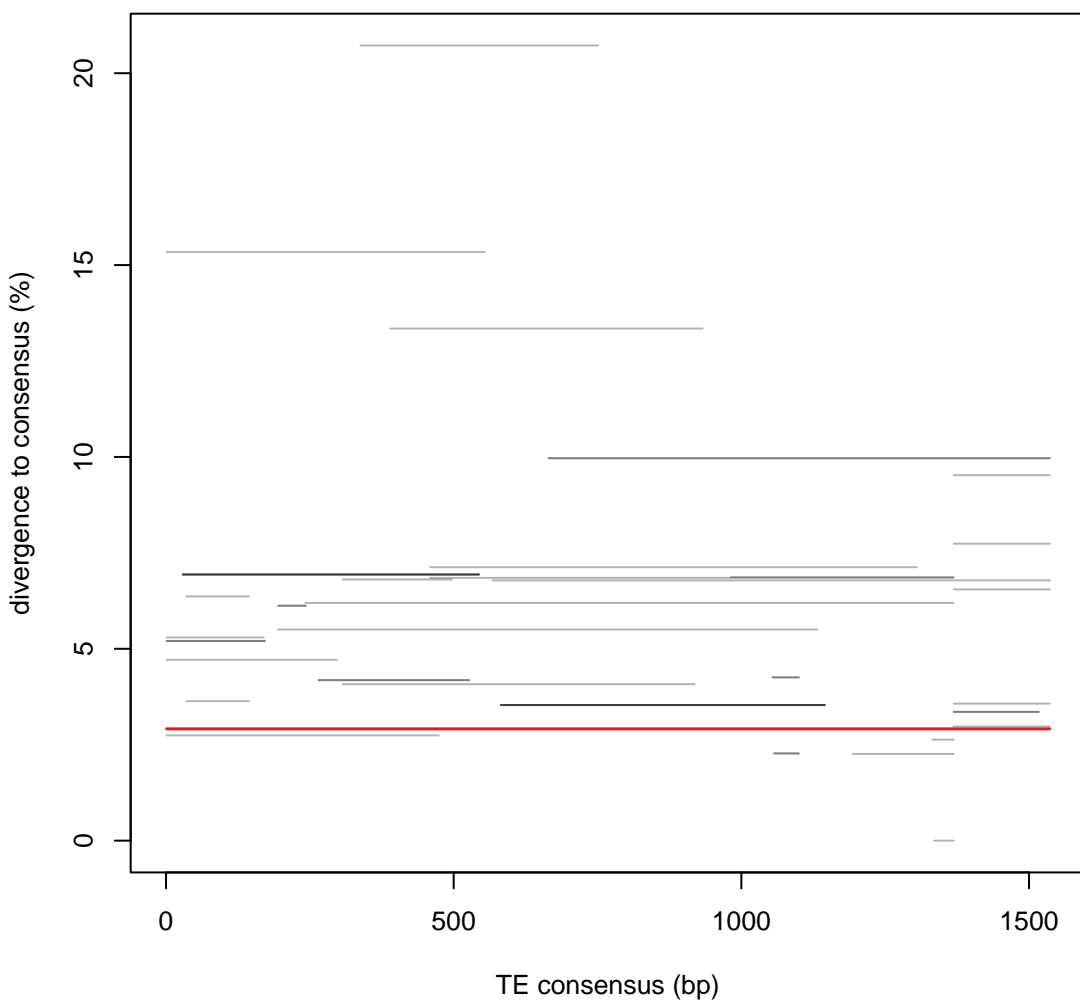
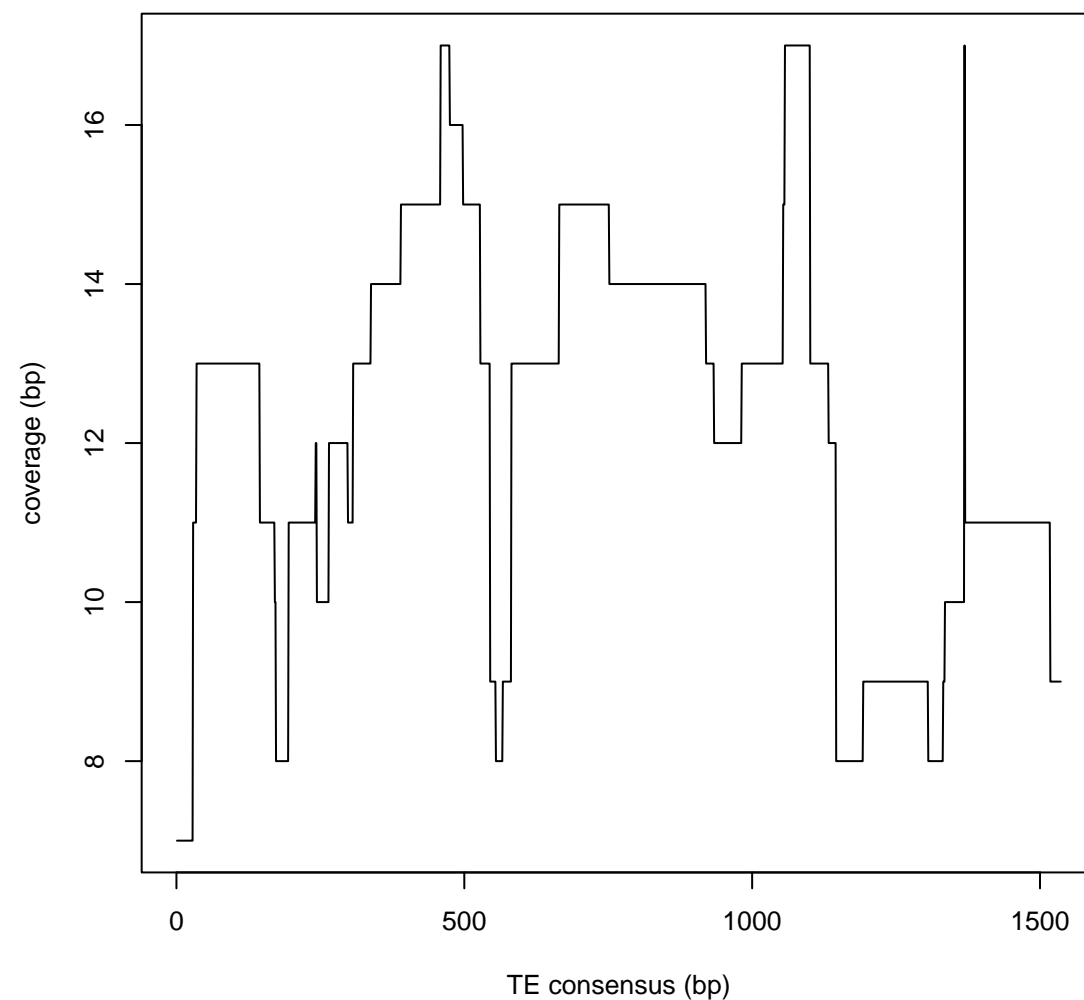


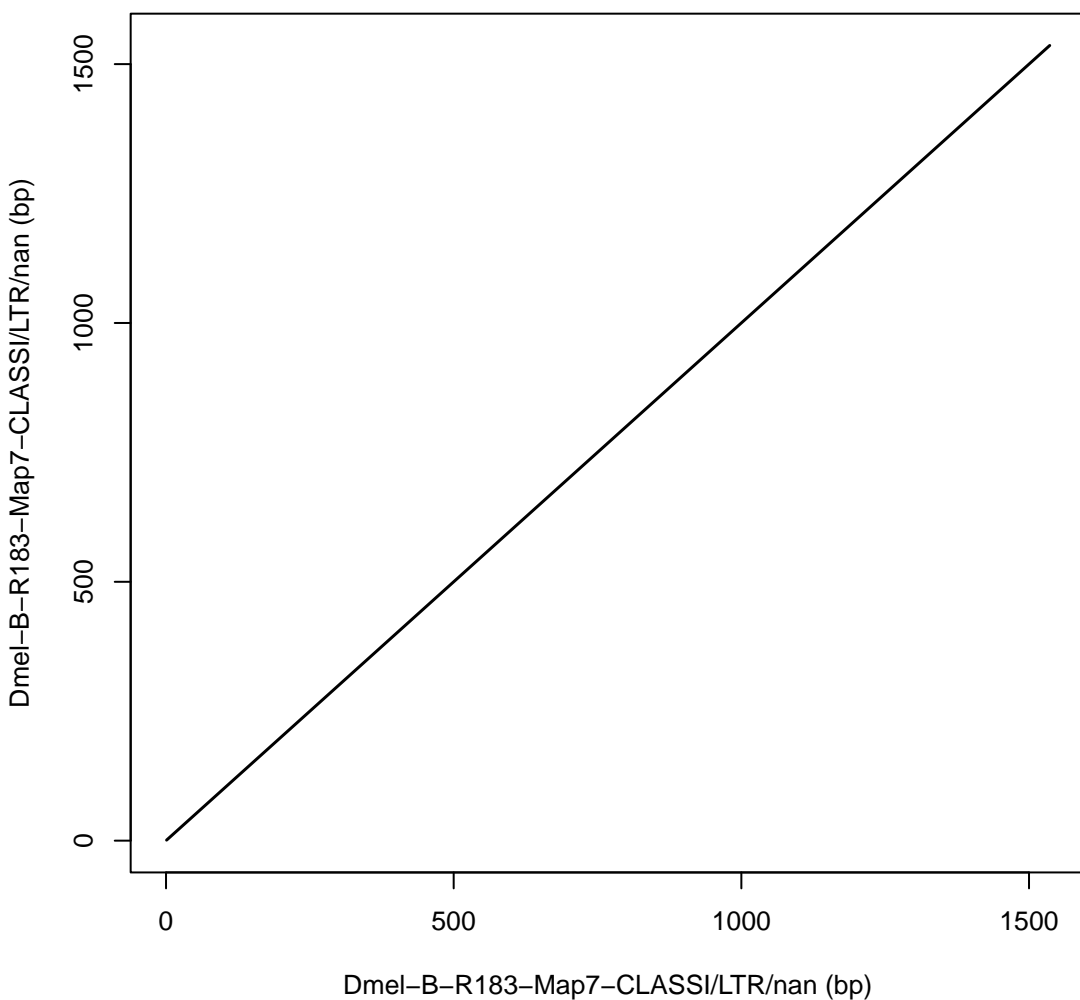
TE: Dmel-B-R183-Map7-CLASSI/LTR/nan  
consensus size: 1536bp; fragments: 47; full length: 1 (>=1382.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

