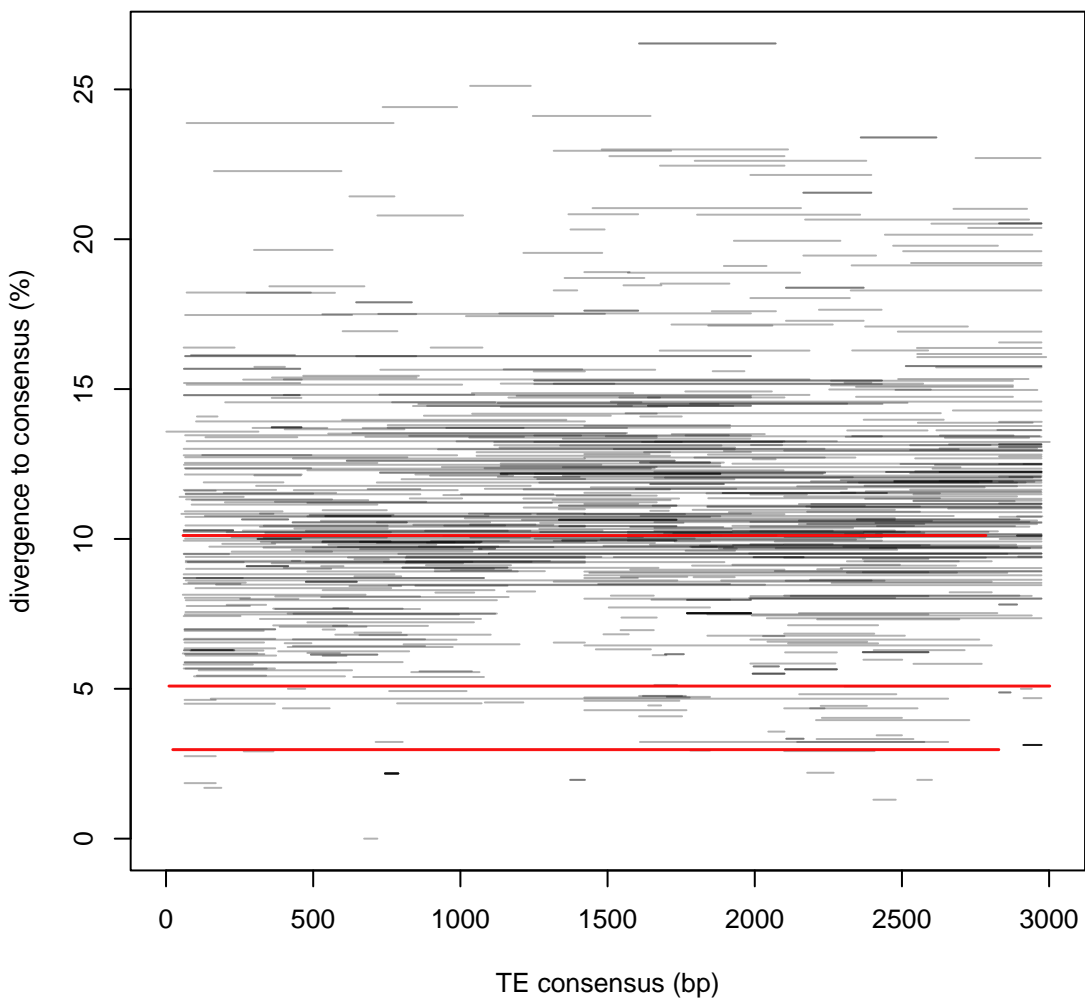
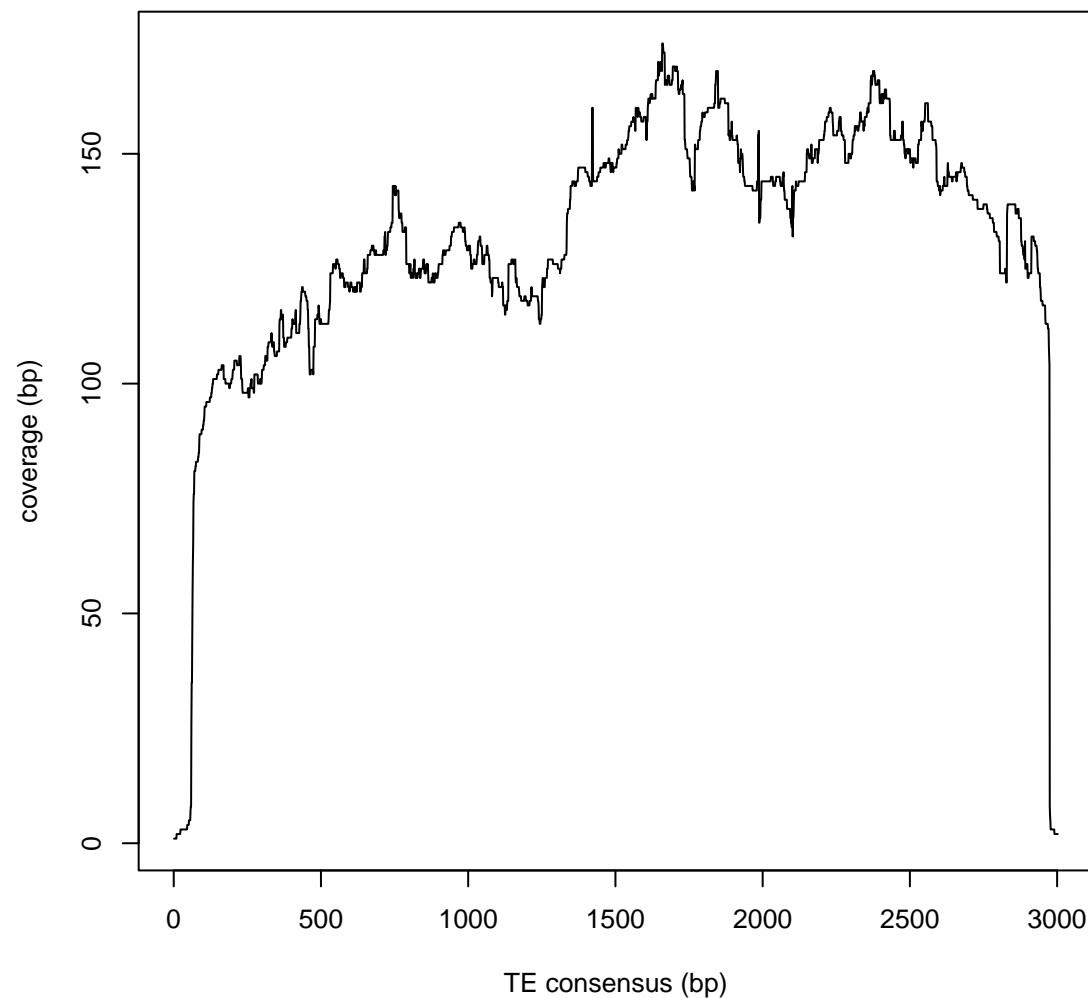


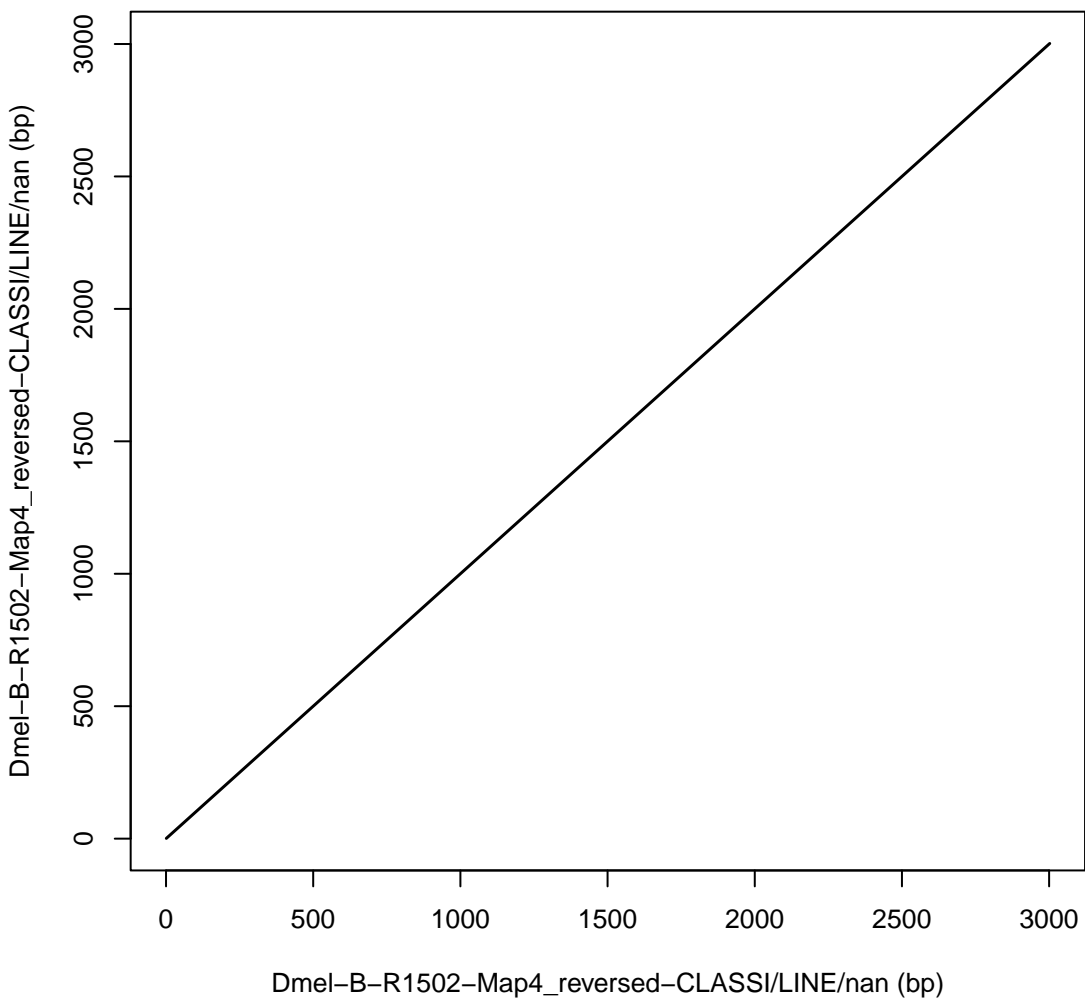
TE: Dmel-B-R1502-Map4_reversed-CLASSI/LINE/nan
consensus size: 3002bp; fragments: 734; full length: 3 (>=2701.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

