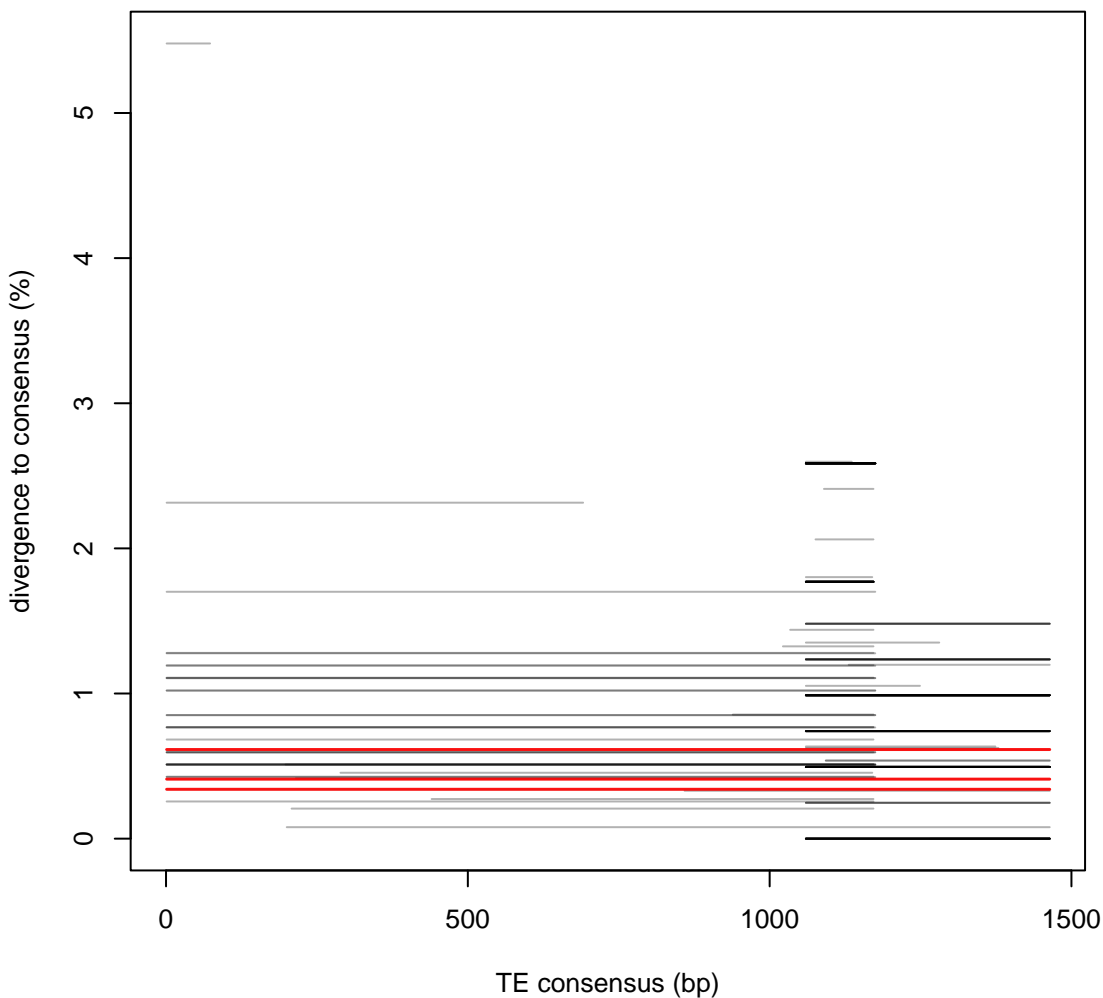
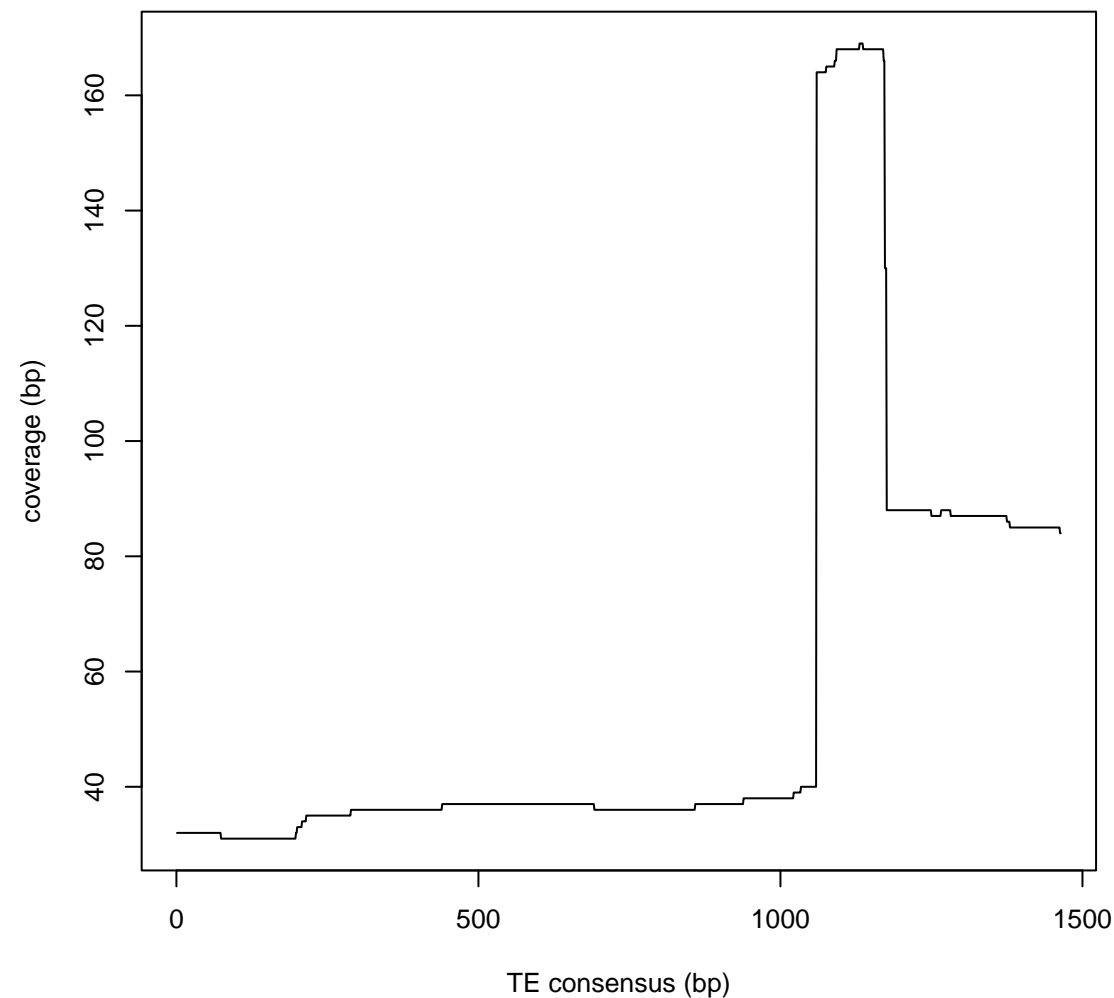


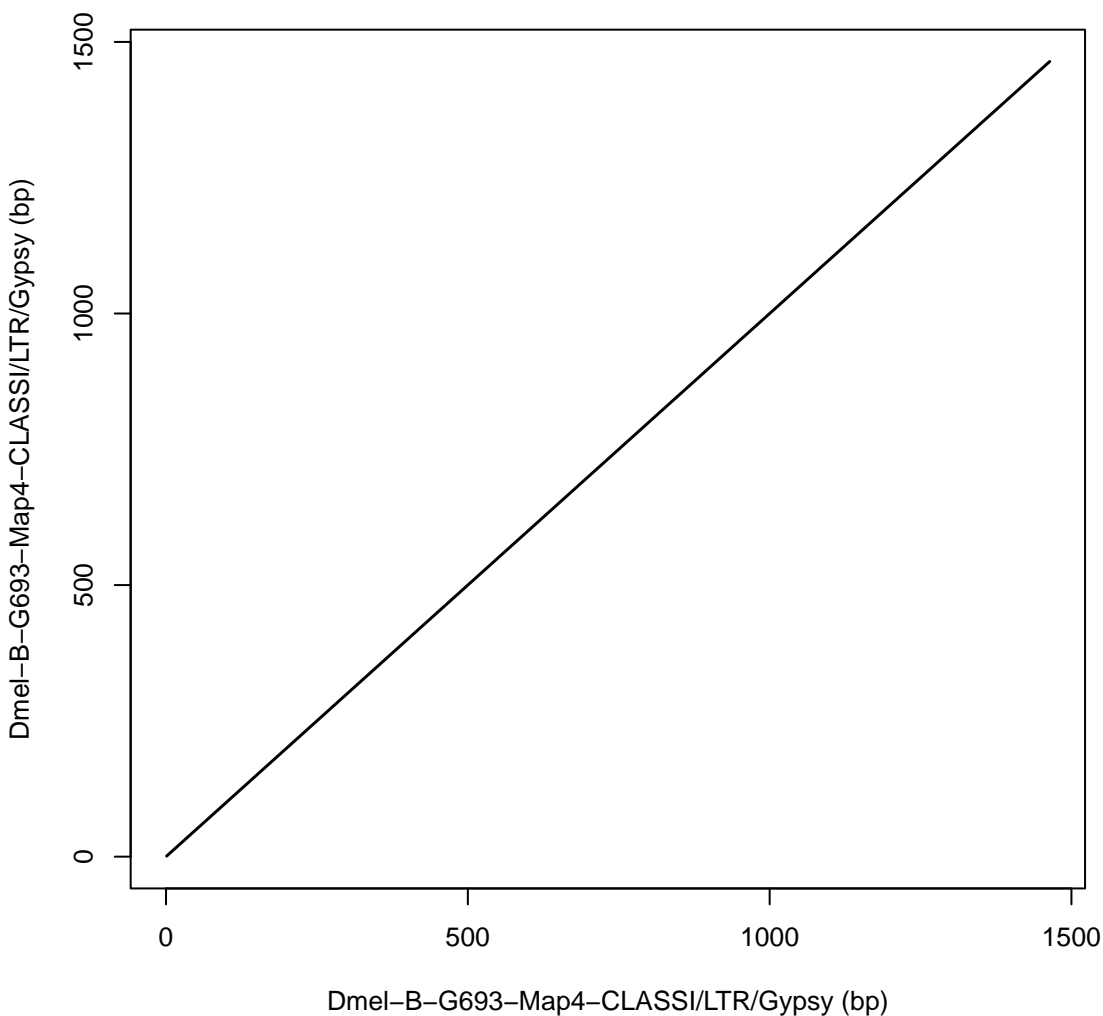
TE: Dmel-B-G693-Map4-CLASSI/LTR/Gypsy  
consensus size: 1464bp; fragments: 172; full length: 3 (>=1317.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

