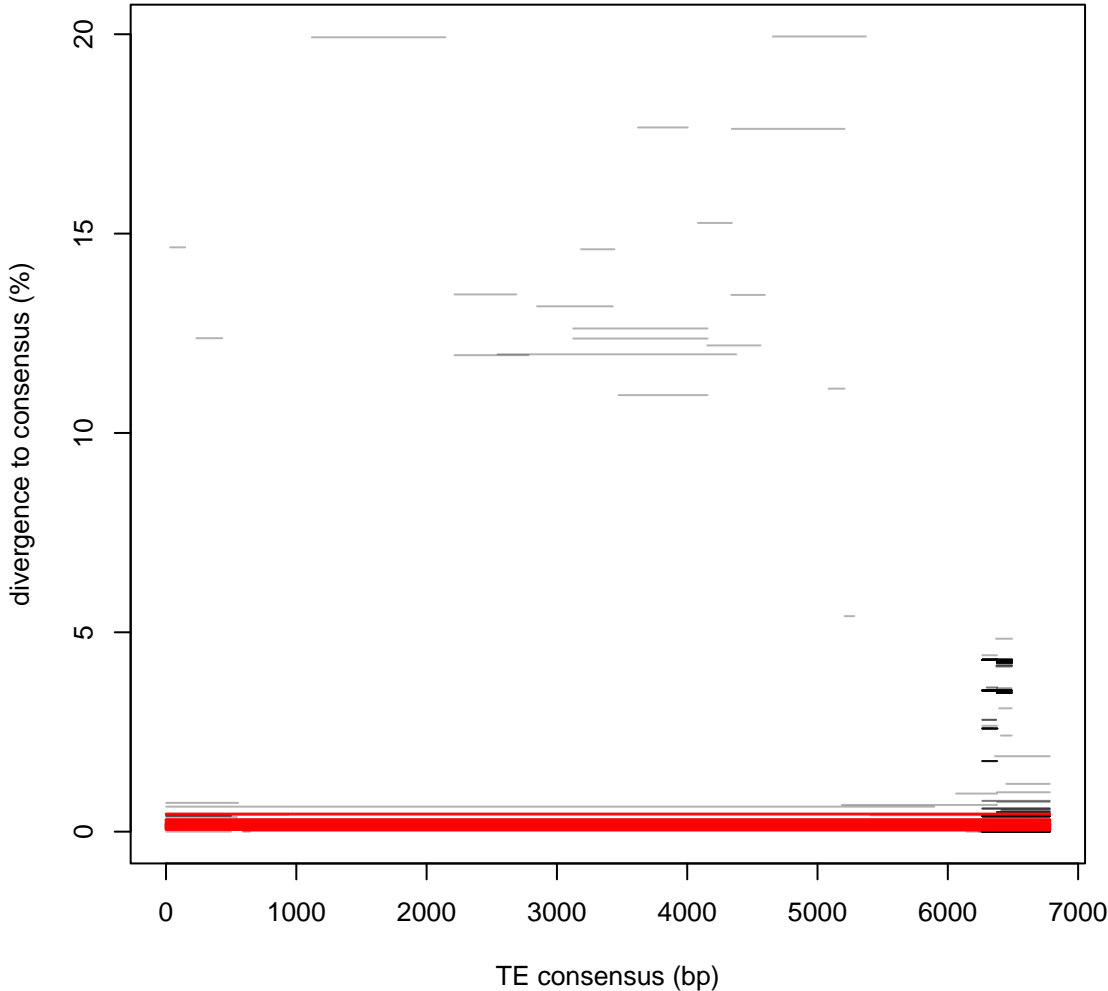
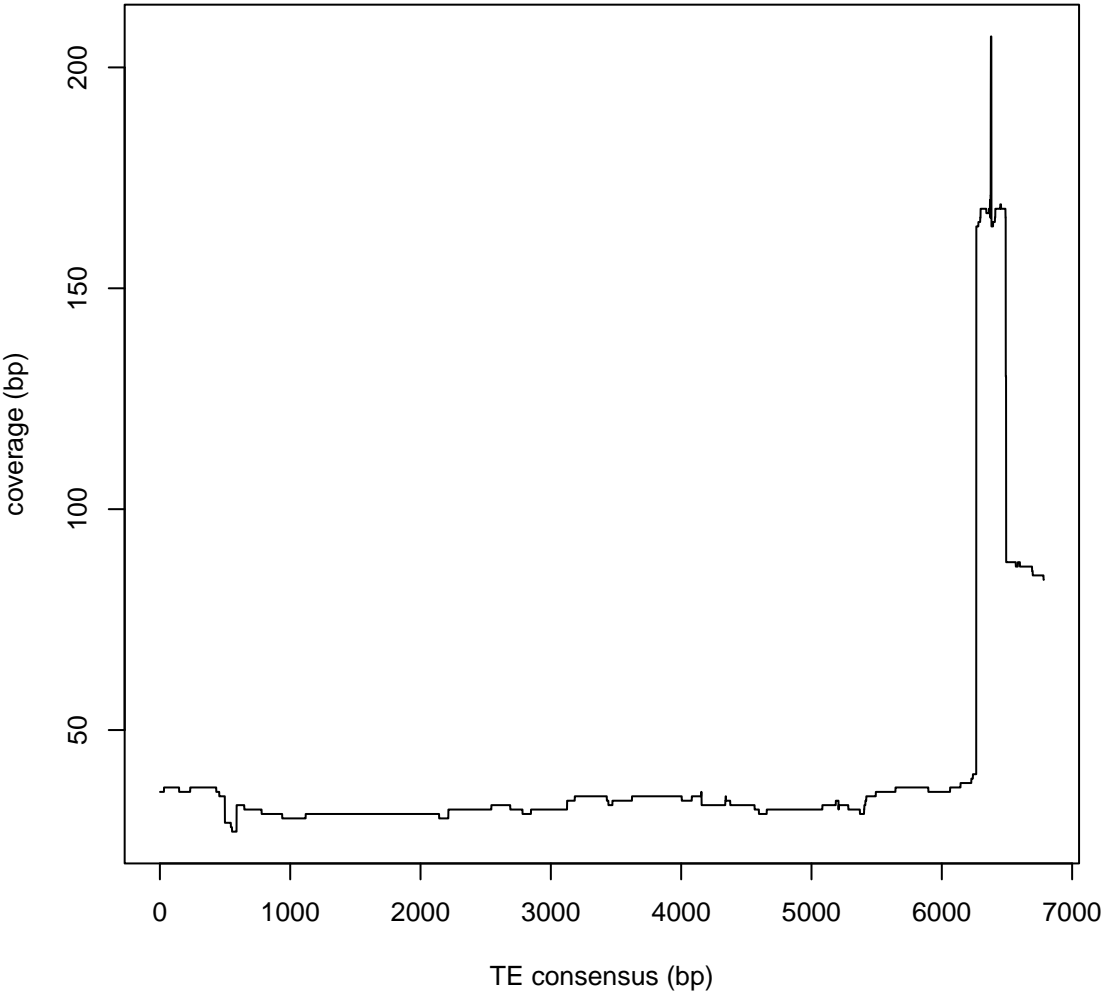


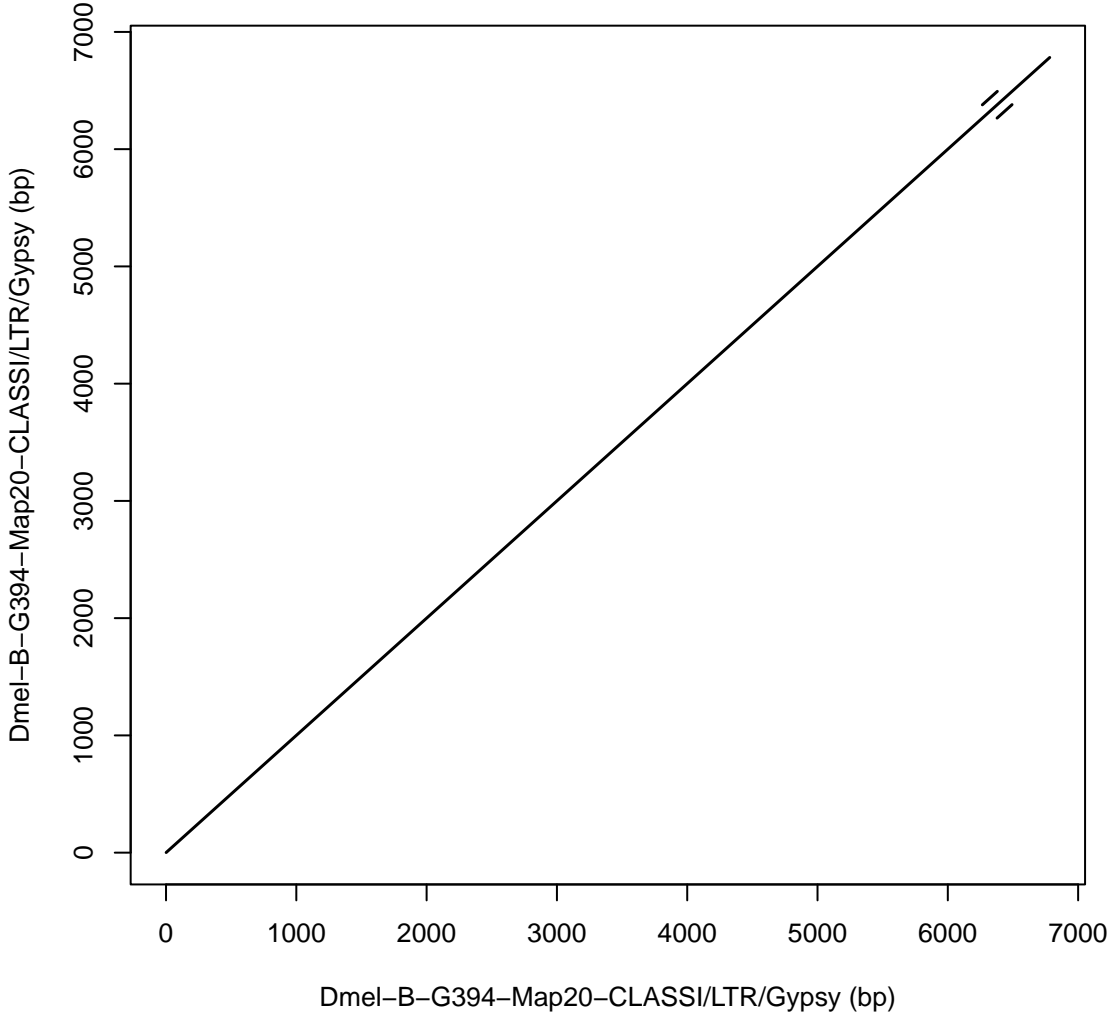
TE: Dmel-B-G394-Map20-CLASSI/LTR/Gypsy  
consensus size: 6782bp; fragments: 292; full length: 28 (>=6103.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

