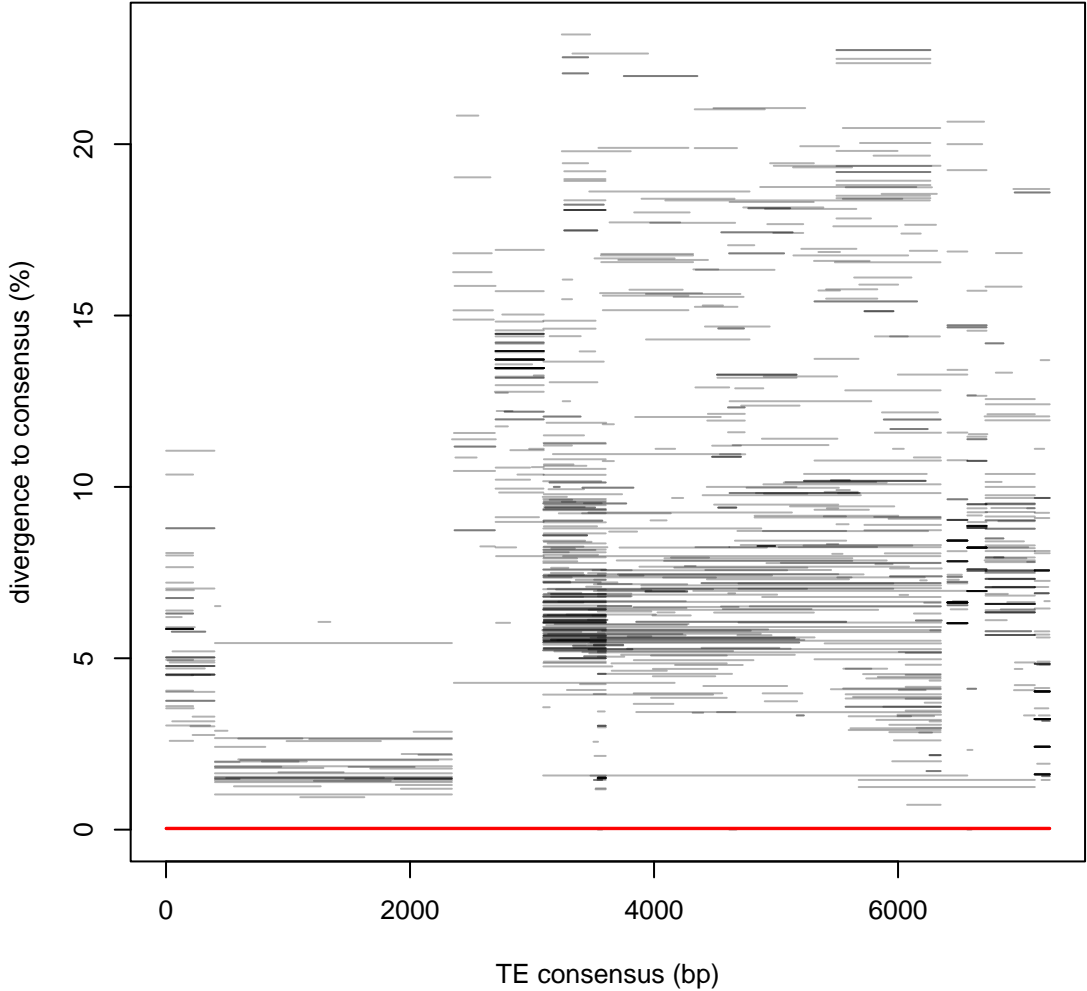
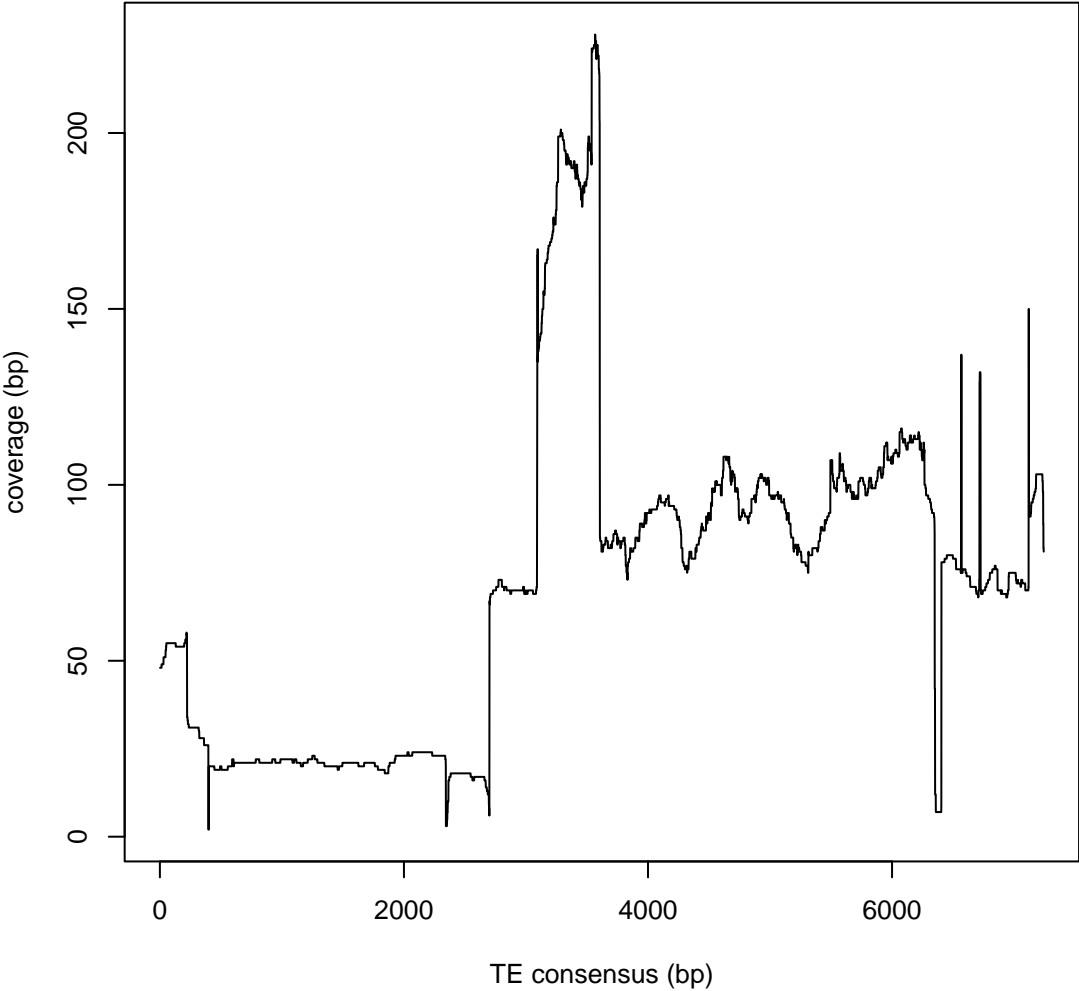


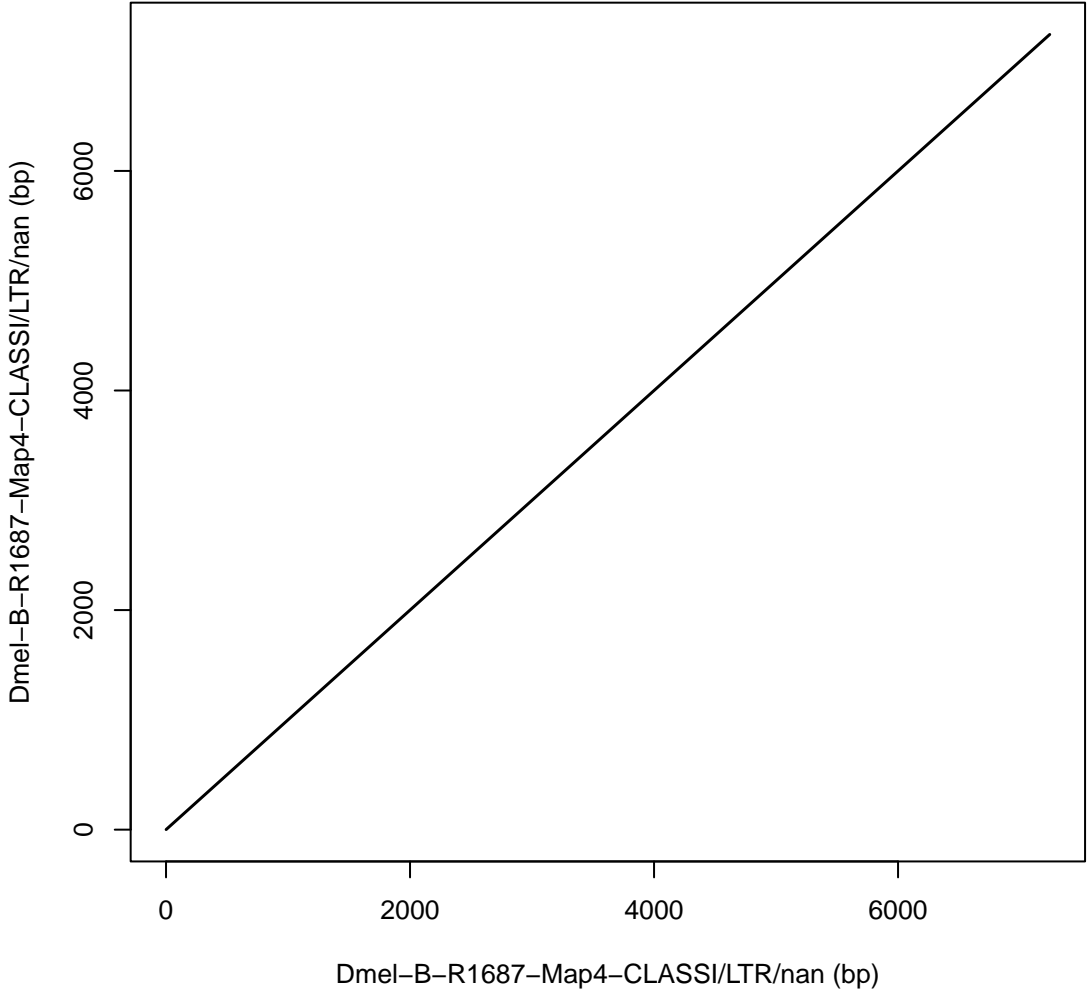
TE: Dmel-B-R1687-Map4-CLASSI/LTR/nan
consensus size: 7243bp; fragments: 1083; full length: 2 (>=6518.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

