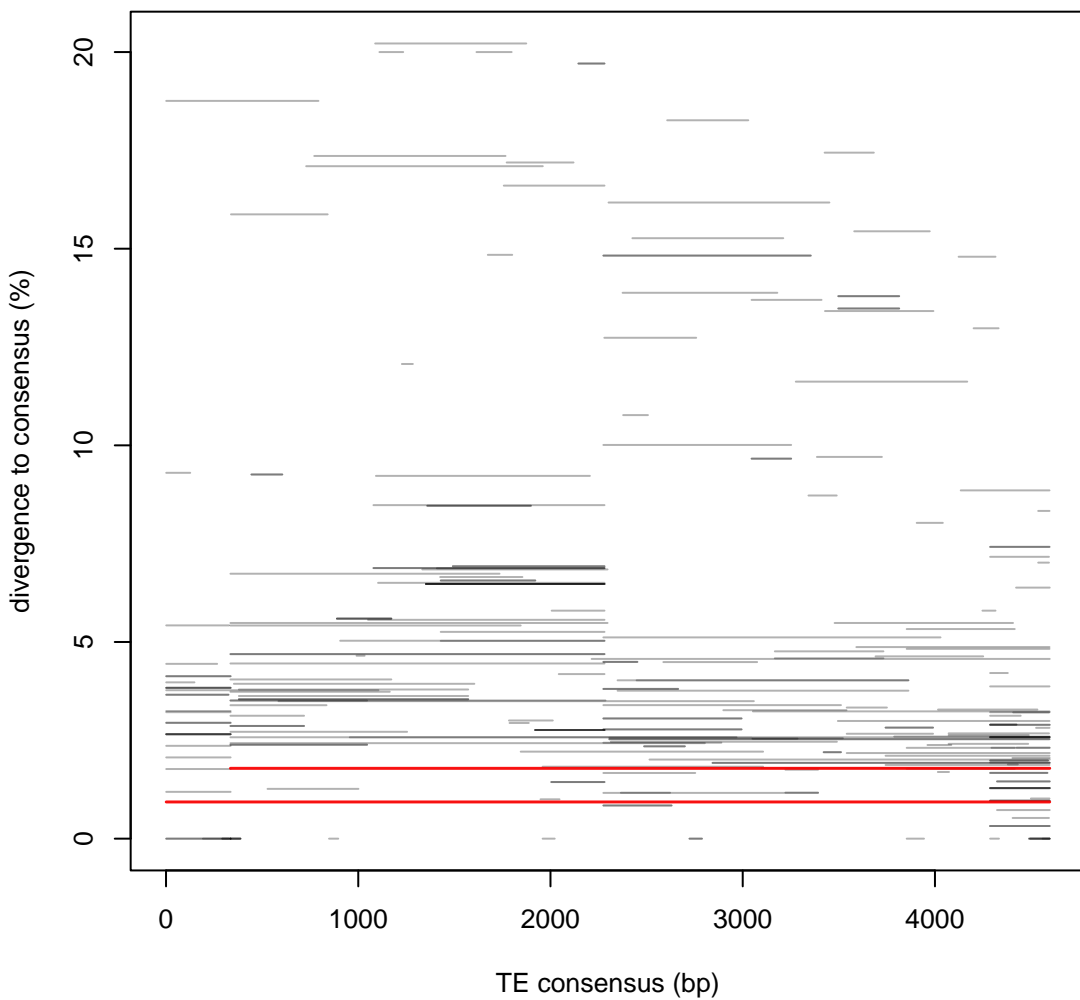
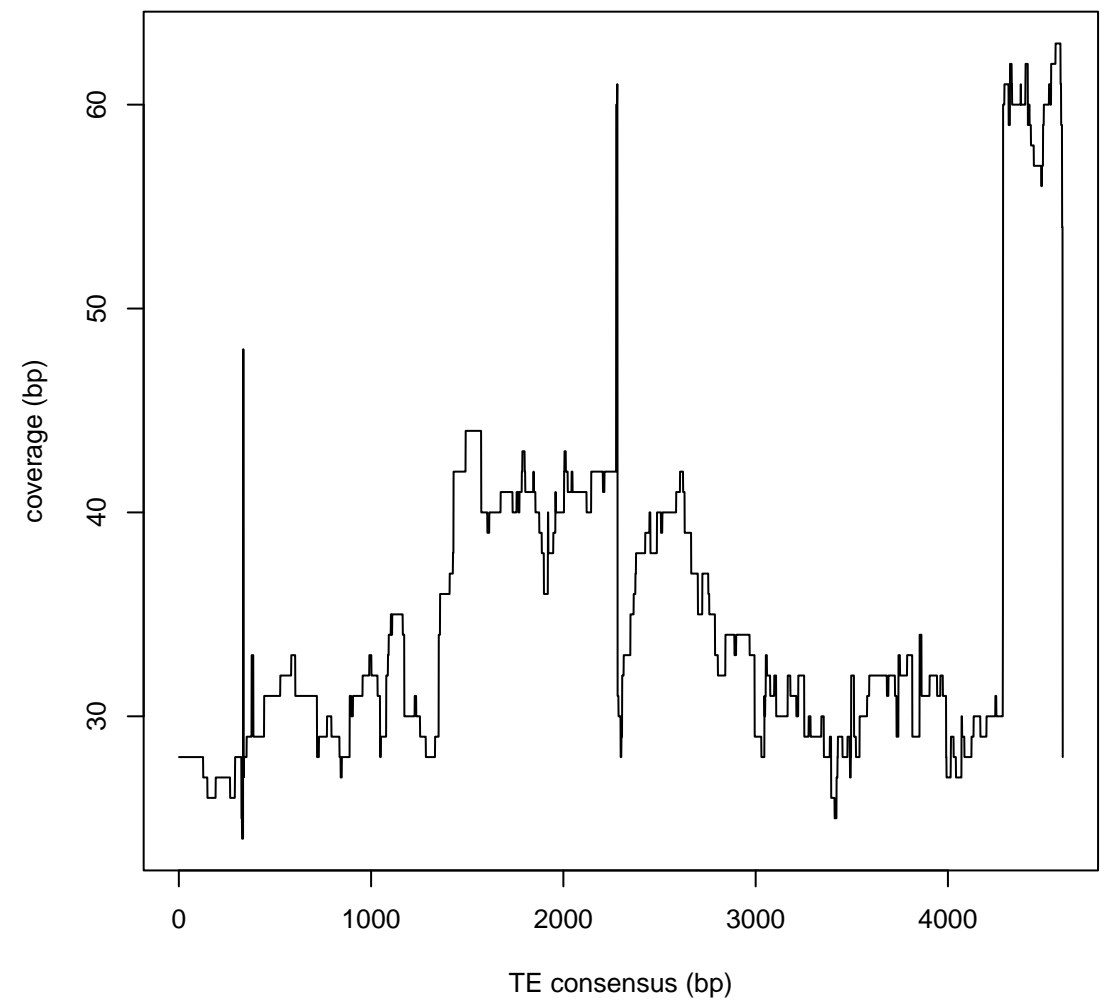


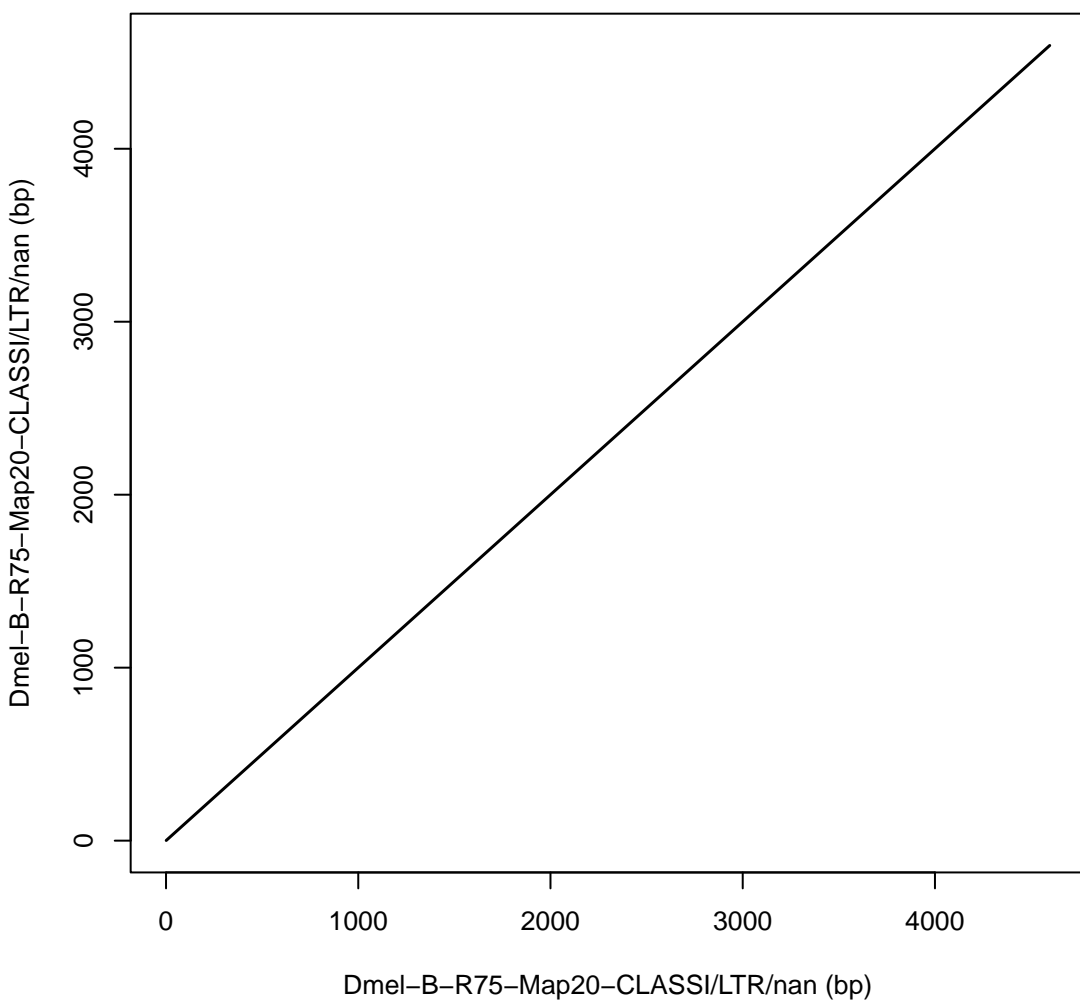
TE: Dmel-B-R75-Map20-CLASSI/LTR/nan
consensus size: 4597bp; fragments: 261; full length: 2 (≥ 4137.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

