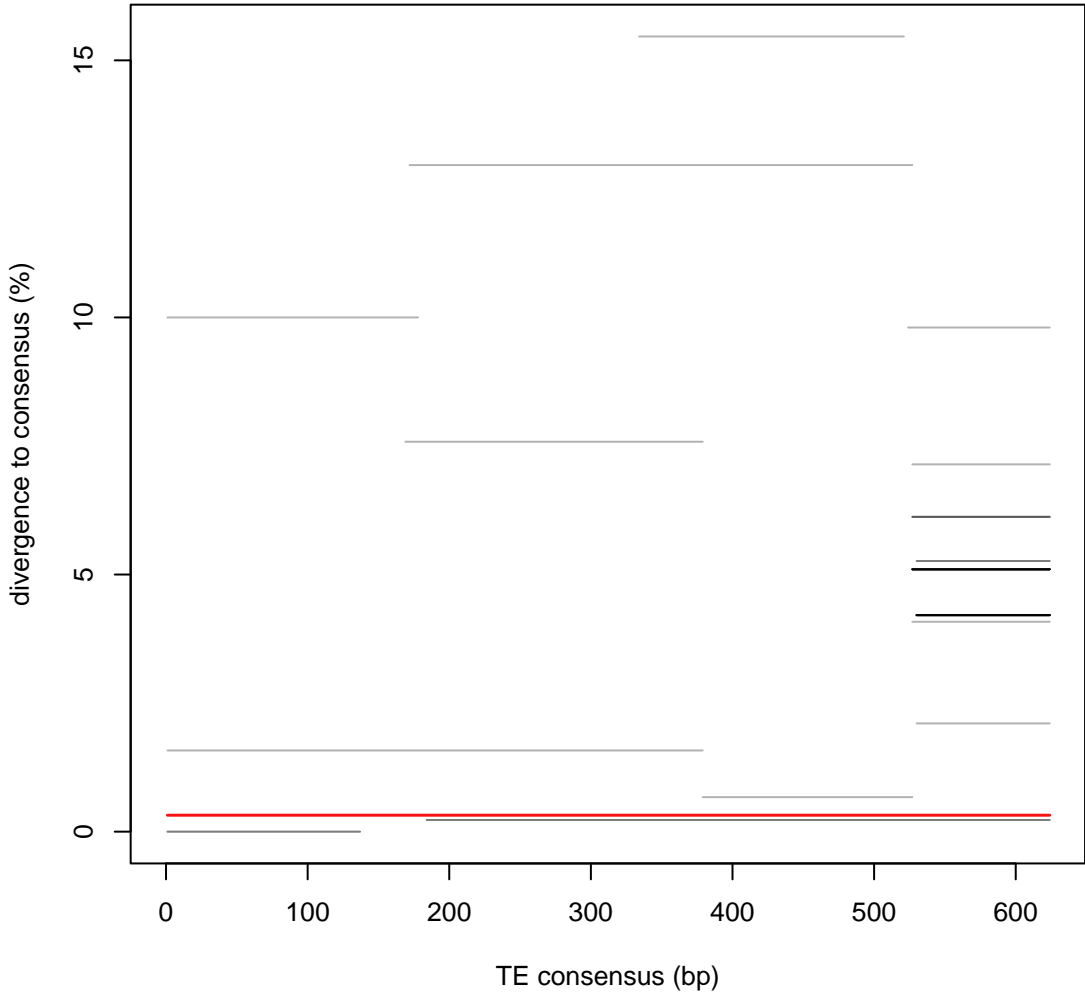
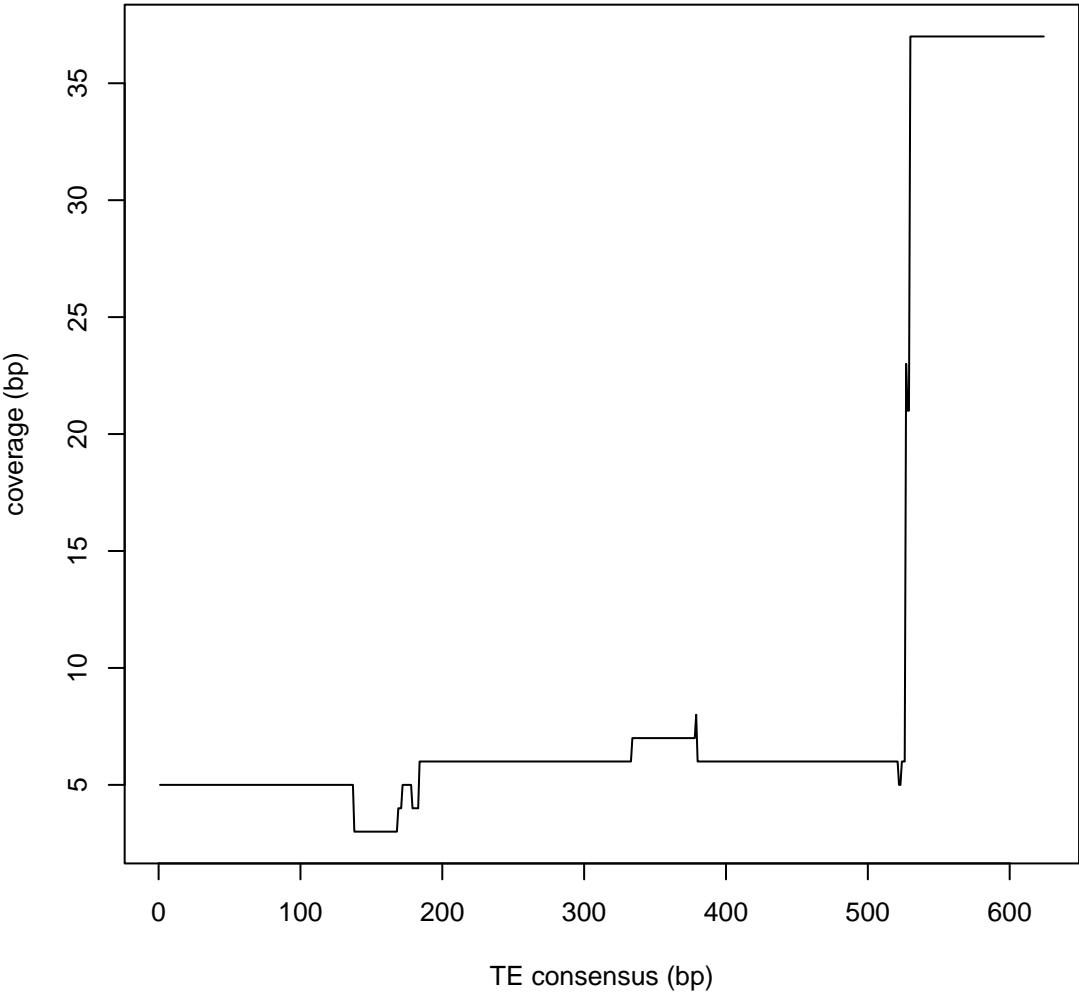


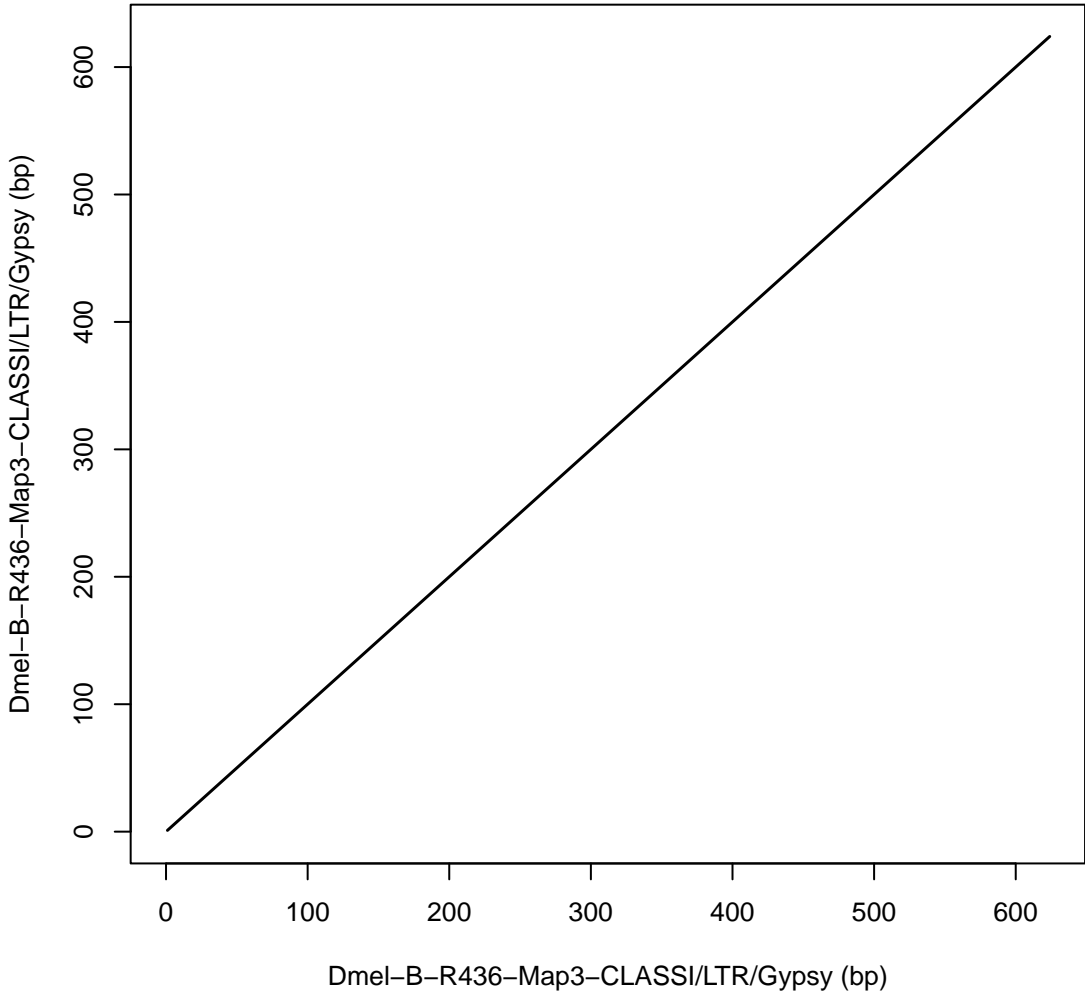
TE: Dmel-B-R436-Map3-CLASSI/LTR/Gypsy
consensus size: 624bp; fragments: 45; full length: 1 (>=561.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

