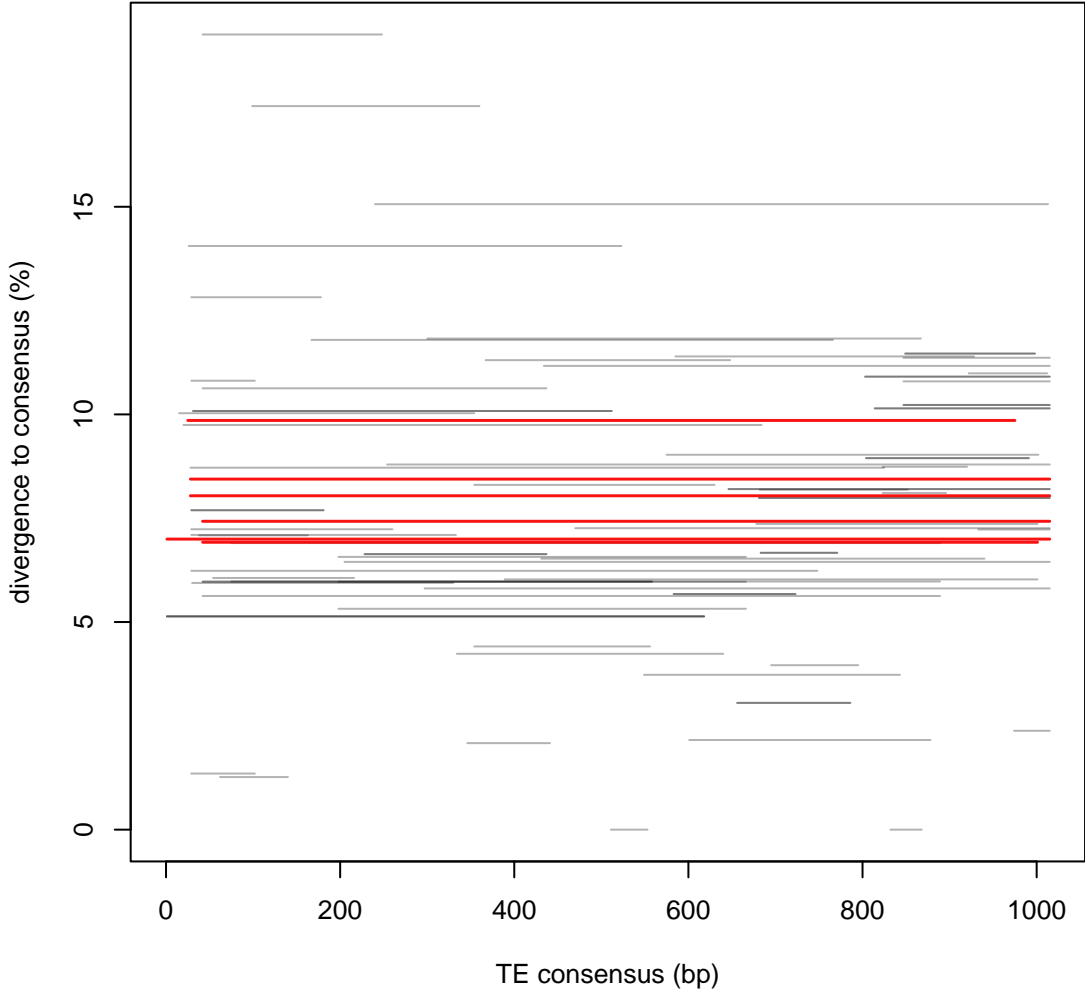
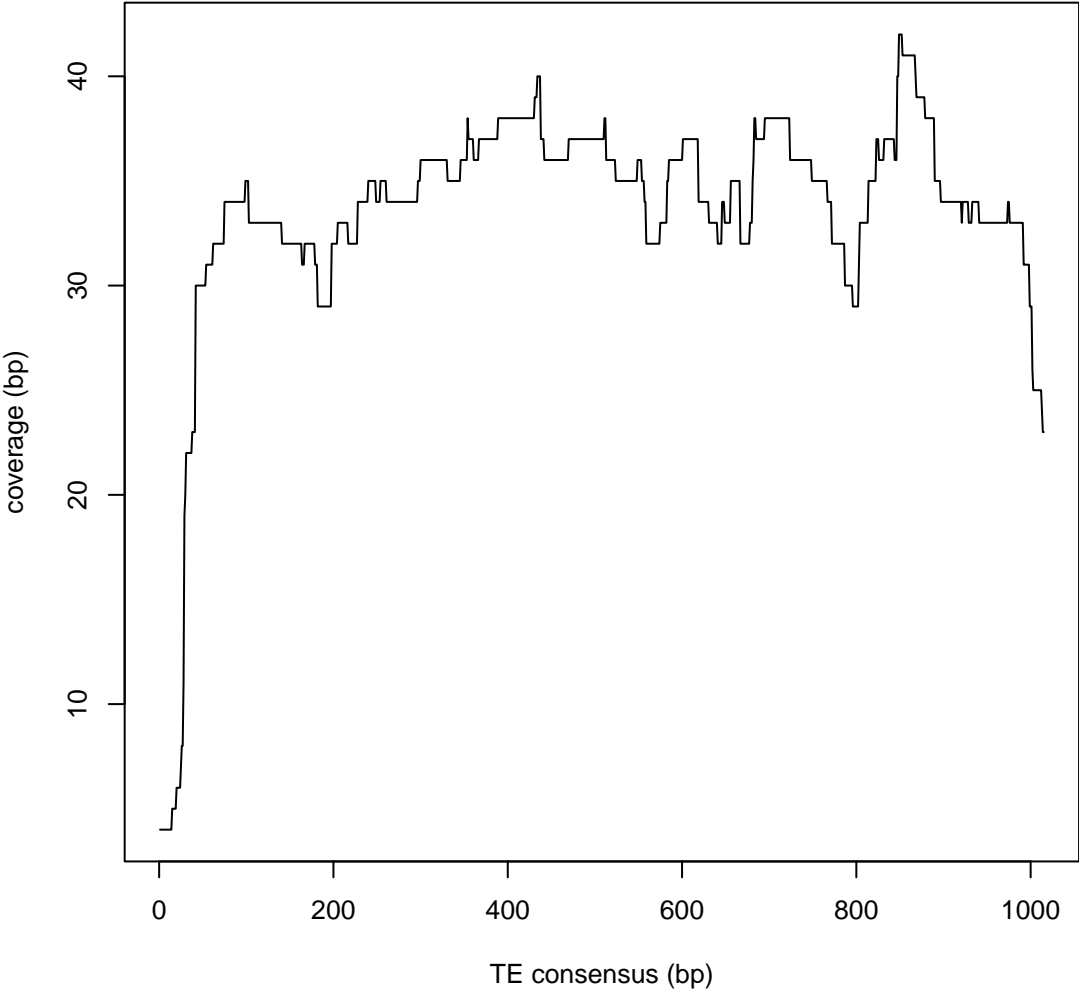


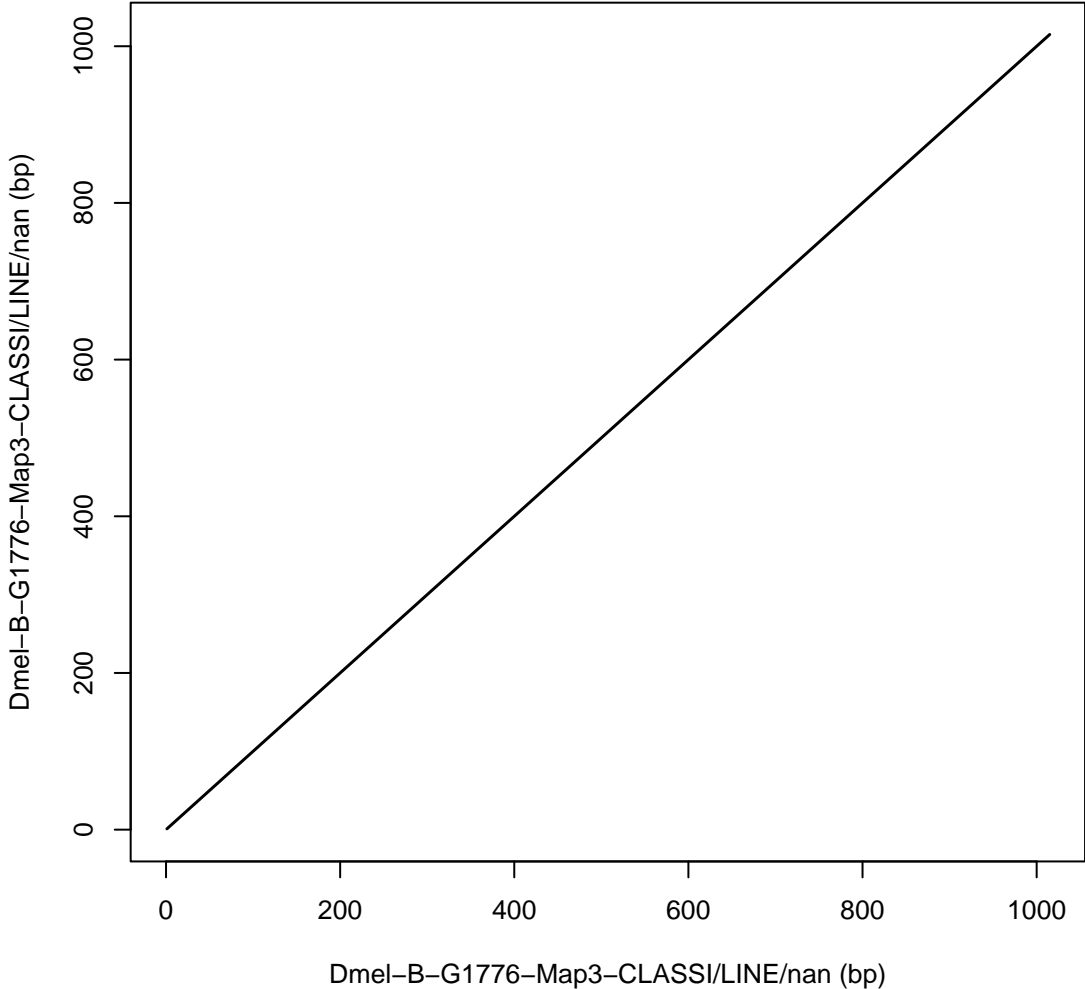
TE: Dmel-B-G1776-Map3-CLASSI/LINE/nan
consensus size: 1015bp; fragments: 91; full length: 6 (>=913.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

