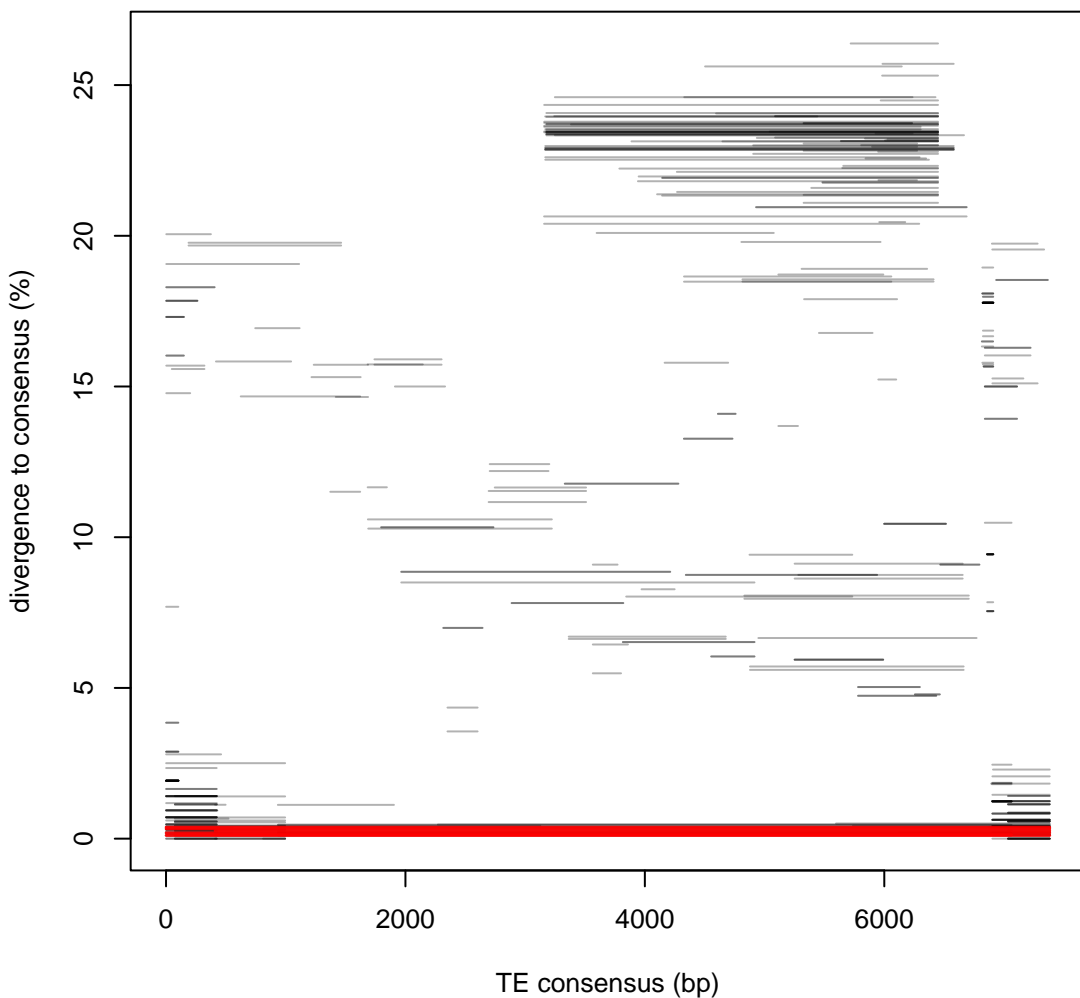
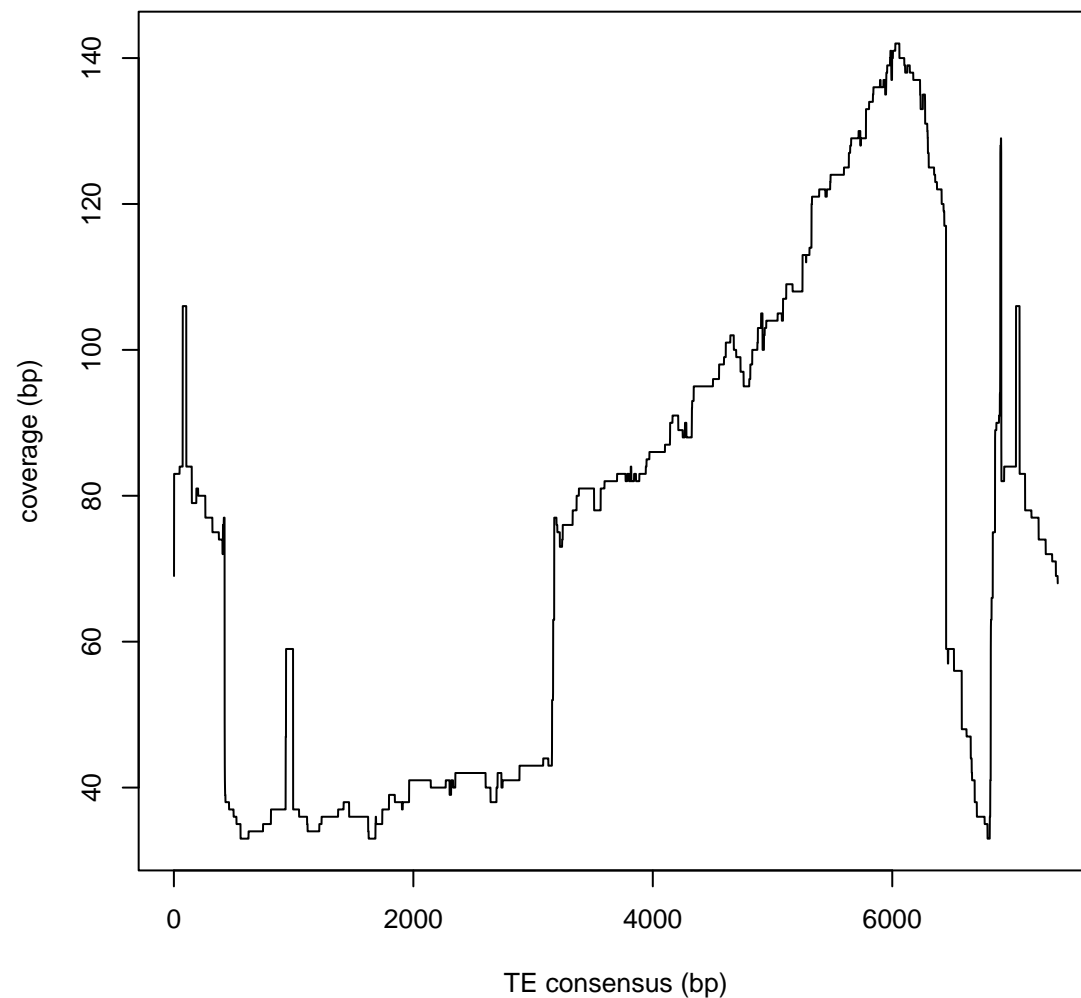


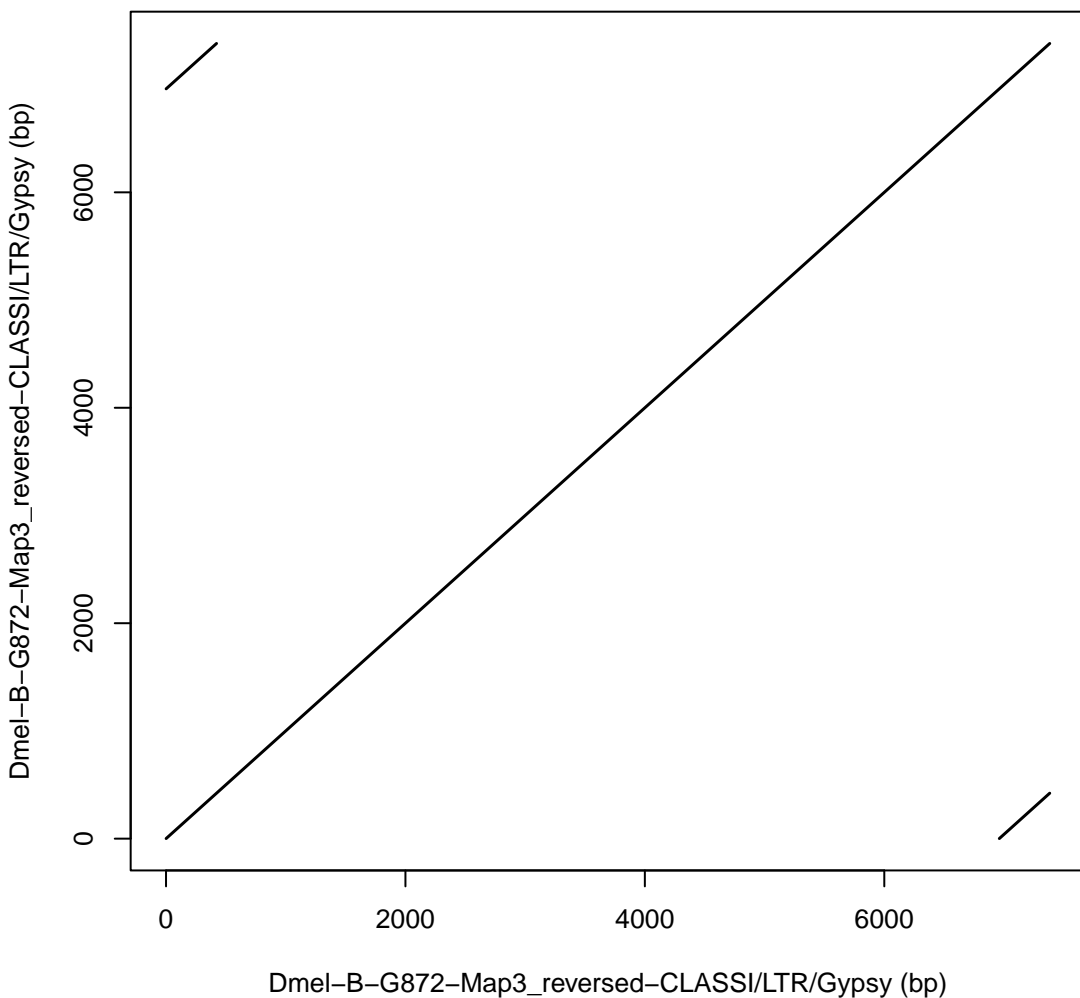
TE: Dmel-B-G872-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 7382bp; fragments: 435; full length: 6 (>=6643.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

