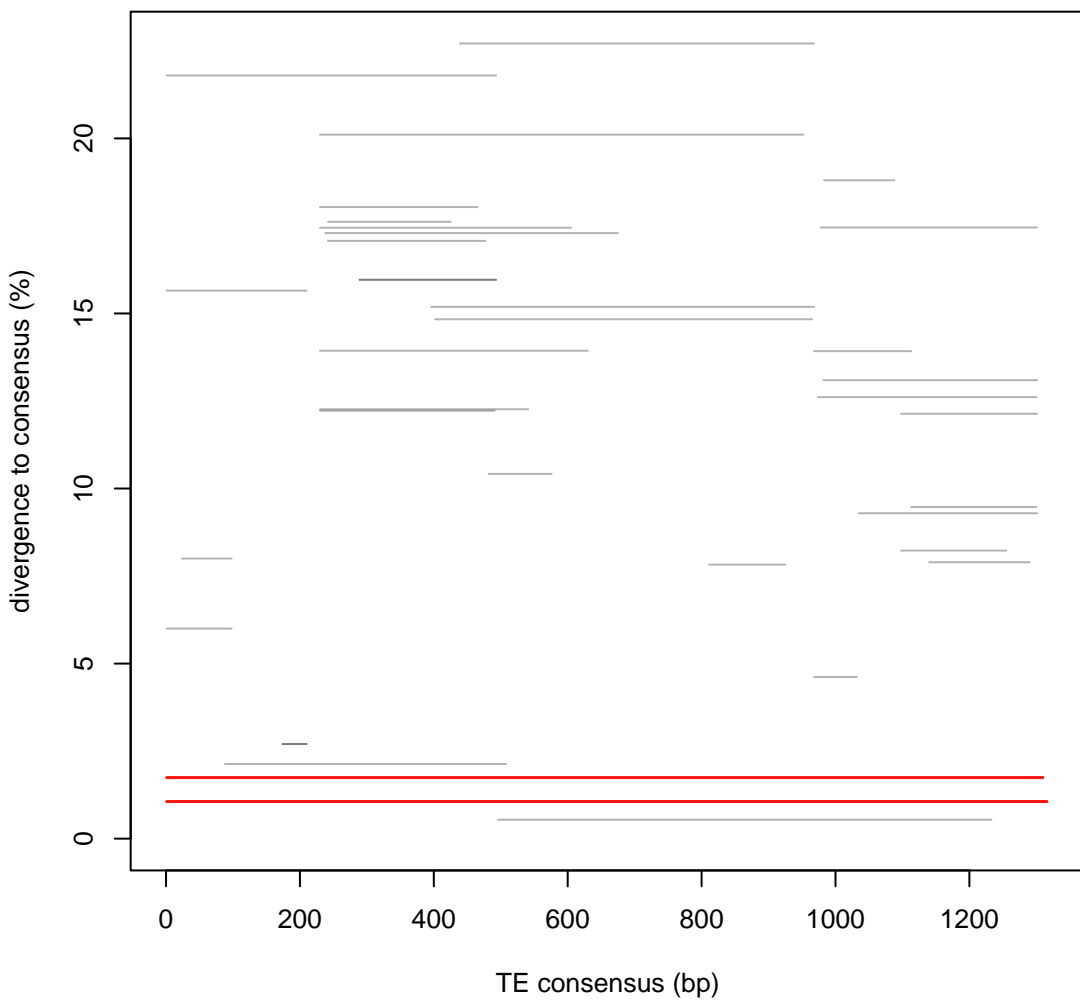
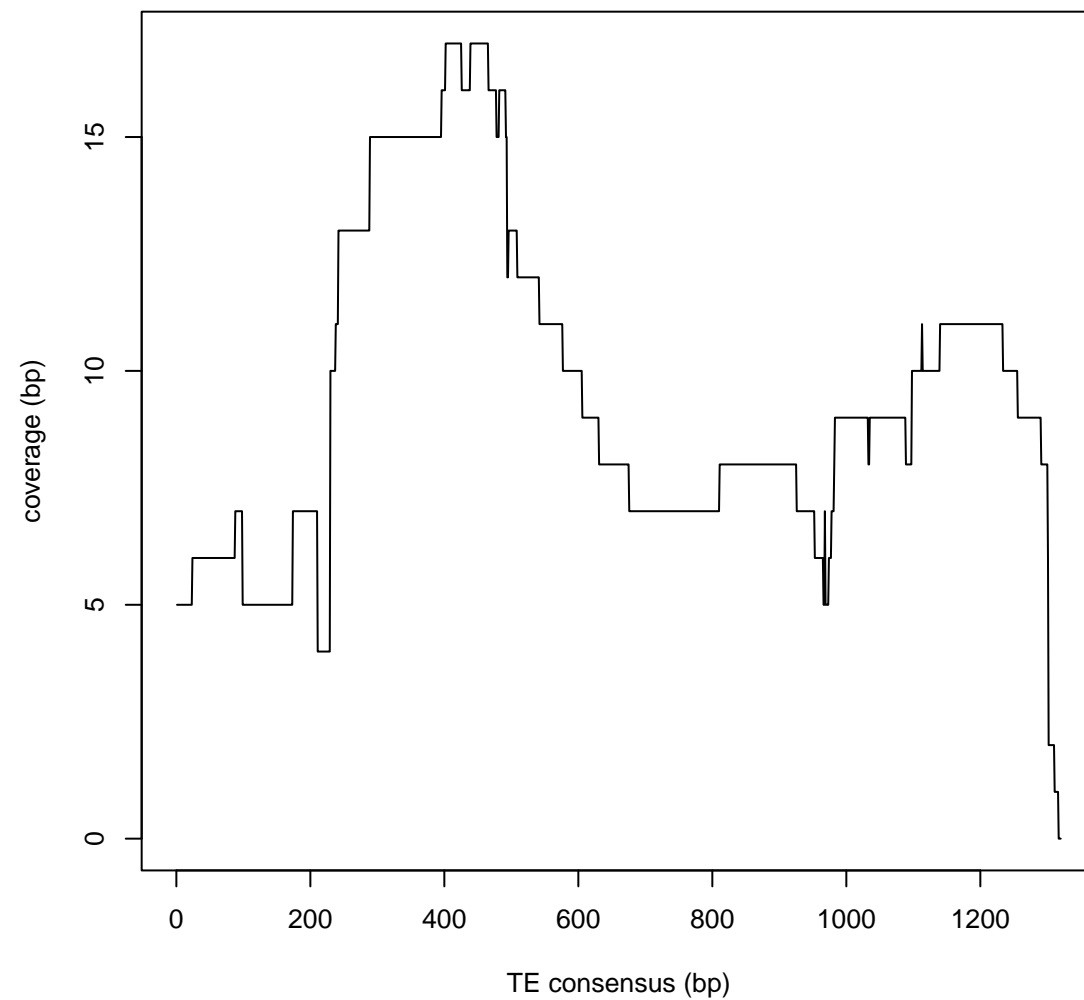


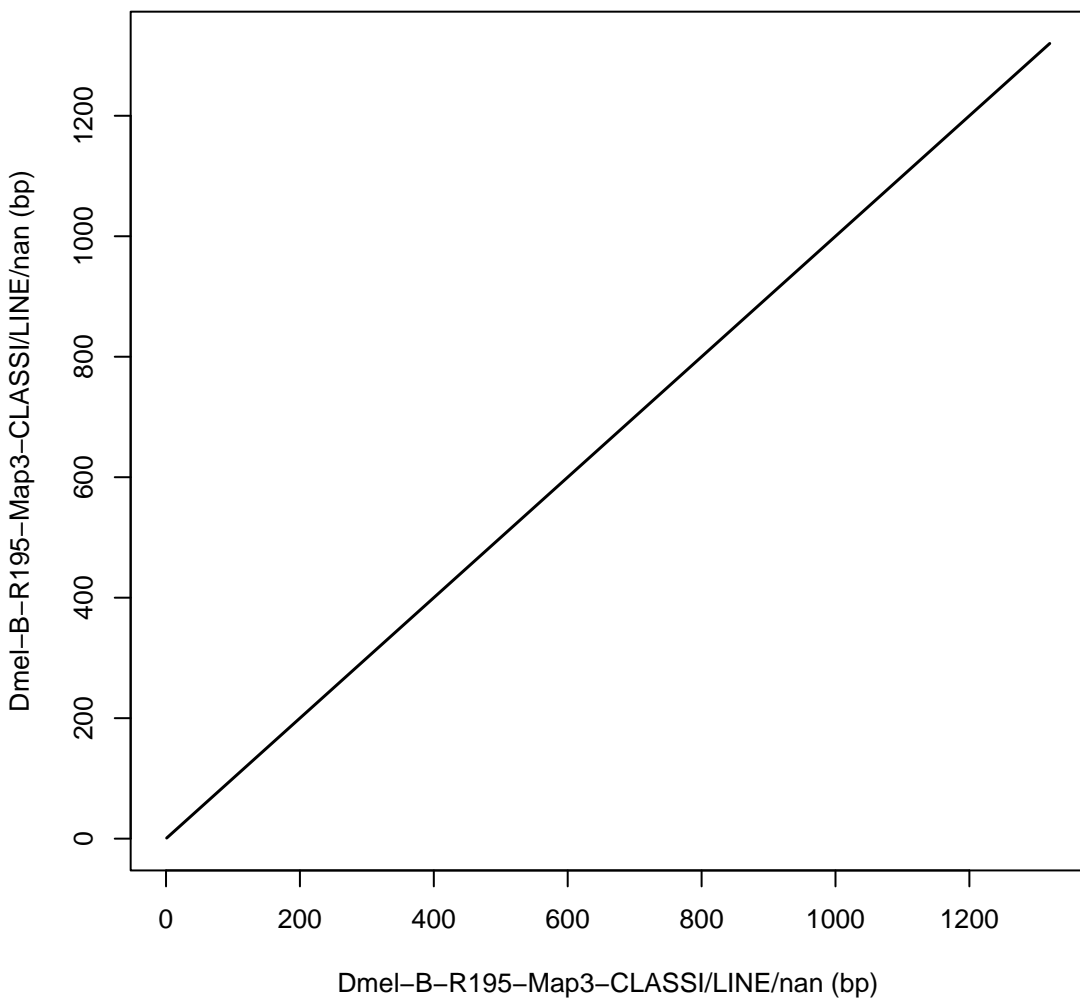
TE: Dmel-B-R195-Map3-CLASSI/LINE/nan  
consensus size: 1320bp; fragments: 37; full length: 2 (>=1188bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

