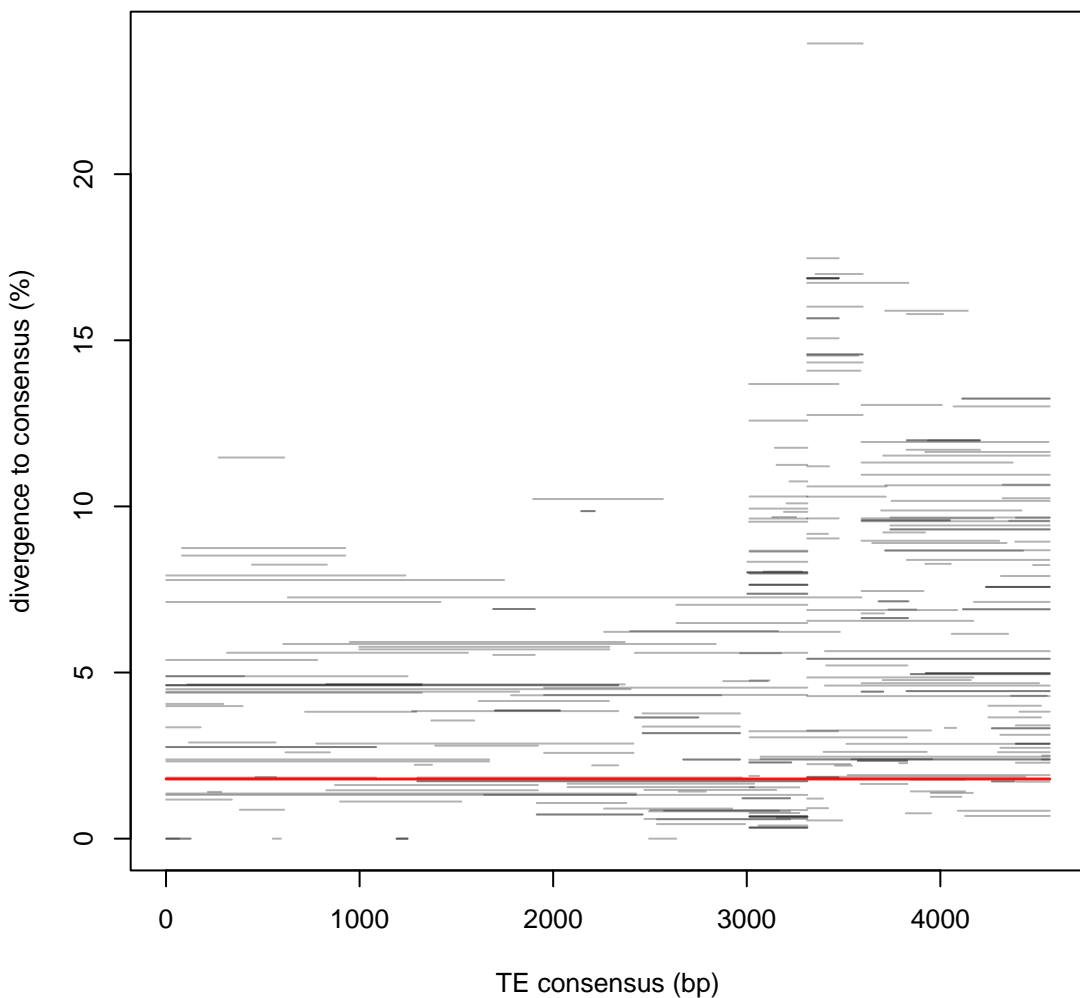
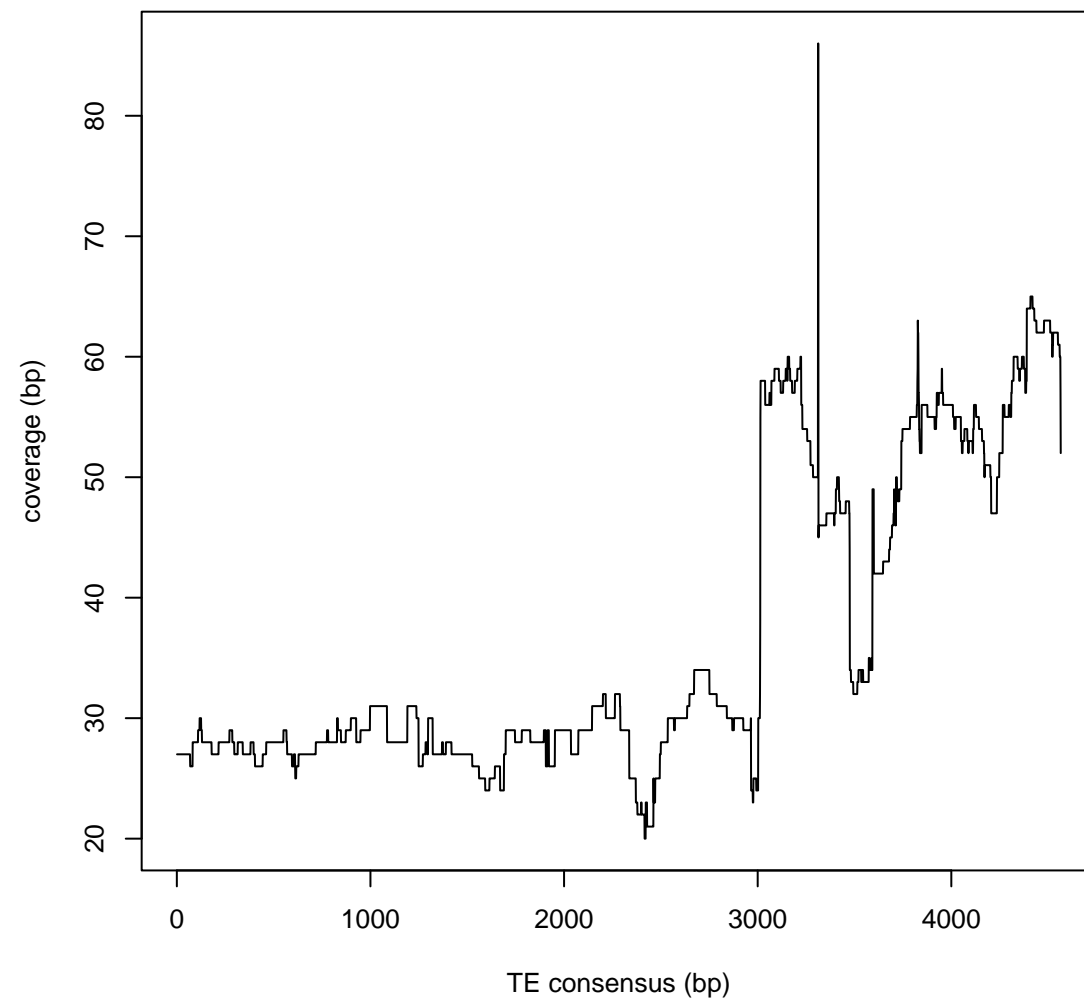


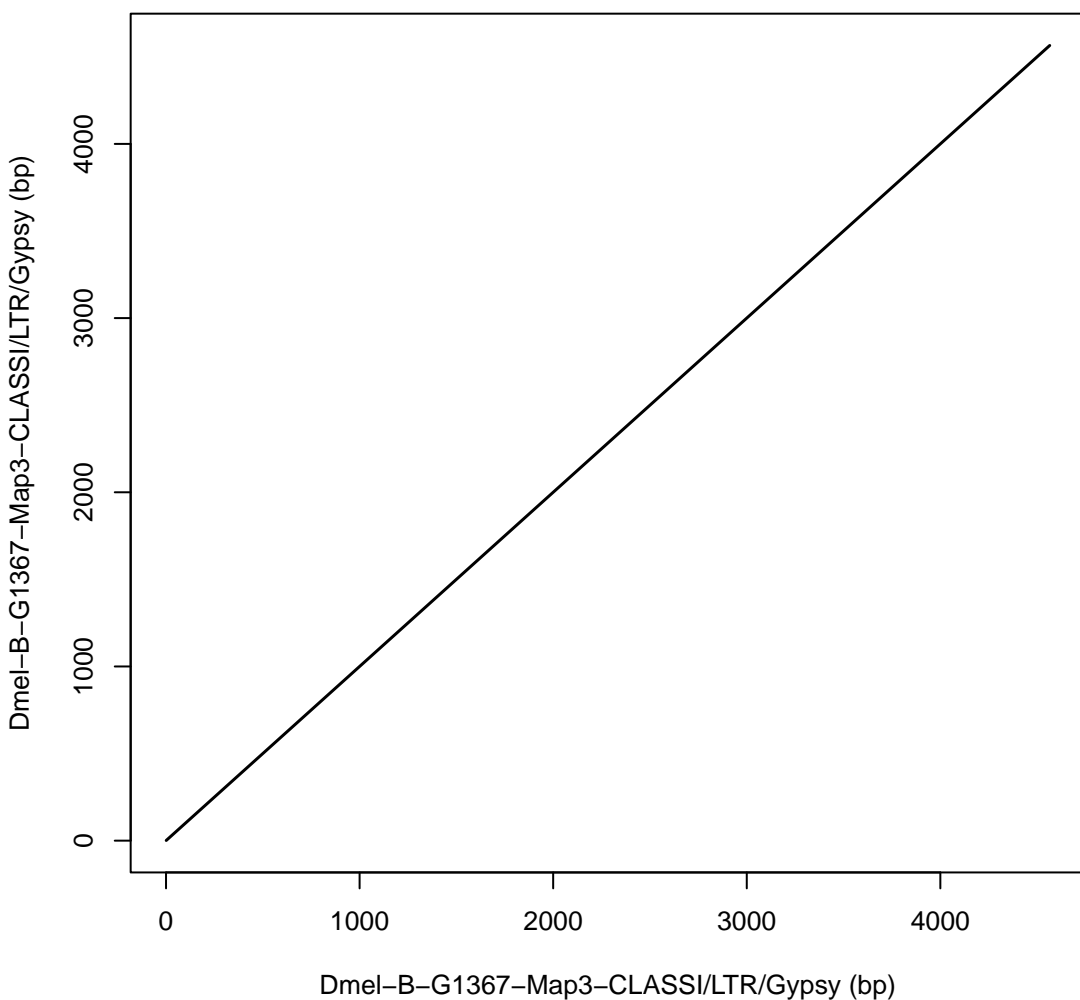
TE: Dmel-B-G1367-Map3-CLASSI/LTR/Gypsy
consensus size: 4565bp; fragments: 292; full length: 1 (>=4108.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

