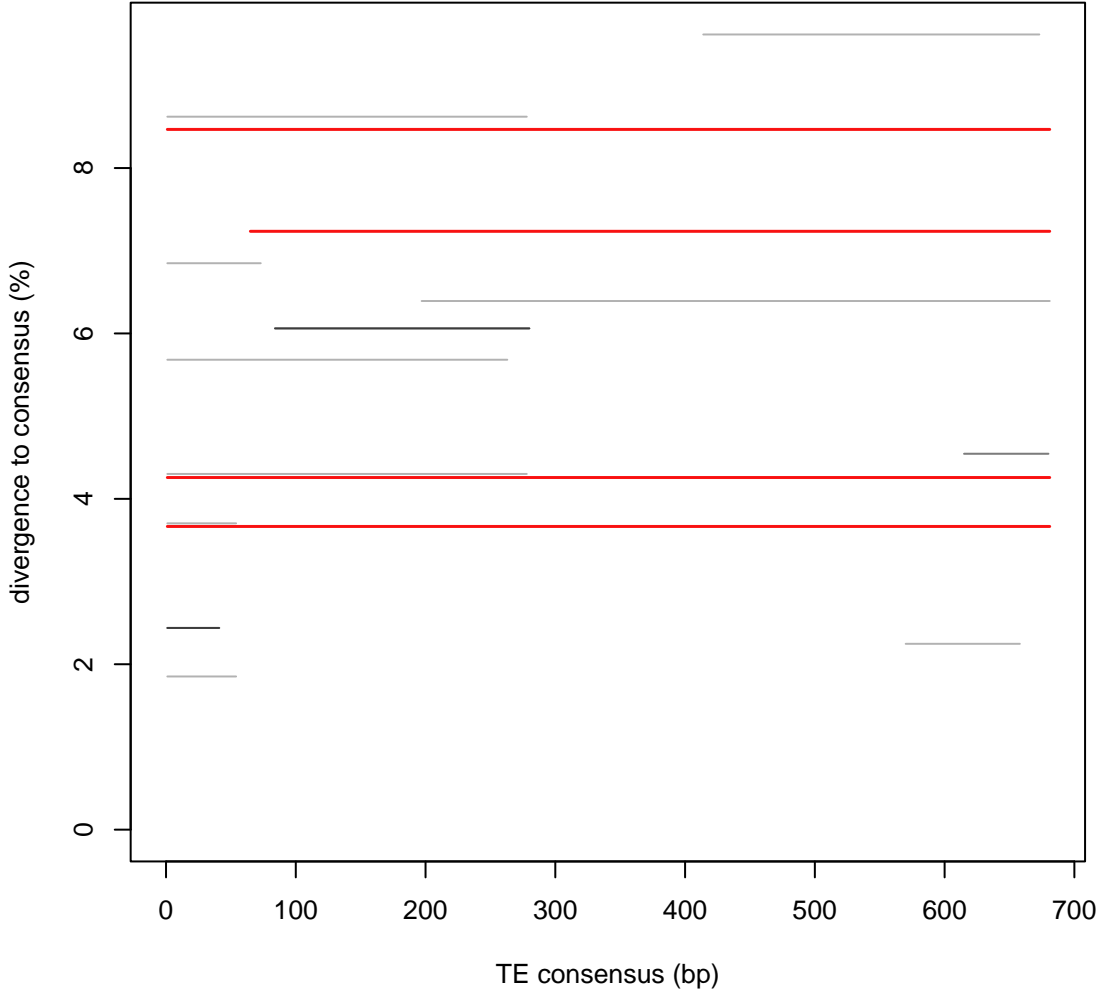
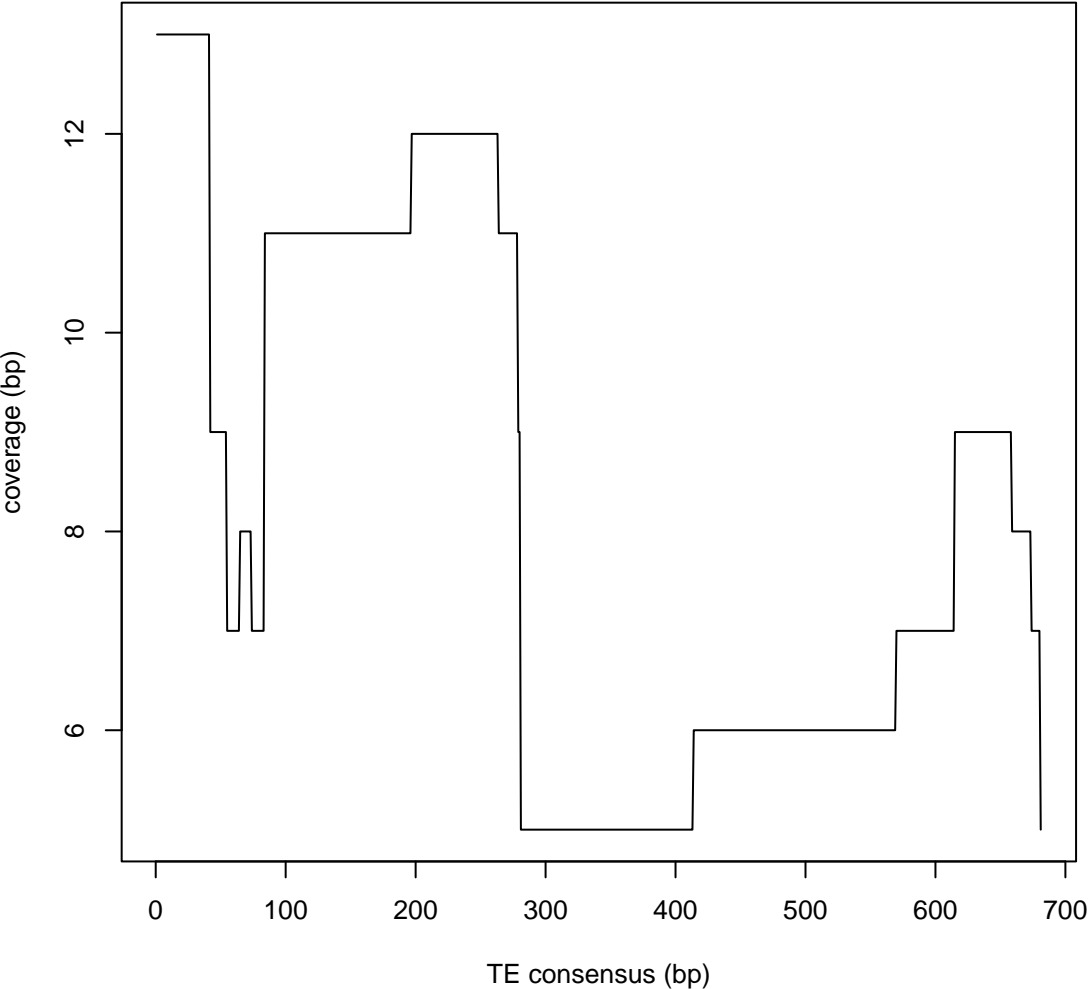


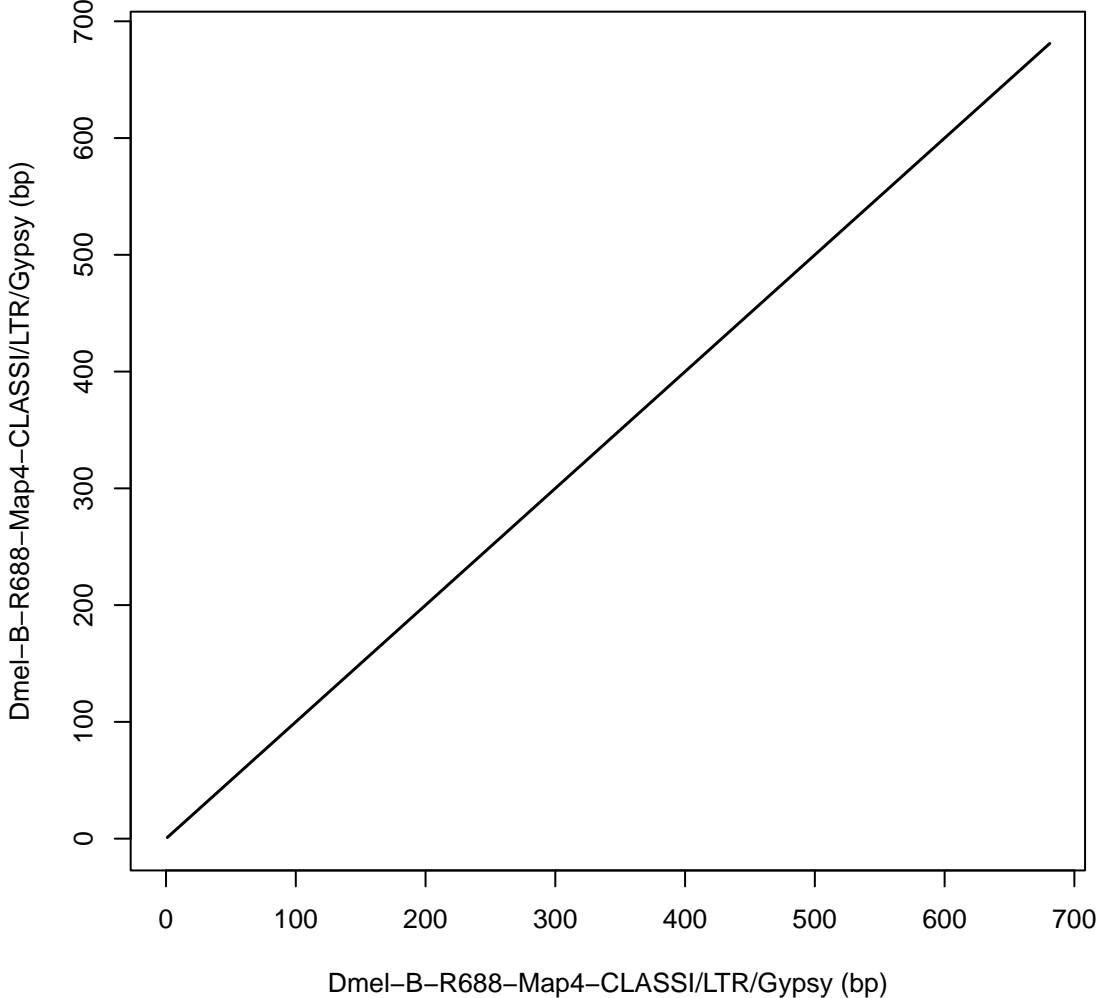
TE: Dmel-B-R688-Map4-CLASSI/LTR/Gypsy  
consensus size: 681bp; fragments: 23; full length: 4 (>=612.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

