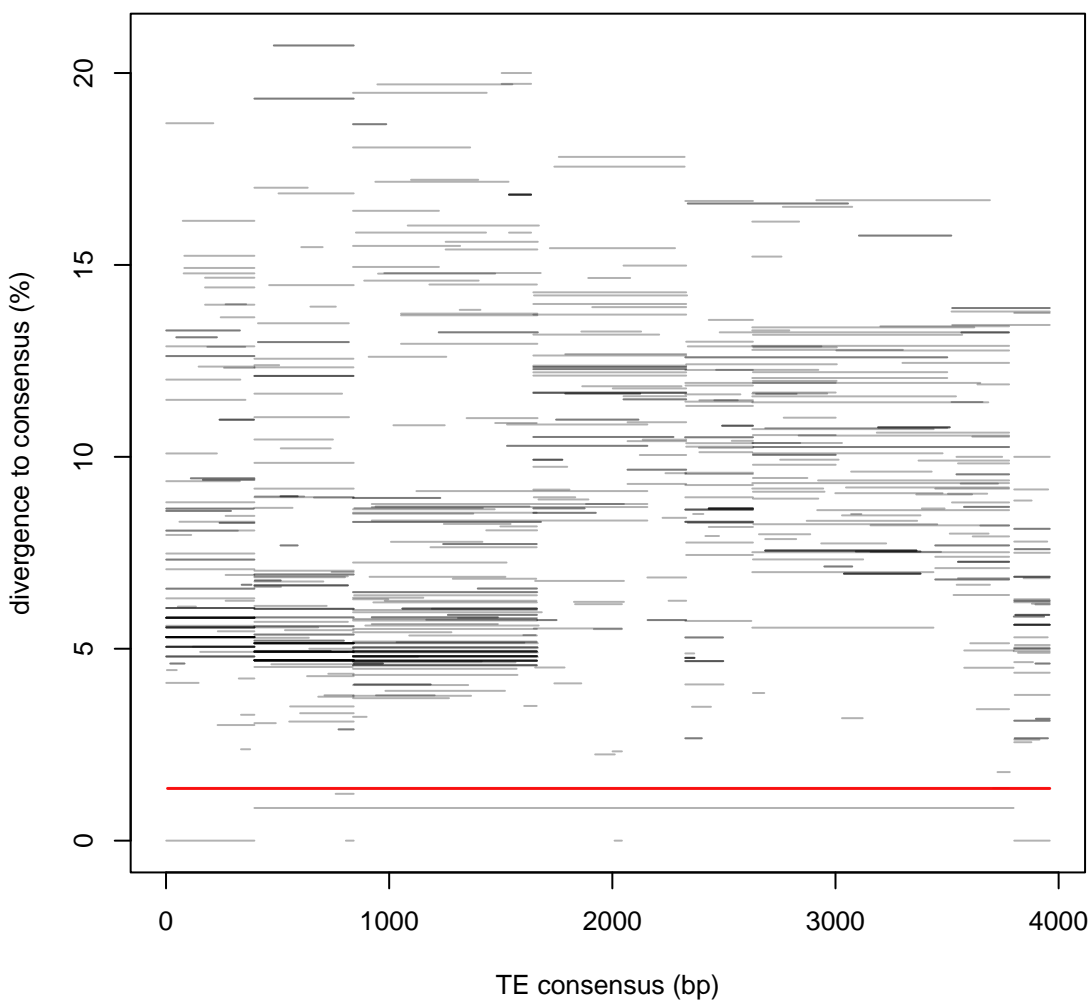
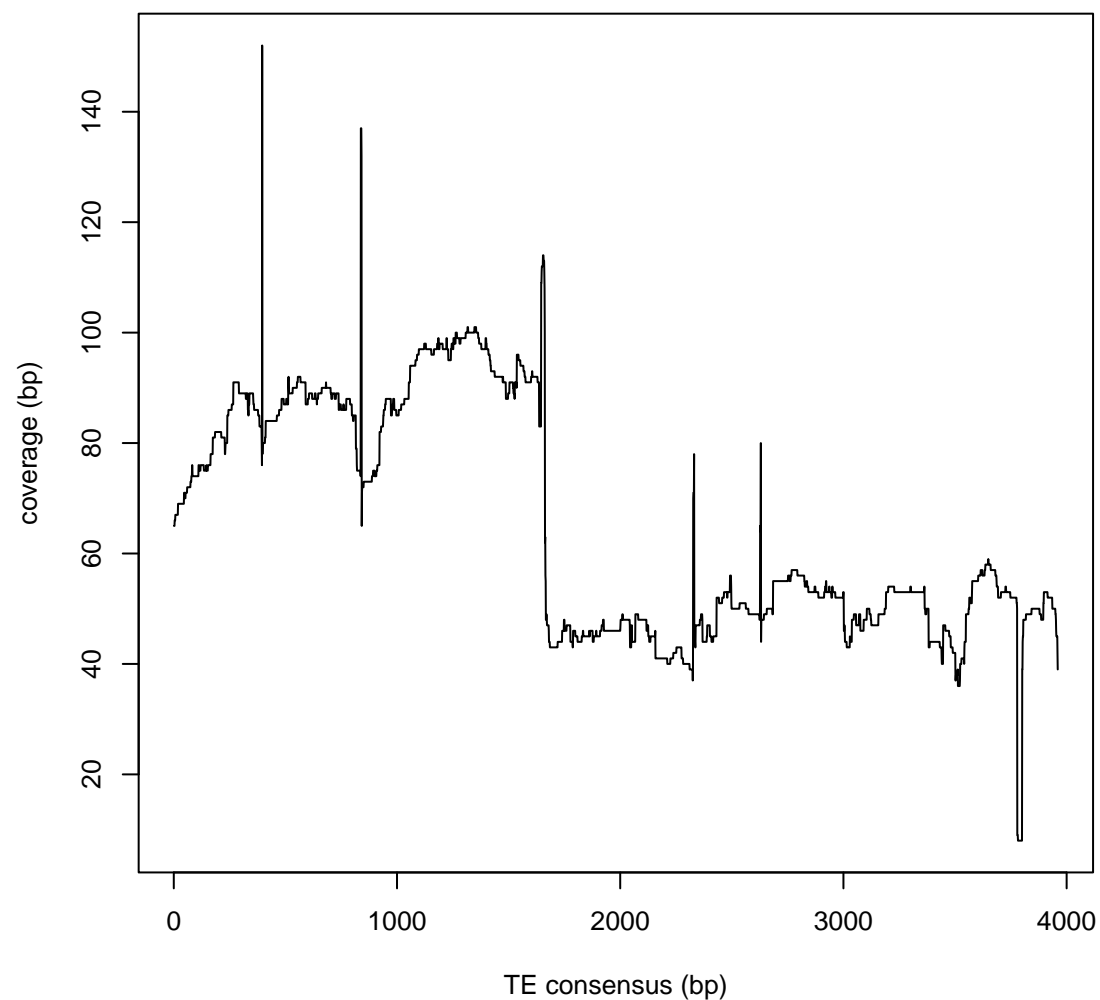


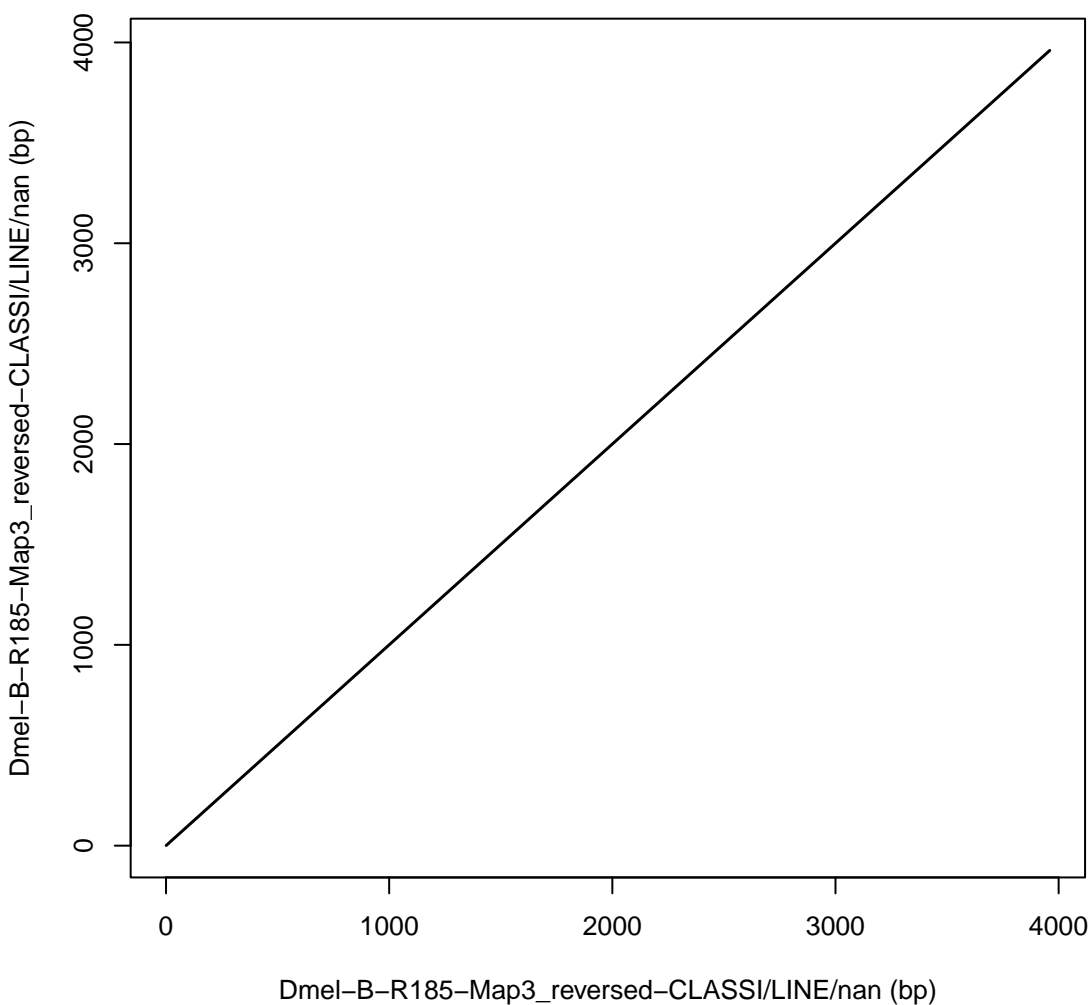
TE: Dmel-B-R185-Map3_reversed-CLASSI/LINE/nan
consensus size: 3960bp; fragments: 664; full length: 1 (>=3564bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

